# formulas Documentation 

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https://github.com/vinci1it2000/formulas
https://pypi.org/project/formulas/
http://formulas.readthedocs.io/
https://github.com/vinci1it2000/formulas/wiki/
http://github.com/vinci1it2000/formulas/releases/
https://donorbox.org/formulas
excel, formulas, interpreter, compiler, dispatch

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## CHAPTER 1

## What is formulas?

formulas implements an interpreter for Excel formulas, which parses and compile Excel formulas expressions.
Moreover, it compiles Excel workbooks to python and executes without using the Excel COM server. Hence, Excel is not needed.

## CHAPTER 2

Installation

To install it use (with root privileges):

```
$ pip install formulas
```

Or download the last git version and use (with root privileges):

```
python setup.py install
```


### 2.1 Install extras

Some additional functionality is enabled installing the following extras:

- excel: enables to compile Excel workbooks to python and execute using: ExcelModel.
- plot: enables to plot the formula ast and the Excel model.

To install formulas and all extras, do:
\$ pip install formulas[all]

### 2.1.1 What is formulas?

formulas implements an interpreter for Excel formulas, which parses and compile Excel formulas expressions.
Moreover, it compiles Excel workbooks to python and executes without using the Excel COM server. Hence, Excel is not needed.

### 2.1.2 Installation

To install it use (with root privileges):
\$ pip install formulas
Or download the last git version and use (with root privileges):

```
$ python setup.py install
```


### 2.1.2.1 Install extras

Some additional functionality is enabled installing the following extras:

- excel: enables to compile Excel workbooks to python and execute using: ExcelModel.
- plot: enables to plot the formula ast and the Excel model.

To install formulas and all extras, do:

```
$ pip install formulas[all]
```


### 2.1.3 Basic Examples

The following sections will show how to:

- parse a Excel formulas;
- load, compile, and execute a Excel workbook;
- extract a sub-model from a Excel workbook;
- add a custom function.


### 2.1.3.1 Parsing formula

An example how to parse and execute an Excel formula is the following:

```
>>> import formulas
>>> func = formulas.Parser().ast('=(1 + 1) + B3 / A2')[1].compile()
```

To visualize formula model and get the input order you can do the following:

```
>>> list(func.inputs)
['A2', 'B3']
>>> func.plot(view=False) # Set view=True to plot in the default browser.
SiteMap([(=((1 + 1) + (B3 / A2)), SiteMap())])
```



Finally to execute the formula and plot the workflow:

```
>>> func(1, 5)
OperatorArray(7.0, dtype=object)
>>> func.plot(workflow=True, view=False) # Set view=True to plot in the
\hookrightarrowdefault browser.
SiteMap([(=((1 + 1) + (B3 / A2)), SiteMap())])
```



### 2.1.3.2 Excel workbook

An example how to load, calculate, and write an Excel workbook is the following:

```
>>> import formulas
>>> fpath = 'file.xlsx'
>>> xl_model = formulas.ExcelModel().loads(fpath).finish()
>>> xl_model.calculate()
Solution(...)
>>> xl_model.write()
{'EXCEL.XLSX': {Book: <openpyxl.workbook.workbook.Workbook ...>}}
```

Tip: If you have or could have circular references, add circular=True to finish method.

To plot the dependency graph that depict relationships between Excel cells:

```
>>> dsp = xl_model.dsp
>>> dsp.plot(view=False) # Set view=True to plot in the default browser.
SiteMap([(Dispatcher ..., SiteMap())])
```



To compile, execute, and plot a Excel sub-model you can do the following:

```
>>> inputs = ["'[EXCEL.XLSX]DATA'!A2"] # input cells
>>> outputs = ["'[EXCEL.XLSX]DATA'!C2"] # output cells
>>> func = xl_model.compile(inputs, outputs)
>>> func(2).value [0,0]
4.0
>>> func.plot(view=False) # Set view=True to plot in the default browser.
SiteMap([(Dispatcher ..., SiteMap())])
```



### 2.1.3.3 Custom functions

An example how to add a custom function to the formula parser is the following:

```
>>> import formulas
>>> FUNCTIONS = formulas.get_functions()
>>> FUNCTIONS['MYFUNC'] = lambda x, y: 1 + y + x
>>> func = formulas.Parser().ast('=MYFUNC(1, 2)') [1].compile()
>>> func()
4
```


### 2.1.4 Next moves

Things yet to do: implement the missing Excel formulas.

### 2.1.5 Contributing to formulas

If you want to contribute to formulas and make it better, your help is very welcome. The contribution should be sent by a pull request. Next sections will explain how to implement and submit a new excel function:

- clone the repository
- implement a new function/functionality
- open a pull request


### 2.1.5.1 Clone the repository

The first step to contribute to formulas is to clone the repository:

- Create a personal fork of the formulas repository on Github.
- Clone the fork on your local machine. Your remote repo on Github is called origin.
- Add the original repository as a remote called upstream, to maintain updated your fork.
- If you created your fork a while ago be sure to pull upstream changes into your local repository.
- Create a new branch to work on! Branch from dev.


### 2.1.5.2 How to implement a new function

Before coding, study the Excel function that you want to implement. If there is something similar implemented in formulas, try to get inspired by the implemented code (I mean, not reinvent the wheel) and to use numpy. Follow the code style of the project, including indentation. Add or change the documentation as needed. Make sure that you have implemented the full function syntax, including the array syntax.
Test cases are very important. This library uses a data-driven testing approach. To implement a new function I recommend the test-driven development cycle. Hence, when you implement a new function, you should write new test cases in test_cell/TestCell.test_output suite to execute in the cycle loop. When you think that the code is ready, add new raw test in test/test_files/test.xlsx (please follow the standard used for other functions) and run the test_excel/TestExcelModel.test_excel_model. This requires more time but is needed to test the array syntax and to check if the Excel documentation respects the reality.

When all test cases are ok (python setup.py test), open a pull request.
Do do list:

- Study the excel function syntax and behaviour when used as array formula.
- Check if there is something similar implemented in formulas.
- Implement/fix your feature, comment your code.
- Write/adapt tests and run them!

Tip: Excel functions are categorized by their functionality. If you are implementing a new functionality group, add a new module in formula/function and in formula.function. SUBMODULES and a new worksheet in test/test_files/test.xlsx (please respect the format).

Note: A pull request without new test case will not be taken into consideration.

### 2.1.5.3 How to open a pull request

Well done! Your contribution is ready to be submitted:

- Squash your commits into a single commit with git's interactive rebase. Create a new branch if necessary. Always write your commit messages in the present tense. Your commit message should describe what the commit, when applied, does to the code - not what you did to the code.
- Push your branch to your fork on Github (i.e., git push origin dev).
- From your fork open a pull request in the correct branch. Target the project's dev branch!
- Once the pull request is approved and merged you can pull the changes from upstream to your local repo and delete your extra branch(es).


### 2.1.6 Donate

If you want to support the formulas development please donate and add your excel function preferences. The selection of the functions to be implemented is done considering the cumulative donation amount per function collected by the campaign.

Note: The cumulative donation amount per function is calculated as the example:

| Function | Donator 1 | Donator 2 | Donator 3 | TOT | Implementation <br> order |
| :--- | :--- | :--- | :--- | :--- | :---: |
| $\cdot$ | $150 €$ | $120 €$ | $50 €$ | $\bullet$ | $\cdot$ |
| SUM | $50 €$ | $40 €$ | $25 €$ | $125 €$ | 1 st |
| SIN | $50 €$ |  | $25 €$ | $75 €$ | 3 rd |
| TAN | $50 €$ | $40 €$ |  | $90 €$ | 2 nd |
| COS |  | $40 €$ |  | $40 €$ | 4 th |

### 2.1.7 API Reference

The core of the library is composed from the following modules:
It contains a comprehensive list of all modules and classes within formulas.
Modules:

| parser | It provides formula parser class. |
| :--- | :--- |
| builder | It provides AstBuilder class. |
| errors | Defines the formulas exception. |
| tokens | It provides tokens needed to parse the Excel formulas. |
| functions | It provides functions implementations to compile the <br>  <br> Excel functions. |

Table 1 - continued from previous page

| ranges | It provides Ranges class. |
| :--- | :--- |
| cell | It provides Cell class. |
| excel | It provides Excel model class. |

### 2.1.7.1 parser

It provides formula parser class.

## Classes

```
Parser
```

Parser
class Parser

Methods

| ast |
| :--- |
| is_formula |

ast

Parser.ast (expression, context=None)
is_formula

Parser.is_formula (value)
__init__()
Initialize self. See help(type(self)) for accurate signature.

## Attributes

```
filters
    formula_check
    filters
    Parser.filters = [<class 'formulas.tokens.operand.Error'>, <class 'formulas.tokens.ope
    formula_check
    Parser.formula_check = regex.Regex('\n (?P<array>^\\s*{\\s*=\\s*(?P<name>\\S.*)\\s*}\\\
```

ast_builder
alias of formulas.builder.AstBuilder

### 2.1.7.2 builder

It provides AstBuilder class.

## Classes

```
AstBuilder
```


## AstBuilder

```
class AstBuilder (*args, \(d s p=\) None, nodes \(=\) None, match \(=\) None, **kwargs)
```


## Methods

| __init__ | Initialize self. |
| :--- | :--- |
| append | Add an element to the right side of the deque. |
| appendleft | Add an element to the left side of the deque. |
| clear | Remove all elements from the deque. |
| compile | Return a shallow copy of a deque. |
| copy | Extend the right side of the deque with elements <br> from the iterable |
| count | Extend the left side of the deque with elements from <br> the iterable |
| extend |  |
| finish | Raises ValueError if the value is not present. |
| get_node_id | D.insert(index, object) - insert object before index |
| index | Remove and return the rightmost element. |
| insert | Remove and return the leftmost element. |
| pop | D.remove(value) - remove first occurrence of value. |
| popleft | D.reverse ()$-$ reverse $I N$ PLACE |
| remove | Rotate the deque $n$ steps to the right (default n=1). |
| reverse |  |
| rotate |  |

$\qquad$
it

AstBuilder.__init__ (*args, dsp=None, nodes=None, match=None, **kwargs)
Initialize self. See help(type(self)) for accurate signature.

## append

AstBuilder.append (token)
Add an element to the right side of the deque.

```
appendleft
AstBuilder.appendleft()
    Add an element to the left side of the deque.
clear
AstBuilder.clear()
    Remove all elements from the deque.
compile
AstBuilder.compile(references=None, **inputs)
copy
AstBuilder.copy()
    Return a shallow copy of a deque.
count
AstBuilder.count (value) }->\mathrm{ integer - return number of occurrences of value
extend
AstBuilder.extend()
    Extend the right side of the deque with elements from the iterable
extendleft
AstBuilder.extendleft()
    Extend the left side of the deque with elements from the iterable
finish
AstBuilder.finish()
get_node_id
AstBuilder.get_node_id(token)
index
```

AstBuilder.index (value $[$, start $[$, stop $]]) \rightarrow$ integer - return first index of value.

Raises ValueError if the value is not present.
insert

AstBuilder.insert()
D.insert(index, object) - insert object before index
pop

AstBuilder.pop()
Remove and return the rightmost element.

## popleft

```
AstBuilder.popleft()
```

Remove and return the leftmost element.
remove

AstBuilder.remove()
D.remove(value) - remove first occurrence of value.
reverse

AstBuilder.reverse()
D.reverse() - reverse IN PLACE
rotate

AstBuilder.rotate()
Rotate the deque n steps to the right (default $\mathrm{n}=1$ ). If n is negative, rotates left.
__init__ (*args, dsp=None, nodes=None, match=None, **kwargs)
Initialize self. See help(type(self)) for accurate signature.

## Attributes

maxlen

AstBuilder.maxlen
maximum size of a deque or None if unbounded
append (token)
Add an element to the right side of the deque.

### 2.1.7.3 errors

Defines the formulas exception.

## Exceptions

| BaseError |
| :--- |
| BroadcastError |
| FormulaError |
| FoundError |
| FunctionError |
| ParenthesesError |
| RangeValueError |
| TokenError |

## BaseError

exception BaseError (*args)

BroadcastError
exception BroadcastError (*args)

## FormulaError

exception FormulaError (*args)

FoundError
exception FoundError (*args, err=None, **kwargs)

## FunctionError

exception FunctionError (*args)

ParenthesesError
exception ParenthesesError (*args)

RangeValueError
exception RangeValueError (*args)

## TokenError

```
exception TokenError(*args)
```


### 2.1.7.4 tokens

It provides tokens needed to parse the Excel formulas.
Sub-Modules:

| function | It provides Function classes. |
| :--- | :--- |
| operand | It provides Operand classes. |
| operator | It provides Operator classes. |
| parenthesis | It provides Parenthesis class. |

## function

It provides Function classes.

## Classes

Array

## Array

class Array (s, context=None)

Methods

| _init__ | Initialize self. |
| :--- | :--- |
| ast |  |
| compile |  |
| match |  |
| process |  |
| set_expr |  |
| update_input_tokens |  |

__init

Array.__init__ ( $s$, context=None)
Initialize self. See help(type(self)) for accurate signature.
ast

Array.ast (tokens, stack, builder, check_n=<function Array.<lambda>>)

```
compile
Array.compile()
match
Array.match(s)
process
Array.process(match, context=None)
set_expr
Array.set_expr(*tokens)
update_input_tokens
Array.update_input_tokens(*tokens)
__init__(s, context=None)
        Initialize self. See help(type(self)) for accurate signature.
```


## Attributes

| name |
| :--- |
| node_id |
| name |
| Array.name |
| node_id |
| Array.node_id |

## Function

class Function ( $s$, context $=$ None)

## Methods

| _init__ | Initialize self. |
| :--- | :--- |
| ast |  |

Table 13 - continued from previous page

| compile |
| :--- |
| match |
| process |
| set_expr |
| update_input_tokens |

__init__
Function.__init__(s, context=None)
Initialize self. See help(type(self)) for accurate signature.
ast

Function.ast (tokens, stack, builder, check_n=<function Function.<lambda>>)
compile

Function.compile()
match

Function.match (s)
process

Function.process (match, context=None)
set_expr
Function.set_expr(*tokens)
update_input_tokens

Function.update_input_tokens (*tokens)
__init__(s, context=None) Initialize self. See help(type(self)) for accurate signature.

## Attributes

| name |
| :--- |
| node_id |

name

Function. name
node_id

Function.node_id
operand

It provides Operand classes.

## Functions

| fast_range2parts |
| :--- |
| fast_range2parts_v1 |
| fast_range2parts_v2 |
| fast_range2parts_v3 |
| range2parts |

fast_range2parts

```
fast_range2parts(**kw)
```

fast_range2parts_v1
fast_range2parts_v1 (rl, cl, excel, sheet=")
fast_range2parts_v2
fast_range2parts_v2 $(r 1, c 1, r 2, c 2$, excel, sheet $=$ " $)$
fast_range2parts_v3
fast_range2parts_v3 ( $r 1, n 1, r 2, n 2$, excel, sheet=")
range2parts
range2parts (outputs, **inputs)

Classes

Table 16 - continued from previous page

| Number |
| :--- |
| Operand |
| Range |
| String |
| XlError |

## Error

class Error (s, context=None)

## Methods

| _init__ | Initialize self. |
| :--- | :--- |
| ast |  |
| compile |  |
| match |  |
| process |  |
| set_expr |  |
| update_input_tokens |  |

__init

Error.__init__(s, context=None) Initialize self. See help(type(self)) for accurate signature.
ast

Error.ast (tokens, stack, builder)
compile

Error.compile()
match

Error.match ( $s$ )
process

Error.process (match, context=None)
set_expr
Error.set_expr (*tokens)
update_input_tokens

Error.update_input_tokens (*tokens)
__init__ ( $s$, context=None)
Initialize self. See help(type(self)) for accurate signature.

Attributes

| errors |
| :--- |
| k |
| name |
| node_id |

errors

Error.errors = \{'\#DIV/O!': \#DIV/O!, '\#N/A': \#N/A, '\#NAME?': \#NAME?, '\#NULL!': \#NULL!,
k

Error.k = '\#N/A'
name

Error. name
node_id

Error.node_id

Number
class Number ( $s$, context $=$ None)

Methods

| _init__ | Initialize self. |
| :--- | :--- |
| ast |  |
| compile |  |
| match |  |
| process |  |
| set_expr |  |
| update_input_tokens |  |

```
    __init
    Number.__init__ ( }s\mathrm{ , context=None)
        Initialize self. See help(type(self)) for accurate signature.
    ast
    Number.ast (tokens, stack, builder)
    compile
    Number.compile()
    match
    Number.match(s)
    process
    Number.process (match, context=None)
    set_expr
    Number.set_expr(*tokens)
    update_input_tokens
    Number.update_input_tokens(*tokens)
    __init__ (s, context=None)
        Initialize self. See help(type(self)) for accurate signature.
```

Attributes
name
node_id
name

Number. name
node_id

Number.node_id

## Operand

class Operand (s, context=None)

Methods

| init__ Initialize self. |  |
| :--- | :--- |
| ast |  |
| match |  |
| process |  |
| set_expr |  |
| update_input_tokens |  |

init $\qquad$
Operand.__init__( $s$, context=None)
Initialize self. See help(type(self)) for accurate signature.
ast

Operand.ast (tokens, stack, builder)
match

Operand.match (s)
process
Operand.process (match, context=None)
set_expr

Operand.set_expr (*tokens)
update_input_tokens
Operand.update_input_tokens (*tokens)
__init__(s, context=None)
Initialize self. See help(type(self)) for accurate signature.

## Attributes

| name |
| :--- | :--- |
| node_id |

name

Operand.name
node_id

Operand.node_id

Range
class Range ( $s$, context=None)

Methods

| _init_ | Initialize self. |
| :--- | :--- |
| ast |  |
| compile |  |
| match |  |
| process |  |
| set_expr |  |
| update_input_tokens |  |

init

Range.__init__ ( $s$, context=None) Initialize self. See help(type(self)) for accurate signature.
ast

Range.ast (tokens, stack, builder)
compile

Range. compile()
match

Range.match (s)
process

Range.process (match, context=None)

```
set_expr
Range.set_expr(*tokens)
update_input_tokens
Range.update_input_tokens(*tokens)
__init__(s, context=None)
        Initialize self. See help(type(self)) for accurate signature.
```


## Attributes

```
name (
node_id
    name
    Range.name
    node_id
    Range.node_id
```


## String

class String ( $s$, context $=$ None )

Methods

| _init__ | Initialize self. |
| :--- | :--- |
| ast |  |
| compile |  |
| match |  |
| process |  |
| set_expr |  |
| update_input_tokens |  |

__init__

String.__init__( $s$, context=None)
Initialize self. See help(type(self)) for accurate signature.
ast

String.ast (tokens, stack, builder)

```
compile
String.compile()
match
String.match(s)
process
String.process(match, context=None)
set_expr
String.set_expr(*tokens)
update_input_tokens
String.update_input_tokens(*tokens)
__init__(s, context=None)
        Initialize self. See help(type(self)) for accurate signature.
```


## Attributes

```
name
node_id
    name
    String.name
    node_id
    String.node_id
```

XIError
class XlError
Methods
capitalize Return a capitalized version of S, i.e.

Table 27 - continued from previous page

| casefold | Return a version of $S$ suitable for caseless comparisons. |
| :---: | :---: |
| center | Return S centered in a string of length width. |
| count | Return the number of non-overlapping occurrences of substring sub in string S[start:end]. |
| encode | Encode S using the codec registered for encoding. |
| endswith | Return True if S ends with the specified suffix, False otherwise. |
| expandtabs | Return a copy of S where all tab characters are expanded using spaces. |
| find | Return the lowest index in $S$ where substring sub is found, such that sub is contained within S [start:end]. |
| format | Return a formatted version of S, using substitutions from args and kwargs. |
| format_map | Return a formatted version of S, using substitutions from mapping. |
| index | Like S.find() but raise ValueError when the substring is not found. |
| isalnum | Return True if all characters in S are alphanumeric and there is at least one character in S, False otherwise. |
| isalpha | Return True if all characters in S are alphabetic and there is at least one character in S, False otherwise. |
| isdecimal | Return True if there are only decimal characters in S, False otherwise. |
| isdigit | Return True if all characters in S are digits and there is at least one character in S, False otherwise. |
| isidentifier | Return True if S is a valid identifier according to the language definition. |
| islower | Return True if all cased characters in S are lowercase and there is at least one cased character in S, False otherwise. |
| isnumeric | Return True if there are only numeric characters in S, False otherwise. |
| isprintable | Return True if all characters in S are considered printable in repr() or S is empty, False otherwise. |
| isspace | Return True if all characters in $S$ are whitespace and there is at least one character in S, False otherwise. |
| istitle | Return True if S is a titlecased string and there is at least one character in S, i.e. |
| isupper | Return True if all cased characters in S are uppercase and there is at least one cased character in S, False otherwise. |
| join | Return a string which is the concatenation of the strings in the iterable. |
| ljust | Return S left-justified in a Unicode string of length width. |
| lower | Return a copy of the string S converted to lowercase. |
| lstrip | Return a copy of the string $S$ with leading whitespace removed. |
| maketrans | Return a translation table usable for str.translate(). |
|  | Continued on next page |

Table 27 - continued from previous page

| partition | Search for the separator sep in S, and return the part <br> before it, the separator itself, and the part after it. |
| :--- | :--- |
| replace | Return a copy of S with all occurrences of substring <br> old replaced by new. |
| rfind | Return the highest index in S where substring sub is <br> found, such that sub is contained within S[start:end]. |
| rindex | Like S.rfind() but raise ValueError when the sub- <br> string is not found. |
| rjust | Return S right-justified in a string of length width. |
| rpartition | Search for the separator sep in S, starting at the end <br> of S, and return the part before it, the separator itself, <br> and the part after it. |
| rsplit | Return a list of the words in S, using sep as the de- <br> limiter string, starting at the end of the string and <br> working to the front. |
| rstrip | Return a copy of the string S with trailing whitespace <br> removed. |
| split | Return a list of the words in S, using sep as the de- <br> limiter string. |
| splitlines | Return a list of the lines in S, breaking at line bound- <br> aries. |
| startswith | Return True if S starts with the specified prefix, False <br> otherwise. |
| strip | Return a copy of the string S with leading and trailing <br> whitespace removed. |
| swapcase | Return a copy of S with uppercase characters con- <br> verted to lowercase and vice versa. |
| translate | Return a titlecased version of S, i.e. |
| zfill | Return a copy of the string S in which each character <br> has been mapped through the given translation table. |
| Return a copy of S converted to uppercase. |  |

## capitalize

XlError.capitalize() $\rightarrow$ str
Return a capitalized version of $S$, i.e. make the first character have upper case and the rest lower case.
casefold

XlError.casefold() $\rightarrow$ str
Return a version of $S$ suitable for caseless comparisons.
center

XlError.center (width[, fillchar]) $\rightarrow \mathrm{str}$
Return $S$ centered in a string of length width. Padding is done using the specified fill character (default is a space)

## count

XlError. count (sub[, start $[$, end $]]) \rightarrow$ int
Return the number of non-overlapping occurrences of substring sub in string S[start:end]. Optional arguments start and end are interpreted as in slice notation.
encode

XlError.encode (encoding='utf-8', errors='strict') $\rightarrow$ bytes
Encode S using the codec registered for encoding. Default encoding is 'utf- 8 '. errors may be given to set a different error handling scheme. Default is 'strict' meaning that encoding errors raise a UnicodeEncodeError. Other possible values are 'ignore', 'replace' and 'xmlcharrefreplace' as well as any other name registered with codecs.register_error that can handle UnicodeEncodeErrors.

## endswith

XlError.endswith (suffix[, start $[$, end $]]) \rightarrow$ bool
Return True if S ends with the specified suffix, False otherwise. With optional start, test $S$ beginning at that position. With optional end, stop comparing $S$ at that position. suffix can also be a tuple of strings to try.

## expandtabs

XlError.expandtabs (tabsize $=8$ ) $\rightarrow$ str
Return a copy of $S$ where all tab characters are expanded using spaces. If tabsize is not given, a tab size of 8 characters is assumed.

## find

XlError.find $(\operatorname{sub}[$, start $[$, end $]]) \rightarrow$ int
Return the lowest index in $S$ where substring sub is found, such that sub is contained within S [start:end]. Optional arguments start and end are interpreted as in slice notation.

Return -1 on failure.

## format

XlError.format (*args, **kwargs) $\rightarrow$ str
Return a formatted version of S, using substitutions from args and kwargs. The substitutions are identified by braces (' $\{$ ' and ' $\}$ ').

## format_map

XlError.format_map (mapping) $\rightarrow$ str
Return a formatted version of $S$, using substitutions from mapping. The substitutions are identified by braces ('\{' and '\}').
index

XlError.index $(\operatorname{sub}[, \operatorname{start}[$, end $]]) \rightarrow$ int
Like S.find() but raise ValueError when the substring is not found.
isalnum

XlError.isalnum() $\rightarrow$ bool
Return True if all characters in $S$ are alphanumeric and there is at least one character in $S$, False otherwise.

## isalpha

XlError.isalpha() $\rightarrow$ bool
Return True if all characters in S are alphabetic and there is at least one character in S , False otherwise.
isdecimal

XlError.isdecimal () $\rightarrow$ bool
Return True if there are only decimal characters in S, False otherwise.
isdigit

XlError.isdigit() $\rightarrow$ bool
Return True if all characters in S are digits and there is at least one character in S , False otherwise.
isidentifier

XlError.isidentifier() $\rightarrow$ bool
Return True if S is a valid identifier according to the language definition.
Use keyword.iskeyword() to test for reserved identifiers such as "def" and "class".
islower

XlError.islower() $\rightarrow$ bool
Return True if all cased characters in $S$ are lowercase and there is at least one cased character in S, False otherwise.
isnumeric

XlError.isnumeric() $\rightarrow$ bool
Return True if there are only numeric characters in S, False otherwise.

## isprintable

XlError.isprintable() $\rightarrow$ bool
Return True if all characters in $S$ are considered printable in repr() or $S$ is empty, False otherwise.
isspace

XlError.isspace () $\rightarrow$ bool
Return True if all characters in $S$ are whitespace and there is at least one character in S, False otherwise.
istitle

XlError.istitle() $\rightarrow$ bool
Return True if $S$ is a titlecased string and there is at least one character in S, i.e. upper- and titlecase characters may only follow uncased characters and lowercase characters only cased ones. Return False otherwise.

## isupper

XlError.isupper () $\rightarrow$ bool
Return True if all cased characters in $S$ are uppercase and there is at least one cased character in S, False otherwise.

## join

XlError.join (iterable) $\rightarrow$ str
Return a string which is the concatenation of the strings in the iterable. The separator between elements is $S$.
ljust

XlError.ljust (width[, fillchar]) $\rightarrow$ str
Return S left-justified in a Unicode string of length width. Padding is done using the specified fill character (default is a space).

Iower

XlError.lower () $\rightarrow$ str
Return a copy of the string $S$ converted to lowercase.

Istrip

XlError.lstrip $([$ chars $]) \rightarrow$ str
Return a copy of the string $S$ with leading whitespace removed. If chars is given and not None, remove characters in chars instead.

## maketrans

## static XlError.maketrans()

Return a translation table usable for str.translate().
If there is only one argument, it must be a dictionary mapping Unicode ordinals (integers) or characters to Unicode ordinals, strings or None. Character keys will be then converted to ordinals. If there are two arguments, they must be strings of equal length, and in the resulting dictionary, each character in x will be mapped to the character at the same position in $y$. If there is a third argument, it must be a string, whose characters will be mapped to None in the result.

## partition

XlError.partition (sep) -> (head, sep, tail)
Search for the separator sep in S, and return the part before it, the separator itself, and the part after it. If the separator is not found, return $S$ and two empty strings.

## replace

XlError. replace (old, new $[$, count $]$ ) $\rightarrow$ str
Return a copy of $S$ with all occurrences of substring old replaced by new. If the optional argument count is given, only the first count occurrences are replaced.
rfind

XlError.rfind (sub[, start $[$, end $]]) \rightarrow$ int
Return the highest index in S where substring sub is found, such that sub is contained within S [start:end]. Optional arguments start and end are interpreted as in slice notation.

Return -1 on failure.
rindex

XlError. rindex $(\operatorname{sub}[$, start $[$, end $]]) \rightarrow$ int
Like S.rfind() but raise ValueError when the substring is not found.
rjust

XlError.rjust (width[, fillchar]) $\rightarrow$ str
Return $S$ right-justified in a string of length width. Padding is done using the specified fill character (default is a space).

## rpartition

XlError.rpartition (sep) -> (head, sep, tail)
Search for the separator sep in S, starting at the end of S, and return the part before it, the separator itself, and the part after it. If the separator is not found, return two empty strings and $S$.
rsplit
XlError.rsplit (sep=None, maxsplit=-1) $\rightarrow$ list of strings
Return a list of the words in S, using sep as the delimiter string, starting at the end of the string and working to the front. If maxsplit is given, at most maxsplit splits are done. If sep is not specified, any whitespace string is a separator.
rstrip

XlError.rstrip $([$ chars $]) \rightarrow$ str
Return a copy of the string S with trailing whitespace removed. If chars is given and not None, remove characters in chars instead.
split
XlError.split (sep=None, maxsplit=-1) $\rightarrow$ list of strings
Return a list of the words in S , using sep as the delimiter string. If maxsplit is given, at most maxsplit splits are done. If sep is not specified or is None, any whitespace string is a separator and empty strings are removed from the result.

## splitlines

XlError.splitlines ([keepends $]$ ) $\rightarrow$ list of strings
Return a list of the lines in S, breaking at line boundaries. Line breaks are not included in the resulting list unless keepends is given and true.

## startswith

XlError.startswith (prefix [, start [, end $]]$ ) $\rightarrow$ bool
Return True if S starts with the specified prefix, False otherwise. With optional start, test S beginning at that position. With optional end, stop comparing $S$ at that position. prefix can also be a tuple of strings to try.
strip
XlError.strip ([chars]) $\rightarrow$ str
Return a copy of the string $S$ with leading and trailing whitespace removed. If chars is given and not None, remove characters in chars instead.

## swapcase

XlError.swapcase () $\rightarrow$ str
Return a copy of S with uppercase characters converted to lowercase and vice versa.
title

XlError.title() $\rightarrow$ str
Return a titlecased version of S , i.e. words start with title case characters, all remaining cased characters have lower case.

## translate

XlError.translate (table) $\rightarrow$ str
Return a copy of the string $S$ in which each character has been mapped through the given translation table. The table must implement lookup/indexing via __getitem_ , for instance a dictionary or list, mapping Unicode ordinals to Unicode ordinals, strings, or None. If this operation raises LookupError, the character is left untouched. Characters mapped to None are deleted.
upper

XlError. upper () $\rightarrow$ str
Return a copy of $S$ converted to uppercase.
zfill

XlError. $\mathbf{z f i l l}$ (width) $\rightarrow$ str
Pad a numeric string $S$ with zeros on the left, to fill a field of the specified width. The string $S$ is never truncated.
__init__()
Initialize self. See help(type(self)) for accurate signature.

## operator

It provides Operator classes.

## Classes

| Intersect |
| :--- |
| Operator |
| OperatorToken |
| Separator |
| Intersect |
| class Intersect $(s$, context $=$ None $)$ |

## Methods

```
\begin{tabular}{ll}
\hline _init__ & Initialize self. \\
\hline ast & \\
\hline compile & \\
\hline match & \\
\hline process & \\
\hline set_expr & \\
\hline update_input_tokens & \\
\hline update_name & \\
\hline
\end{tabular}
    __init
```

$\qquad$

```
    Intersect.__init__ (s,context=None)
        Initialize self. See help(type(self)) for accurate signature.
    ast
    Intersect.ast (tokens, stack, builder)
    compile
    Intersect.compile()
    match
    Intersect.match(s)
    process
    Intersect.process (match, context=None)
    set_expr
    Intersect.set_expr(*tokens)
    update_input_tokens
    Intersect.update_input_tokens(*tokens)
    update_name
    Intersect.update_name(tokens, stack)
    __init__(s,context=None)
        Initialize self. See help(type(self)) for accurate signature.
```


## Attributes

| get_n_args |
| :--- |
| name |
| node_id |
| pred |

get_n_args

Intersect.get_n_args
name

Intersect. name
node_id

Intersect.node_id
pred

Intersect.pred

## Operator

class Operator $(s$, context $=$ None $)$

Methods

| _init__ | Initialize self. |
| :--- | :--- |
| ast |  |
| compile |  |
| match |  |
| process |  |
| set_expr |  |
| update_input_tokens |  |
| update_name |  |

__init

Operator.__init__ $(s$, context $=$ None $)$
Initialize self. See help(type(self)) for accurate signature.
ast

Operator.ast (tokens, stack, builder)
compile

Operator.compile()
match

Operator.match $(s)$
process

Operator.process (match, context=None)
set_expr

Operator.set_expr(*tokens)
update_input_tokens

Operator.update_input_tokens (*tokens)
update_name

Operator.update_name (tokens, stack)
_init__( $s$, context $=$ None) Initialize self. See help(type(self)) for accurate signature.

## Attributes

| get_n_args |
| :--- |
| name |
| node_id |
| pred |

get_n_args

Operator.get_n_args
name

Operator.name
node_id

Operator.node_id
pred

Operator.pred

## OperatorToken

class OperatorToken (s, context=None)

## Methods

| _init__ | Initialize self. |
| :--- | :--- |
| ast |  |
| compile |  |
| match |  |
| process |  |
| set_expr |  |
| update_input_tokens |  |
| update_name |  |

init $\qquad$

OperatorToken.__init__( $s$, context $=$ None $)$ Initialize self. See help(type(self)) for accurate signature.
ast

OperatorToken.ast (tokens, stack, builder)
compile

OperatorToken.compile()
match

OperatorToken.match ( $s$ )
process

OperatorToken.process (match, context=None)

```
set_expr
OperatorToken.set_expr(*tokens)
update_input_tokens
OperatorToken.update_input_tokens(*tokens)
update_name
OperatorToken.update_name (tokens, stack)
__init__(s, context=None)
        Initialize self. See help(type(self)) for accurate signature.
```


## Attributes

get_n_args
name
node_id
pred
get_n_args
OperatorToken.get_n_args
name
OperatorToken.name
node_id
OperatorToken.node_id
pred
OperatorToken.pred
Separator
class Separator ( $s$, context=None)

## Methods

| init__ Initialize self. |
| :---: |
| ast |
| compile |
| match |
| process |
| set_expr |
| update_input_tokens |
| update_name |
| __init__ |
| Separator. $\qquad$ init $\qquad$ ( $s$, context $=$ None) Initialize self. See help(type(self)) for accurate signature. |
| ast |
| Separator.ast (tokens, stack, builder) |
| compile |
| Separator.compile() |
| match |
| Separator.match (s) |
| process |
| Separator.process ( match, context=None) |
| set_expr |
| Separator.set_expr (*tokens) |
| update_input_tokens |
| Separator.update_input_tokens (*tokens) |
| update_name |
| Separator.update_name (tokens, stack) |
| $\qquad$ init $\qquad$ ( $s$, context $=$ None) <br> Initialize self. See help(type(self)) for accurate signature. |

## Attributes

| get_n_args |
| :--- |
| name |
| node_id |
| pred |

get_n_args

Separator.get_n_args
name

Separator. name
node_id

Separator.node_id
pred

Separator.pred
parenthesis

It provides Parenthesis class.

Classes

Parenthesis

Parenthesis
class Parenthesis ( $s$, context=None)

Methods

| _init__ | Initialize self. |
| :--- | :--- |
| ast |  |
| match |  |
| process |  |
| set_expr |  |
| update_input_tokens |  |

```
__init
Parenthesis.__init__ (s, context=None)
    Initialize self. See help(type(self)) for accurate signature.
ast
Parenthesis.ast(tokens, stack, builder)
match
Parenthesis.match(s)
process
Parenthesis.process(match, context=None)
set_expr
Parenthesis.set_expr(*tokens)
update_input_tokens
Parenthesis.update_input_tokens(*tokens)
__init__(s, context=None)
    Initialize self. See help(type(self)) for accurate signature.
```


## Attributes

| n_args |
| :--- |
| name |
| node_id |
| opens |

n_args

Parenthesis.n_args $=0$
name

Parenthesis.name

```
node_id
Parenthesis.node_id
opens
Parenthesis.opens = {')': '('}
```


## Classes

```
Token
```


## Token

```
class Token (s, context=None)
```

Methods

| _init__ | Initialize self. |
| :--- | :--- |
| ast |  |
| match |  |
| process |  |
| set_expr |  |
| update_input_tokens |  |

__init

Token.__init__(s, context=None) Initialize self. See help(type(self)) for accurate signature.
ast

Token.ast (tokens, stack, builder)
match

Token.match (s)
process
Token.process (match, context=None)
set_expr

Token.set_expr (*tokens)
update_input_tokens
Token. update_input_tokens (*tokens)
__init__(s, context=None)
Initialize self. See help(type(self)) for accurate signature.

Attributes

| name |
| :--- | :--- |
| node_id |

name

Token.name
node_id

Token.node_id

### 2.1.7.5 functions

It provides functions implementations to compile the Excel functions.
Sub-Modules:

| eng | Python equivalents of engineering Excel functions. |
| :--- | :--- |
| financial | Python equivalents of financial Excel functions. |
| info | Python equivalents of information Excel functions. |
| logic | Python equivalents of logical Excel functions. |
| look | Python equivalents of lookup and reference Excel func- <br> tions. |
| math | Python equivalents of math and trigonometry Excel <br> functions. |
| operators | Python equivalents of Excel operators. |
| stat | Python equivalents of statistical Excel functions. |
| text | Python equivalents of text Excel functions. |

## eng

Python equivalents of engineering Excel functions.
financial

Python equivalents of financial Excel functions.

## Functions

$\qquad$
xirr
xirr ( $x$, guess=0.1)
info

Python equivalents of information Excel functions.

## Functions

| iserr |
| :--- |
| iserror |

iserr
iserr (val)
iserror
iserror (val)

Classes
IsErrArray
IsErrorArray

## IsErrArray

class IsErrArray

## Methods

| all | Returns True if all elements evaluate to True. |
| :--- | :--- |
| any | Returns True if any of the elements of $a$ evaluate to <br> True. |
| argmax | Return indices of the maximum values along the <br> given axis. |
| argmin | Return indices of the minimum values along the <br> given axis of $a$. |
| argpartition | Returns the indices that would partition this array. |
|  | Continued on next page |

Table 47 - continued from previous page

| argsort | Returns the indices that would sort this array. |
| :---: | :---: |
| astype | Copy of the array, cast to a specified type. |
| byteswap | Swap the bytes of the array elements |
| choose | Use an index array to construct a new array from a set of choices. |
| clip | Return an array whose values are limited to [min, max]. |
| collapse |  |
| compress | Return selected slices of this array along given axis. |
| conj | Complex-conjugate all elements. |
| conjugate | Return the complex conjugate, element-wise. |
| copy | Return a copy of the array. |
| cumprod | Return the cumulative product of the elements along the given axis. |
| cumsum | Return the cumulative sum of the elements along the given axis. |
| diagonal | Return specified diagonals. |
| dot | Dot product of two arrays. |
| dump | Dump a pickle of the array to the specified file. |
| dumps | Returns the pickle of the array as a string. |
| fill | Fill the array with a scalar value. |
| flatten | Return a copy of the array collapsed into one dimension. |
| getfield | Returns a field of the given array as a certain type. |
| item | Copy an element of an array to a standard Python scalar and return it. |
| itemset | Insert scalar into an array (scalar is cast to array's dtype, if possible) |
| max | Return the maximum along a given axis. |
| mean | Returns the average of the array elements along given axis. |
| min | Return the minimum along a given axis. |
| newbyteorder | Return the array with the same data viewed with a different byte order. |
| nonzero | Return the indices of the elements that are non-zero. |
| partition | Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. |
| prod | Return the product of the array elements over the given axis |
| ptp | Peak to peak (maximum - minimum) value along a given axis. |
| put | Set a.flat [n] = values [n] for all $n$ in indices. |
| ravel | Return a flattened array. |
| repeat | Repeat elements of an array. |
| reshape | Returns an array containing the same data with a new shape. |
| resize | Change shape and size of array in-place. |
| round | Return $a$ with each element rounded to the given number of decimals. |

Continued on next page

Table 47 - continued from previous page

| searchsorted | Find indices where elements of v should be inserted <br> in a to maintain order. |
| :--- | :--- |
| setfield | Put a value into a specified place in a field defined by <br> a data-type. |
| setflags | Set array flags WRITEABLE, ALIGNED, and UP- <br> DATEIFCOPY, respectively. |
| sort | Sort an array, in-place. <br> Remove single-dimensional entries from the shape <br> of $a$. |
| squeeze | Returns the standard deviation of the array elements <br> along given axis. |
| std | Return the sum of the array elements over the given <br> axis. |
| sum | Return a view of the array with axisl and $a x i s 2$ in- <br> terchanged. |
| swapaxes | Return an array formed from the elements of $a$ at the <br> given indices. |
| take | Construct Python bytes containing the raw data bytes <br> in the array. |
| tobytes | Write array to a file as text or binary (default). <br> Return the array as a (possibly nested) list. <br> tofileConstruct Python bytes containing the raw data bytes <br> in the array. |
| tolist | Return the sum along diagonals of the array. |
| tostring | Returns a view of the array with axes transposed. <br> Returns the variance of the array elements, along <br> given axis. |
| trace | New view of array with the same data. |
| transpose |  |

## all

IsErrArray.all (axis=None, out=None, keepdims=False)
Returns True if all elements evaluate to True.
Refer to numpy.all for full documentation.
numpy.all : equivalent function
any

IsErrArray . any (axis=None, out=None, keepdims=False)
Returns True if any of the elements of $a$ evaluate to True.
Refer to numpy.any for full documentation.
numpy.any : equivalent function
argmax

IsErrArray.argmax (axis=None, out=None)
Return indices of the maximum values along the given axis.

Refer to numpy.argmax for full documentation.
numpy.argmax : equivalent function
argmin

IsErrArray. argmin (axis=None, out=None)
Return indices of the minimum values along the given axis of $a$.
Refer to numpy.argmin for detailed documentation.
numpy.argmin : equivalent function

## argpartition

IsErrArray. argpartition (kth, axis=-1, kind='introselect', order=None)
Returns the indices that would partition this array.
Refer to numpy.argpartition for full documentation.
New in version 1.8.0.
numpy.argpartition : equivalent function

## argsort

IsErrArray. argsort (axis=-1, kind='quicksort', order=None)
Returns the indices that would sort this array.
Refer to numpy.argsort for full documentation.
numpy.argsort : equivalent function
astype

IsErrArray. astype (dtype, order='K', casting='unsafe', subok=True, copy=True)
Copy of the array, cast to a specified type.
dtype [str or dtype] Typecode or data-type to which the array is cast.
order [ $\{$ ' C ', ' F ', ' A ', ' K '\}, optional] Controls the memory layout order of the result. ' C ' means C order, ' F ' means Fortran order, ' $A$ ' means ' F ' order if all the arrays are Fortran contiguous, ' C ' order otherwise, and ' K ' means as close to the order the array elements appear in memory as possible. Default is ' K '.
casting [\{ 'no', 'equiv', 'safe', 'same_kind', 'unsafe'\}, optional] Controls what kind of data casting may occur. Defaults to 'unsafe' for backwards compatibility.

- 'no' means the data types should not be cast at all.
- 'equiv’ means only byte-order changes are allowed.
- 'safe' means only casts which can preserve values are allowed.
- 'same_kind' means only safe casts or casts within a kind, like float64 to float32, are allowed.
- 'unsafe' means any data conversions may be done.
subok [bool, optional] If True, then sub-classes will be passed-through (default), otherwise the returned array will be forced to be a base-class array.
copy [bool, optional] By default, astype always returns a newly allocated array. If this is set to false, and the dtype, order, and subok requirements are satisfied, the input array is returned instead of a copy.
arr_t [ndarray] Unless copy is False and the other conditions for returning the input array are satisfied (see description for copy input parameter), arr_t is a new array of the same shape as the input array, with dtype, order given by dtype, order.

Starting in NumPy 1.9, astype method now returns an error if the string dtype to cast to is not long enough in 'safe' casting mode to hold the max value of integer/float array that is being casted. Previously the casting was allowed even if the result was truncated.

ComplexWarning When casting from complex to float or int. To avoid this, one should use a.real. astype(t).

```
>>> x = np.array([1, 2, 2.5])
>>> x
array([ 1. , 2. , 2.5])
```

```
>>> x.astype(int)
array([1, 2, 2])
```


## byteswap

## IsErrArray .byteswap (inplace)

Swap the bytes of the array elements
Toggle between low-endian and big-endian data representation by returning a byteswapped array, optionally swapped in-place.
inplace [bool, optional] If True, swap bytes in-place, default is False.
out [ndarray] The byteswapped array. If inplace is True, this is a view to self.

```
>>> A = np.array([1, 256, 8755], dtype=np.int16)
>>> map (hex, A)
['0x1', '0x100', '0x2233']
>>> A.byteswap(True)
array([ 256, 1, 13090], dtype=int16)
>>> map (hex, A)
['0x100', '0x1', '0x3322']
```

Arrays of strings are not swapped

```
>>> A = np.array(['ceg', 'fac'])
>>> A.byteswap()
array(['ceg', 'fac'],
    dtype='|S3')
```

choose

IsErrArray. choose (choices, out=None, mode='raise')
Use an index array to construct a new array from a set of choices.
Refer to numpy.choose for full documentation.
numpy.choose : equivalent function
clip

IsErrArray.clip (min=None, max=None, out=None)
Return an array whose values are limited to [ $\min , \max$ ]. One of max or min must be given.
Refer to numpy.clip for full documentation.
numpy.clip : equivalent function
collapse

IsErrArray.collapse (shape)
compress

IsErrArray. compress (condition, axis=None, out=None)
Return selected slices of this array along given axis.
Refer to numpy.compress for full documentation.
numpy.compress : equivalent function
conj

IsErrArray.conj()
Complex-conjugate all elements.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function
conjugate

IsErrArray.conjugate()
Return the complex conjugate, element-wise.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function

## copy

IsErrArray. copy (order=' ${ }^{\prime}$ ')
Return a copy of the array.
order [\{ 'C', ' F ', ' A ', ' K ' \}, optional] Controls the memory layout of the copy. ' C ' means C -order, ' F ' means F-order, 'A' means ' F ' if $a$ is Fortran contiguous, ' C ' otherwise. ' K ' means match the layout of $a$ as closely as possible. (Note that this function and :func:numpy.copy are very similar, but have different default values for their order= arguments.)
numpy.copy numpy.copyto
>>> $x=n p$.array $([1,2,3],[4,5,6]]$, order='F')

```
>>> y = x.copy()
```

```
>>> x.fill(0)
```

```
>>> x
array([[0, 0, 0],
    [0, 0, 0]])
```

```
>>> y
array([[1, 2, 3],
    [4, 5, 6]])
```

>>> y.flags['C_CONTIGUOUS']
True

## cumprod

IsErrArray. cumprod (axis=None, dtype=None, out=None)
Return the cumulative product of the elements along the given axis.
Refer to numpy.cumprod for full documentation.
numpy.cumprod : equivalent function

## cumsum

IsErrArray. cumsum (axis=None, dtype=None, out=None)
Return the cumulative sum of the elements along the given axis.
Refer to numpy.cumsum for full documentation.
numpy.cumsum : equivalent function

## diagonal

## IsErrArray.diagonal (offset=0, axis $1=0$, axis $2=1$ )

Return specified diagonals. In NumPy 1.9 the returned array is a read-only view instead of a copy as in previous NumPy versions. In a future version the read-only restriction will be removed.

Refer to numpy. diagonal () for full documentation.
numpy.diagonal : equivalent function
dot

IsErrArray.dot ( $b$, out=None)
Dot product of two arrays.
Refer to numpy.dot for full documentation.
numpy.dot : equivalent function

```
>>> a = np.eye(2)
>>> b = np.ones((2, 2)) * 2
>>> a.dot(b)
array([[ 2., 2.],
    [ 2., 2.]])
```

This array method can be conveniently chained:

```
>>> a.dot(b).dot (b)
array([[ 8., 8.],
    [ 8., 8.]])
```

dump

IsErrArray.dump (file)
Dump a pickle of the array to the specified file. The array can be read back with pickle.load or numpy.load.
file [str] A string naming the dump file.
dumps

IsErrArray.dumps()
Returns the pickle of the array as a string. pickle.loads or numpy.loads will convert the string back to an array.
None
fill

IsErrArray.fill (value)
Fill the array with a scalar value.
value [scalar] All elements of $a$ will be assigned this value.

```
>>> a = np.array([1, 2])
>>> a.fill(0)
>>> a
array([0, 0])
>>> a = np.empty(2)
>>> a.fill(1)
```

```
>>> a
array([ 1., 1.])
```


## flatten

IsErrArray.flatten (order='C')
Return a copy of the array collapsed into one dimension.
order [\{ 'C', ' F ', ' A ', ' K '\}, optional] ' C ' means to flatten in row-major ( C -style) order. ' F ' means to flatten in column-major (Fortran- style) order. 'A' means to flatten in column-major order if $a$ is Fortran contiguous in memory, row-major order otherwise. ' K ' means to flatten $a$ in the order the elements occur in memory. The default is ' C '.
$\mathbf{y}$ [ndarray] A copy of the input array, flattened to one dimension.
ravel : Return a flattened array. flat: A 1-D flat iterator over the array.

```
>>> a = np.array([[1,2], [3,4]])
>>> a.flatten()
array([1, 2, 3, 4])
>>> a.flatten('F')
array([1, 3, 2, 4])
```


## getfield

IsErrArray.getfield (dtype, offset=0)
Returns a field of the given array as a certain type.
A field is a view of the array data with a given data-type. The values in the view are determined by the given type and the offset into the current array in bytes. The offset needs to be such that the view dtype fits in the array dtype; for example an array of dtype complex 128 has 16-byte elements. If taking a view with a 32-bit integer ( 4 bytes), the offset needs to be between 0 and 12 bytes.
dtype [str or dtype] The data type of the view. The dtype size of the view can not be larger than that of the array itself.
offset [int] Number of bytes to skip before beginning the element view.

```
>>> x = np.diag([1.+1.j]*2)
>>> x[1, 1] = 2 + 4.j
>>> x
array([[ 1.+1.j, 0.+0.j],
    [ 0.+0.j, 2.+4.j]])
>>> x.getfield(np.float64)
array([[ 1., 0.],
    [ 0., 2.]])
```

By choosing an offset of 8 bytes we can select the complex part of the array for our view:

```
>>> x.getfield(np.float64, offset=8)
array([[ 1., 0.],
    [0., 4.]])
```

item

IsErrArray.item (*args)
Copy an element of an array to a standard Python scalar and return it.
*args : Arguments (variable number and type)

- none: in this case, the method only works for arrays with one element (a.size $==1$ ), which element is copied into a standard Python scalar object and returned.
- int_type: this argument is interpreted as a flat index into the array, specifying which element to copy and return.
- tuple of int_types: functions as does a single int_type argument, except that the argument is interpreted as an nd-index into the array.
$\mathbf{z}$ [Standard Python scalar object] A copy of the specified element of the array as a suitable Python scalar
When the data type of $a$ is longdouble or clongdouble, item() returns a scalar array object because there is no available Python scalar that would not lose information. Void arrays return a buffer object for item(), unless fields are defined, in which case a tuple is returned.
item is very similar to a[args], except, instead of an array scalar, a standard Python scalar is returned. This can be useful for speeding up access to elements of the array and doing arithmetic on elements of the array using Python's optimized math.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.item(3)
2
>>> x.item(7)
5
>>> x.item((0, 1))
1
>>> x.item((2, 2))
3
```


## itemset

IsErrArray.itemset (*args)
Insert scalar into an array (scalar is cast to array's dtype, if possible)
There must be at least 1 argument, and define the last argument as item. Then, a.itemset (*args) is equivalent to but faster than a [args] = item. The item should be a scalar value and args must select a single item in the array $a$.
\*args [Arguments] If one argument: a scalar, only used in case $a$ is of size 1 . If two arguments: the last argument is the value to be set and must be a scalar, the first argument specifies a single array element location. It is either an int or a tuple.

Compared to indexing syntax, itemset provides some speed increase for placing a scalar into a particular location in an ndarray, if you must do this. However, generally this is discouraged: among other problems, it complicates the appearance of the code. Also, when using itemset (and item) inside a loop, be sure to assign the methods to a local variable to avoid the attribute look-up at each loop iteration.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.itemset (4, 0)
>>> x.itemset ((2, 2), 9)
>>> x
array([[3, 1, 7],
    [2, 0, 3],
    [8, 5, 9]])
```

max

IsErrArray.max (axis=None, out=None)
Return the maximum along a given axis.
Refer to numpy.amax for full documentation.
numpy.amax : equivalent function
mean

IsErrArray .mean (axis=None, dtype=None, out=None, keepdims=False)
Returns the average of the array elements along given axis.
Refer to numpy.mean for full documentation.
numpy.mean : equivalent function
min

IsErrArray.min (axis=None, out=None, keepdims=False)
Return the minimum along a given axis.
Refer to numpy.amin for full documentation.
numpy.amin : equivalent function
newbyteorder

IsErrArray. newbyteorder (new_order='S')
Return the array with the same data viewed with a different byte order.
Equivalent to:

```
arr.view(arr.dtype.newbytorder(new_order))
```

Changes are also made in all fields and sub-arrays of the array data type.
new_order [string, optional] Byte order to force; a value from the byte order specifications below. new_order codes can be any of:

- 'S' - swap dtype from current to opposite endian
- $\{$ ' $<$ ', 'L' $\}$ - little endian
- \{'>', 'B'\} - big endian
- $\{$ ' $=$ ', ' N ' $\}$ - native order
- \{' 1 ', ' I ' $\}$ - ignore (no change to byte order)

The default value ('S') results in swapping the current byte order. The code does a case-insensitive check on the first letter of new_order for the alternatives above. For example, any of ' B ' or ' $b$ ' or 'biggish' are valid to specify big-endian.
new_arr [array] New array object with the dtype reflecting given change to the byte order.

## nonzero

## IsErrArray.nonzero()

Return the indices of the elements that are non-zero.
Refer to numpy.nonzero for full documentation.
numpy.nonzero : equivalent function

## partition

IsErrArray.partition(kth, axis=-l, kind='introselect',order=None)
Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. All elements smaller than the kth element are moved before this element and all equal or greater are moved behind it. The ordering of the elements in the two partitions is undefined.

New in version 1.8.0.
kth [int or sequence of ints] Element index to partition by. The kth element value will be in its final sorted position and all smaller elements will be moved before it and all equal or greater elements behind it. The order all elements in the partitions is undefined. If provided with a sequence of kth it will partition all elements indexed by kth of them into their sorted position at once.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'introselect'\}, optional] Selection algorithm. Default is 'introselect'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.partition : Return a parititioned copy of an array. argpartition : Indirect partition. sort : Full sort.
See np.partition for notes on the different algorithms.

```
>>> a = np.array([3, 4, 2, 1])
>>> a.partition(a, 3)
>>> a
array([2, 1, 3, 4])
```

```
>>> a.partition((1, 3))
array([1, 2, 3, 4])
```


## prod

IsErrArray.prod (axis=None, dtype=None, out=None, keepdims $=$ False )
Return the product of the array elements over the given axis
Refer to numpy.prod for full documentation.
numpy.prod : equivalent function
ptp

IsErrArray.ptp (axis=None, out=None)
Peak to peak (maximum - minimum) value along a given axis.
Refer to numpy.ptp for full documentation.
numpy.ptp : equivalent function
put

IsErrArray.put (indices, values, mode='raise')
Set a.flat $[\mathrm{n}]=$ values $[\mathrm{n}]$ for all $n$ in indices.
Refer to numpy.put for full documentation.
numpy.put : equivalent function
ravel

IsErrArray. ravel ([order])
Return a flattened array.
Refer to numpy.ravel for full documentation.
numpy.ravel : equivalent function
ndarray.flat : a flat iterator on the array.
repeat

IsErrArray.repeat (repeats, axis=None)
Repeat elements of an array.
Refer to numpy.repeat for full documentation.
numpy.repeat : equivalent function
reshape

IsErrArray.reshape (shape, order='C')
Returns an array containing the same data with a new shape.
Refer to numpy.reshape for full documentation.
numpy.reshape : equivalent function
resize

IsErrArray. resize (new_shape, refcheck=True)
Change shape and size of array in-place.
new_shape [tuple of ints, or $n$ ints] Shape of resized array.
refcheck [bool, optional] If False, reference count will not be checked. Default is True.
None
ValueError If $a$ does not own its own data or references or views to it exist, and the data memory must be changed.
SystemError If the order keyword argument is specified. This behaviour is a bug in NumPy.
resize : Return a new array with the specified shape.
This reallocates space for the data area if necessary.
Only contiguous arrays (data elements consecutive in memory) can be resized.
The purpose of the reference count check is to make sure you do not use this array as a buffer for another Python object and then reallocate the memory. However, reference counts can increase in other ways so if you are sure that you have not shared the memory for this array with another Python object, then you may safely set refcheck to False.
Shrinking an array: array is flattened (in the order that the data are stored in memory), resized, and reshaped:

```
>>>}a=np.\operatorname{array([[0, 1], [2, 3]], order='C')
>>> a.resize((2, 1))
>>> a
array([[0],
    [1]])
```

```
>>> a = np.array([[0, 1], [2, 3]], order='F')
>>> a.resize((2, 1))
>>> a
array([[0],
    [2]])
```

Enlarging an array: as above, but missing entries are filled with zeros:

```
>>> b = np.array([[0, 1], [2, 3]])
>>> b.resize(2, 3) # new_shape parameter doesn't have to be a tuple
>>> b
array([[0, 1, 2],
    [3, 0, 0]])
```

Referencing an array prevents resizing...

```
>>> c = a
>>> a.resize((1, 1))
Traceback (most recent call last):
ValueError: cannot resize an array that has been referenced ...
```

Unless refcheck is False:

```
>>> a.resize((1, 1), refcheck=False)
>>> a
array([[0]])
>>> c
array([[0]])
```


## round

## IsErrArray.round (decimals=0, out=None)

Return $a$ with each element rounded to the given number of decimals.
Refer to numpy.around for full documentation.
numpy.around : equivalent function

## searchsorted

IsErrArray. searchsorted ( $v$, side='left', sorter=None)
Find indices where elements of v should be inserted in a to maintain order.
For full documentation, see numpy.searchsorted
numpy.searchsorted : equivalent function

## setfield

IsErrArray.setfield (val, dtype, offset=0)
Put a value into a specified place in a field defined by a data-type.
Place val into $a$ 's field defined by dtype and beginning offset bytes into the field.
val [object] Value to be placed in field.
dtype [dtype object] Data-type of the field in which to place val.
offset [int, optional] The number of bytes into the field at which to place val.
None
getfield

```
>>> x = np.eye(3)
>>> x.getfield(np.float64)
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
>>> x.setfield(3, np.int32)
>>> x.getfield(np.int32)
array([[3, 3, 3],
    [3, 3, 3],
    [3, 3, 3]])
>>> x
array([[ 1.00000000e+000, 1.48219694e-323, 1.48219694e-323],
    [ 1.48219694e-323, 1.00000000e+000, 1.48219694e-323],
    [ 1.48219694e-323, 1.48219694e-323, 1.00000000e+000]])
>>> x.setfield(np.eye(3), np.int32)
```

```
>>> x
```

array([[ 1., 0., 0.],
$\left[\begin{array}{lll}10 ., & 1 ., & 0 .],\end{array}\right.$
$\left[\begin{array}{lll}{[0 .,} & 0 ., & 1 .]\end{array}\right)$

## setflags

IsErrArray.setflags (write=None, align=None, uic=None)
Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively.
These Boolean-valued flags affect how numpy interprets the memory area used by $a$ (see Notes below). The ALIGNED flag can only be set to True if the data is actually aligned according to the type. The UPDATEIFCOPY flag can never be set to True. The flag WRITEABLE can only be set to True if the array owns its own memory, or the ultimate owner of the memory exposes a writeable buffer interface, or is a string. (The exception for string is made so that unpickling can be done without copying memory.)
write [bool, optional] Describes whether or not $a$ can be written to.
align [bool, optional] Describes whether or not $a$ is aligned properly for its type.
uic [bool, optional] Describes whether or not $a$ is a copy of another "base" array.
Array flags provide information about how the memory area used for the array is to be interpreted. There are 6 Boolean flags in use, only three of which can be changed by the user: UPDATEIFCOPY, WRITEABLE, and ALIGNED.

WRITEABLE (W) the data area can be written to;
ALIGNED (A) the data and strides are aligned appropriately for the hardware (as determined by the compiler);

UPDATEIFCOPY (U) this array is a copy of some other array (referenced by .base). When this array is deallocated, the base array will be updated with the contents of this array.

All flags can be accessed using their first (upper case) letter as well as the full name.

```
>>> y
array([[3, 1, 7],
    [2, 0, 0],
    [8, 5, 9]])
>>> y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : True
    ALIGNED : True
    UPDATEIFCOPY : False
>>> Y.setflags(write=0, align=0)
>>> Y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : False
    ALIGNED : False
    UPDATEIFCOPY : False
>>> y.setflags(uic=1)
Traceback (most recent call last):
```

```
File "<stdin>", line 1, in <module>
ValueError: cannot set UPDATEIFCOPY flag to True
```


## sort

IsErrArray.sort (axis=-1, kind='quicksort', order=None)
Sort an array, in-place.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'quicksort', 'mergesort', 'heapsort'\}, optional] Sorting algorithm. Default is 'quicksort'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.sort : Return a sorted copy of an array. argsort : Indirect sort. lexsort : Indirect stable sort on multiple keys. searchsorted : Find elements in sorted array. partition: Partial sort.

See sort for notes on the different sorting algorithms.

```
>>> a = np.array([[1,4], [3,1]])
>>> a.sort(axis=1)
>>> a
array([[1, 4],
    [1, 3]])
>>> a.sort(axis=0)
>>> a
array([[1, 3],
    [1, 4]])
```

Use the order keyword to specify a field to use when sorting a structured array:

```
>>> a = np.array([('a', 2), ('c', 1)], dtype=[('x', 'S1'), ('y', int)])
>>> a.sort(order='y')
>>> a
array([('c', 1), ('a', 2)],
    dtype=[('x', '|S1'), ('y', '<i4')])
```


## squeeze

IsErrArray.squeeze (axis=None)
Remove single-dimensional entries from the shape of $a$.
Refer to numpy.squeeze for full documentation.
numpy.squeeze : equivalent function
std

IsErrArray.std (axis=None, dtype=None, out=None, ddof=0, keepdims=False)
Returns the standard deviation of the array elements along given axis.

Refer to numpy.std for full documentation.
numpy.std : equivalent function

## sum

IsErrArray. sum (axis=None, dtype $=$ None, out $=$ None, keepdims $=$ False )
Return the sum of the array elements over the given axis.
Refer to numpy.sum for full documentation.
numpy.sum : equivalent function

## swapaxes

IsErrArray.swapaxes (axis1, axis2)
Return a view of the array with axis 1 and axis 2 interchanged.
Refer to numpy.swapaxes for full documentation.
numpy.swapaxes: equivalent function

## take

IsErrArray.take (indices, axis=None, out=None, mode='raise')
Return an array formed from the elements of $a$ at the given indices.
Refer to numpy.take for full documentation.
numpy.take : equivalent function

## tobytes

IsErrArray.tobytes (order='C')
Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.
New in version 1.9.0.
order [ $\{$ ' $C$ ', ' $F$ ', None $\}$, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01\x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01\x00\x00\x00\x03\x00\x00\x00'
```


## tofile

IsErrArray.tofile (fid, sep="", format="\%s")
Write array to a file as text or binary (default).
Data is always written in ' C ' order, independent of the order of $a$. The data produced by this method can be recovered using the function fromfile().
fid [file or str] An open file object, or a string containing a filename.
sep [str] Separator between array items for text output. If "" (empty), a binary file is written, equivalent to file.write (a.tobytes()).
format [str] Format string for text file output. Each entry in the array is formatted to text by first converting it to the closest Python type, and then using "format" \% item.

This is a convenience function for quick storage of array data. Information on endianness and precision is lost, so this method is not a good choice for files intended to archive data or transport data between machines with different endianness. Some of these problems can be overcome by outputting the data as text files, at the expense of speed and file size.

## tolist

IsErrArray.tolist()
Return the array as a (possibly nested) list.
Return a copy of the array data as a (nested) Python list. Data items are converted to the nearest compatible Python type.
none
$\mathbf{y}$ [list] The possibly nested list of array elements.
The array may be recreated, $a=n p$.array(a.tolist()).

```
>>> a = np.array([1, 2])
>>> a.tolist()
[1, 2]
>>> a = np.array([[1, 2], [3, 4]])
>>> list(a)
[array([1, 2]), array([3, 4])]
>>> a.tolist()
[[1, 2], [3, 4]]
```


## tostring

## IsErrArray.tostring (order='C')

Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.

This function is a compatibility alias for tobytes. Despite its name it returns bytes not strings.
order [ $\{$ ' C ', ' F ', None \}, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01\x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01 \x00\x00\x00\x03\x00\x00\x00'
```


## trace

## IsErrArray.trace (offset=0, axis $1=0$, axis $2=1$, dtype=None, out=None)

Return the sum along diagonals of the array.
Refer to numpy.trace for full documentation.
numpy.trace : equivalent function

## transpose

IsErrArray.transpose (*axes)
Returns a view of the array with axes transposed.
For a 1-D array, this has no effect. (To change between column and row vectors, first cast the 1-D array into a matrix object.) For a 2-D array, this is the usual matrix transpose. For an n-D array, if axes are given, their order indicates how the axes are permuted (see Examples). If axes are not provided and a.shape $=(i[0], i[1], \ldots i[n-2], i[n-1])$, then a.transpose(). shape $=$ (i[n-1], i[n-2], ... i[1], i[0]).
axes : None, tuple of ints, or $n$ ints

- None or no argument: reverses the order of the axes.
- tuple of ints: $i$ in the $j$-th place in the tuple means $a$ 's $i$-th axis becomes a.transpose()'s $j$-th axis.
- $n$ ints: same as an $n$-tuple of the same ints (this form is intended simply as a "convenience" alternative to the tuple form)
out [ndarray] View of $a$, with axes suitably permuted.
ndarray.T : Array property returning the array transposed.

```
>>> a = np.array([[1, 2], [3, 4]])
>>> a
array([[1, 2],
    [3, 4]])
>>> a.transpose()
array([[1, 3],
        [2, 4]])
>>> a.transpose((1, 0))
array([[1, 3],
            [2, 4]])
>>> a.transpose(1, 0)
array([[1, 3],
    [2, 4]])
```


## var

IsErrArray. $\operatorname{var}($ axis=None, dtype=None, out=None, ddof=0, keepdims $=$ False $)$
Returns the variance of the array elements, along given axis.
Refer to numpy.var for full documentation.
numpy.var : equivalent function
view

IsErrArray.view (dtype=None, type=None)
New view of array with the same data.
dtype [data-type or ndarray sub-class, optional] Data-type descriptor of the returned view, e.g., float 32 or int16. The default, None, results in the view having the same data-type as $a$. This argument can also be specified as an ndarray sub-class, which then specifies the type of the returned object (this is equivalent to setting the type parameter).
type [Python type, optional] Type of the returned view, e.g., ndarray or matrix. Again, the default None results in type preservation.
a.view () is used two different ways:
a.view (some_dtype) or a.view (dtype=some_dtype) constructs a view of the array's memory with a different data-type. This can cause a reinterpretation of the bytes of memory.
a.view (ndarray_subclass) or a.view(type=ndarray_subclass) just returns an instance of ndarray_subclass that looks at the same array (same shape, dtype, etc.) This does not cause a reinterpretation of the memory.

For a.view (some_dtype), if some_dtype has a different number of bytes per entry than the previous dtype (for example, converting a regular array to a structured array), then the behavior of the view cannot be predicted just from the superficial appearance of a (shown by print (a)). It also depends on exactly how a is stored in memory. Therefore if a is C-ordered versus fortran-ordered, versus defined as a slice or transpose, etc., the view may give different results.

```
>>> x = np.array([(1, 2)], dtype=[('a', np.int8), ('b', np.int8)])
```

Viewing array data using a different type and dtype:

```
>>> y = x.view(dtype=np.int16, type=np.matrix)
>>> y
matrix([[513]], dtype=int16)
>>> print(type(y))
<class 'numpy.matrixlib.defmatrix.matrix'>
```

Creating a view on a structured array so it can be used in calculations

```
>>> x = np.array([(1, 2), (3,4)], dtype=[('a', np.int8), ('b', np.int8)])
>>> xv = x.view(dtype=np.int8).reshape(-1,2)
>>> xv
array([[1, 2],
    [3, 4]], dtype=int8)
>>> xv.mean(0)
array([ 2., 3.])
```

Making changes to the view changes the underlying array

```
>>> xv[0,1] = 20
>>> print(x)
[(1, 20) (3, 4)]
```

Using a view to convert an array to a recarray:

```
>>> z = x.view(np.recarray)
>>> z.a
array([1], dtype=int8)
```

Views share data:

```
>>> x[0] = (9, 10)
>>> z[0]
(9, 10)
```

Views that change the dtype size (bytes per entry) should normally be avoided on arrays defined by slices, transposes, fortran-ordering, etc.:

```
>>> x = np.array([[1,2,3],[4,5,6]], dtype=np.int16)
>>> y = x[:, 0:2]
>>> y
array([[1, 2],
    [4, 5]], dtype=int16)
>>> y.view(dtype=[('width', np.int16), ('length', np.int16)])
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: new type not compatible with array.
>>> z = y.copy()
>>> z.view(dtype=[('width', np.int16), ('length', np.int16)])
array([[(1, 2)],
    [(4, 5)]], dtype=[('width', '<i2'), ('length', '<i2')])
```

init__()
Initialize self. See help(type(self)) for accurate signature.

## Attributes

| T | Same as self.transpose(), except that self is returned <br> if self.ndim < 2. |
| :--- | :--- |
| base | Base object if memory is from some other object. |
| ctypes | An object to simplify the interaction of the array with <br> the ctypes module. |
| data | Python buffer object pointing to the start of the ar- <br> ray's data. |
| dtype | Data-type of the array's elements. |
| flags | Information about the memory layout of the array. |
| flat | A 1-D iterator over the array. |
| imag | The imaginary part of the array. |
| itemsize | Length of one array element in bytes. |
| nbytes | Total bytes consumed by the elements of the array. |
| ndim | Number of array dimensions. |
| real | The real part of the array. |

Table 48 - continued from previous page

| shape | Tuple of array dimensions. |
| :--- | :--- |
| size | Number of elements in the array. |
| strides | Tuple of bytes to step in each dimension when <br> traversing an array. |

T

IsErrArray.T
Same as self.transpose(), except that self is returned if self.ndim $<2$.

```
>>> x = np.array([[1.,2.],[3.,4.]])
>>> x
array([[ 1., 2.],
                    [ 3., 4.]])
>>> x.T
array([[ 1., 3.],
    [ 2., 4.]])
>>> x = np.array([1.,2.,3.,4.])
>>> X
array([ 1., 2., 3., 4.])
>>> x.T
array([ 1., 2., 3., 4.])
```

base

## IsErrArray.base

Base object if memory is from some other object.
The base of an array that owns its memory is None:

```
>>> x = np.array([1,2,3,4])
>>> x.base is None
True
```

Slicing creates a view, whose memory is shared with x :

```
>>> y = x[2:]
>>> y.base is x
True
```

ctypes

IsErrArray. ctypes
An object to simplify the interaction of the array with the ctypes module.
This attribute creates an object that makes it easier to use arrays when calling shared libraries with the ctypes module. The returned object has, among others, data, shape, and strides attributes (see Notes below) which themselves return ctypes objects that can be used as arguments to a shared library.

None
c [Python object] Possessing attributes data, shape, strides, etc.

## numpy.ctypeslib

Below are the public attributes of this object which were documented in "Guide to NumPy" (we have omitted undocumented public attributes, as well as documented private attributes):

- data: A pointer to the memory area of the array as a Python integer. This memory area may contain data that is not aligned, or not in correct byte-order. The memory area may not even be writeable. The array flags and data-type of this array should be respected when passing this attribute to arbitrary C-code to avoid trouble that can include Python crashing. User Beware! The value of this attribute is exactly the same as self._array_interface_['data'][0].
- shape (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the C-integer corresponding to dtype(' p ') on this platform. This base-type could be c_int, c _long, or c_longlong depending on the platform. The c_intp type is defined accordingly in numpy.ctypeslib. The ctypes array contains the shape of the underlying array.
- strides (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the same as for the shape attribute. This ctypes array contains the strides information from the underlying array. This strides information is important for showing how many bytes must be jumped to get to the next element in the array.
- data_as(obj): Return the data pointer cast to a particular c-types object. For example, calling self._as_parameter_ is equivalent to self.data_as(ctypes.c_void_p). Perhaps you want to use the data as a pointer to a ctypes array of floating-point data: self.data_as(ctypes.POINTER(ctypes.c_double)).
- shape_as(obj): Return the shape tuple as an array of some other c-types type. For example: self.shape_as(ctypes.c_short).
- strides_as(obj): Return the strides tuple as an array of some other c-types type. For example: self.strides_as(ctypes.c_longlong).
Be careful using the ctypes attribute - especially on temporary arrays or arrays constructed on the fly. For example, calling ( $a+b$ ). ctypes.data_as (ctypes.c_void_p) returns a pointer to memory that is invalid because the array created as $(\mathrm{a}+\mathrm{b})$ is deallocated before the next Python statement. You can avoid this problem using either $c=a+b$ or $c t=(a+b)$. ctypes. In the latter case, ct will hold a reference to the array until ct is deleted or re-assigned.

If the ctypes module is not available, then the ctypes attribute of array objects still returns something useful, but ctypes objects are not returned and errors may be raised instead. In particular, the object will still have the as parameter attribute which will return an integer equal to the data attribute.

```
>>> import ctypes
>>> x
array([[0, 1],
    [2, 3]])
>>> x.ctypes.data
30439712
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long))
<ctypes.LP_c_long object at 0x01F01300>
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long)).contents
c_long(0)
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_longlong)).contents
C_longlong(4294967296L)
>>> x.ctypes.shape
<numpy.core._internal.c_long_Array_2 object at 0x01FFD580>
>>> x.ctypes.shape_as(ctypes.c_long)
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides
```

(continues on next page)

```
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides_as(ctypes.c_longlong)
<numpy.core._internal.c_longlong_Array_2 object at 0x01F01300>
```


## data

## IsErrArray.data

Python buffer object pointing to the start of the array's data.

## dtype

IsErrArray.dtype
Data-type of the array's elements.
None
d : numpy dtype object
numpy.dtype

```
>>> x
array([[0, 1],
        [2, 3]])
>>> x.dtype
dtype('int32')
>>> type(x.dtype)
<type 'numpy.dtype'>
```


## flags

## IsErrArray.flags

Information about the memory layout of the array.
C_CONTIGUOUS (C) The data is in a single, C-style contiguous segment.
F_CONTIGUOUS (F) The data is in a single, Fortran-style contiguous segment.
OWNDATA (O) The array owns the memory it uses or borrows it from another object.
WRITEABLE (W) The data area can be written to. Setting this to False locks the data, making it readonly. A view (slice, etc.) inherits WRITEABLE from its base array at creation time, but a view of a writeable array may be subsequently locked while the base array remains writeable. (The opposite is not true, in that a view of a locked array may not be made writeable. However, currently, locking a base object does not lock any views that already reference it, so under that circumstance it is possible to alter the contents of a locked array via a previously created writeable view onto it.) Attempting to change a non-writeable array raises a RuntimeError exception.

ALIGNED (A) The data and all elements are aligned appropriately for the hardware.
UPDATEIFCOPY (U) This array is a copy of some other array. When this array is deallocated, the base array will be updated with the contents of this array.

FNC F_CONTIGUOUS and not C_CONTIGUOUS.
FORC F_CONTIGUOUS or C_CONTIGUOUS (one-segment test).

BEHAVED (B) ALIGNED and WRITEABLE.
CARRAY (CA) BEHAVED and C_CONTIGUOUS.
FARRAY (FA) BEHAVED and F_CONTIGUOUS and not C_CONTIGUOUS.
The flags object can be accessed dictionary-like (as in a.flags ['WRITEABLE']), or by using lowercased attribute names (as in a.flags.writeable). Short flag names are only supported in dictionary access.

Only the UPDATEIFCOPY, WRITEABLE, and ALIGNED flags can be changed by the user, via direct assignment to the attribute or dictionary entry, or by calling ndarray.setflags.
The array flags cannot be set arbitrarily:

- UPDATEIFCOPY can only be set False.
- ALIGNED can only be set True if the data is truly aligned.
- WRITEABLE can only be set True if the array owns its own memory or the ultimate owner of the memory exposes a writeable buffer interface or is a string.

Arrays can be both C-style and Fortran-style contiguous simultaneously. This is clear for 1-dimensional arrays, but can also be true for higher dimensional arrays.

Even for contiguous arrays a stride for a given dimension arr.strides [dim] may be arbitrary if arr.shape [dim] == 1 or the array has no elements. It does not generally hold that self. strides[-1] == self.itemsize for C-style contiguous arrays or self.strides[0] == self.itemsize for Fortran-style contiguous arrays is true.
flat

IsErrArray.flat
A 1-D iterator over the array.
This is a numpy.flatiter instance, which acts similarly to, but is not a subclass of, Python's built-in iterator object.
flatten : Return a copy of the array collapsed into one dimension.
flatiter

```
>>> x = np.arange(1, 7).reshape(2, 3)
>>> x
array([[1, 2, 3],
    [4, 5, 6]])
>>> x.flat[3]
4
>>> x.T
array([[1, 4],
    [2, 5],
    [3, 6]])
>>> x.T.flat[3]
5
>>> type(x.flat)
<type 'numpy.flatiter'>
```

An assignment example:

```
>>> x.flat = 3; x
array([[3, 3, 3],
    [3, 3, 3]])
>>> x.flat[[1,4]] = 1; x
array([[3, 1, 3],
    [3, 1, 3]])
```

imag

## IsErrArray.imag

The imaginary part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.imag
array([ 0. , 0.70710678])
>>> x.imag.dtype
dtype('float64')
```


## itemsize

IsErrArray.itemsize
Length of one array element in bytes.

```
>>> x = np.array([1,2,3], dtype=np.float64)
>>> x.itemsize
8
>>> x = np.array([1,2,3], dtype=np.complex128)
>>> x.itemsize
16
```

nbytes

IsErrArray.nbytes
Total bytes consumed by the elements of the array.
Does not include memory consumed by non-element attributes of the array object.

```
>>> x = np.zeros((3,5,2), dtype=np.complex128)
>>> x.nbytes
480
>>> np.prod(x.shape) * x.itemsize
480
```

ndim

IsErrArray.ndim
Number of array dimensions.

```
>>> x = np.array([1, 2, 3])
>>> x.ndim
1
>>> y = np.zeros((2, 3, 4))
>>> y.ndim
3
```

real

## IsErrArray.real

The real part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.real
array([ 1. , 0.70710678])
>>> x.real.dtype
dtype('float64')
```

numpy.real : equivalent function

## shape

IsErrArray.shape
Tuple of array dimensions.
May be used to "reshape" the array, as long as this would not require a change in the total number of elements

```
>>> x = np.array([1, 2, 3, 4])
>>> x.shape
(4,)
>>> y = np.zeros((2, 3, 4))
>>> y.shape
(2, 3, 4)
>>> y.shape = (3, 8)
>>> y
array([[[ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.]])
>>> y.shape = (3, 6)
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: total size of new array must be unchanged
```

size

IsErrArray.size
Number of elements in the array.
Equivalent to np.prod(a.shape), i.e., the product of the array's dimensions.

```
>>> x = np.zeros((3, 5, 2), dtype=np.complex128)
>>> x.size
30
>>> np.prod(x.shape)
30
```

strides

## IsErrArray.strides

Tuple of bytes to step in each dimension when traversing an array.
The byte offset of element (i[0], i[1], ..., i[n]) in an array $a$ is:

```
offset = sum(np.array(i) * a.strides)
```

A more detailed explanation of strides can be found in the "ndarray.rst" file in the NumPy reference guide.
Imagine an array of 32-bit integers (each 4 bytes):

```
x = np.array([[0, 1, 2, 3, 4],
    [5, 6, 7, 8, 9]], dtype=np.int32)
```

This array is stored in memory as 40 bytes, one after the other (known as a contiguous block of memory). The strides of an array tell us how many bytes we have to skip in memory to move to the next position along a certain axis. For example, we have to skip 4 bytes ( 1 value) to move to the next column, but 20 bytes ( 5 values) to get to the same position in the next row. As such, the strides for the array $x$ will be (20, 4).
numpy.lib.stride_tricks.as_strided

```
>>> y = np.reshape(np.arange(2*3*4), (2, 3,4))
>>> y
array([[[ 0, 1, 2, 3],
    [4, 5, 6, 7],
    [ 8, 9, 10, 11]],
    [[12, 13, 14, 15],
    [16, 17, 18, 19],
    [20, 21, 22, 23]]])
>>> y.strides
(48, 16, 4)
>>> y[1,1,1]
17
>>> offset=sum(y.strides * np.array((1,1,1)))
>>> offset/y.itemsize
17
```

```
>>> x = np.reshape(np.arange(5*6*7*8), (5, 6,7,8)).transpose(2,3,1,0)
>>> x.strides
(32, 4, 224, 1344)
>>> i = np.array([3,5,2,2])
>>> offset = sum(i * x.strides)
>>> x[3,5,2,2]
813
>>> offset / x.itemsize
813
```


## IsErrorArray

class IsErrorArray

## Methods

| all | Returns True if all elements evaluate to True. |
| :---: | :---: |
| any | Returns True if any of the elements of $a$ evaluate to True. |
| argmax | Return indices of the maximum values along the given axis. |
| argmin | Return indices of the minimum values along the given axis of $a$. |
| argpartition | Returns the indices that would partition this array. |
| argsort | Returns the indices that would sort this array. |
| astype | Copy of the array, cast to a specified type. |
| byteswap | Swap the bytes of the array elements |
| choose | Use an index array to construct a new array from a set of choices. |
| clip | Return an array whose values are limited to [min, max]. |
| collapse |  |
| compress | Return selected slices of this array along given axis. |
| conj | Complex-conjugate all elements. |
| conjugate | Return the complex conjugate, element-wise. |
| copy | Return a copy of the array. |
| cumprod | Return the cumulative product of the elements along the given axis. |
| cumsum | Return the cumulative sum of the elements along the given axis. |
| diagonal | Return specified diagonals. |
| dot | Dot product of two arrays. |
| dump | Dump a pickle of the array to the specified file. |
| dumps | Returns the pickle of the array as a string. |
| fill | Fill the array with a scalar value. |
| flatten | Return a copy of the array collapsed into one dimension. |
| getfield | Returns a field of the given array as a certain type. |
| item | Copy an element of an array to a standard Python scalar and return it. |
| itemset | Insert scalar into an array (scalar is cast to array's dtype, if possible) |
| max | Return the maximum along a given axis. |
| mean | Returns the average of the array elements along given axis. |
| min | Return the minimum along a given axis. |
| newbyteorder | Return the array with the same data viewed with a different byte order. |
| nonzero | Return the indices of the elements that are non-zero. |

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| partition | Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. |
| :---: | :---: |
| prod | Return the product of the array elements over the given axis |
| ptp | Peak to peak (maximum - minimum) value along a given axis. |
| put | Set a.flat $[\mathrm{n}]=$ values $[\mathrm{n}]$ for all $n$ in indices. |
| ravel | Return a flattened array. |
| repeat | Repeat elements of an array. |
| reshape | Returns an array containing the same data with a new shape. |
| resize | Change shape and size of array in-place. |
| round | Return $a$ with each element rounded to the given number of decimals. |
| searchsorted | Find indices where elements of v should be inserted in a to maintain order. |
| setfield | Put a value into a specified place in a field defined by a data-type. |
| setflags | Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively. |
| sort | Sort an array, in-place. |
| squeeze | Remove single-dimensional entries from the shape of $a$. |
| std | Returns the standard deviation of the array elements along given axis. |
| sum | Return the sum of the array elements over the given axis. |
| swapaxes | Return a view of the array with axis1 and axis2 interchanged. |
| take | Return an array formed from the elements of $a$ at the given indices. |
| tobytes | Construct Python bytes containing the raw data bytes in the array. |
| tofile | Write array to a file as text or binary (default). |
| tolist | Return the array as a (possibly nested) list. |
| tostring | Construct Python bytes containing the raw data bytes in the array. |
| trace | Return the sum along diagonals of the array. |
| transpose | Returns a view of the array with axes transposed. |
| var | Returns the variance of the array elements, along given axis. |
| view | New view of array with the same data. |

all

IsErrorArray.all (axis=None, out=None, keepdims=False) Returns True if all elements evaluate to True.

Refer to numpy.all for full documentation.
numpy.all : equivalent function

## any

IsErrorArray. any (axis=None, out=None, keepdims=False)
Returns True if any of the elements of $a$ evaluate to True.
Refer to numpy.any for full documentation.
numpy.any : equivalent function
$\operatorname{argmax}$

IsErrorArray. argmax (axis=None, out=None)
Return indices of the maximum values along the given axis.
Refer to numpy.argmax for full documentation.
numpy.argmax : equivalent function
argmin

IsErrorArray.argmin (axis=None, out=None)
Return indices of the minimum values along the given axis of $a$.
Refer to numpy.argmin for detailed documentation.
numpy.argmin : equivalent function

## argpartition

IsErrorArray.argpartition (kth, axis=-1, kind='introselect', order=None)
Returns the indices that would partition this array.
Refer to numpy.argpartition for full documentation.
New in version 1.8.0.
numpy.argpartition : equivalent function
argsort

IsErrorArray.argsort (axis=-1, kind='quicksort', order=None)
Returns the indices that would sort this array.
Refer to numpy.argsort for full documentation.
numpy.argsort : equivalent function
astype

IsErrorArray .astype (dtype, order='K', casting='unsafe', subok=True, copy=True)
Copy of the array, cast to a specified type.
dtype [str or dtype] Typecode or data-type to which the array is cast.
order [ $\{$ ' C ', ' F ', ' A ', ' K '\}, optional] Controls the memory layout order of the result. ' C ' means C order, ' F ' means Fortran order, ' $A$ ' means ' F ' order if all the arrays are Fortran contiguous, ' C ' order otherwise, and ' K ' means as close to the order the array elements appear in memory as possible. Default is ' K '.
casting [\{'no', 'equiv', 'safe', 'same_kind', 'unsafe' \}, optional] Controls what kind of data casting may occur. Defaults to 'unsafe' for backwards compatibility.

- 'no' means the data types should not be cast at all.
- 'equiv' means only byte-order changes are allowed.
- 'safe' means only casts which can preserve values are allowed.
- 'same_kind' means only safe casts or casts within a kind, like float64 to float32, are allowed.
- 'unsafe' means any data conversions may be done.
subok [bool, optional] If True, then sub-classes will be passed-through (default), otherwise the returned array will be forced to be a base-class array.
copy [bool, optional] By default, astype always returns a newly allocated array. If this is set to false, and the dtype, order, and subok requirements are satisfied, the input array is returned instead of a copy.
arr_t [ndarray] Unless copy is False and the other conditions for returning the input array are satisfied (see description for copy input parameter), arr_t is a new array of the same shape as the input array, with dtype, order given by dtype, order.

Starting in NumPy 1.9, astype method now returns an error if the string dtype to cast to is not long enough in 'safe' casting mode to hold the max value of integer/float array that is being casted. Previously the casting was allowed even if the result was truncated.

ComplexWarning When casting from complex to float or int. To avoid this, one should use a. real. astype (t).

```
>>> x = np.array([1, 2, 2.5])
>>> x
array([ 1. , 2. , 2.5])
```

```
>>> x.astype(int)
array([1, 2, 2])
```


## byteswap

## IsErrorArray.byteswap (inplace)

Swap the bytes of the array elements
Toggle between low-endian and big-endian data representation by returning a byteswapped array, optionally swapped in-place.
inplace [bool, optional] If True, swap bytes in-place, default is False.
out [ndarray] The byteswapped array. If inplace is True, this is a view to self.

```
>>> A = np.array([1, 256, 8755], dtype=np.int16)
>>> map (hex, A)
['0x1', '0x100', '0x2233']
```

(continued from previous page)

```
>>> A.byteswap (True)
array([ 256, 1, 13090], dtype=int16)
>>> map (hex, A)
['0x100', '0x1', '0x3322']
```

Arrays of strings are not swapped

```
>>> A = np.array(['ceg', 'fac'])
>>> A.byteswap()
array(['ceg', 'fac'],
    dtype='|S3')
```

choose

IsErrorArray. choose (choices, out=None, mode='raise')
Use an index array to construct a new array from a set of choices.
Refer to numpy.choose for full documentation.
numpy.choose : equivalent function
clip

IsErrorArray.clip (min=None, max=None, out=None)
Return an array whose values are limited to [min, max]. One of max or min must be given.
Refer to numpy.clip for full documentation.
numpy.clip : equivalent function
collapse

IsErrorArray.collapse (shape)
compress

IsErrorArray. compress (condition, axis=None, out=None)
Return selected slices of this array along given axis.
Refer to numpy.compress for full documentation.
numpy.compress : equivalent function
conj

IsErrorArray.conj()
Complex-conjugate all elements.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function

## conjugate

IsErrorArray.conjugate()
Return the complex conjugate, element-wise.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function
copy

IsErrorArray. copy (order='C')
Return a copy of the array.
order [ $\{$ ' C ', ' F ', ' A ', ' K '\}, optional] Controls the memory layout of the copy. ' C ' means C -order, ' F ' means F-order, 'A' means ' F ' if $a$ is Fortran contiguous, ' C ' otherwise. ' K ' means match the layout of $a$ as closely as possible. (Note that this function and :func:numpy.copy are very similar, but have different default values for their order $=$ arguments.)
numpy.copy numpy.copyto

```
>>> x = np.array([[1,2,3],[4,5,6]], order='F')
```

```
>>> y = x.copy()
```

>>> x.fill(0)
>>> $x$
array ([[0, 0, 0],
$[0,0,0]])$

```
>>> y
array([[1, 2, 3],
    [4, 5, 6]])
```

```
>>> y.flags['C_CONTIGUOUS']
True
```


## cumprod

IsErrorArray. Cumprod (axis=None, dtype=None, out=None)
Return the cumulative product of the elements along the given axis.
Refer to numpy.cumprod for full documentation.
numpy.cumprod : equivalent function

## cumsum

IsErrorArray. cumsum (axis=None, dtype=None, out=None)
Return the cumulative sum of the elements along the given axis.
Refer to numpy.cumsum for full documentation.
numpy.cumsum : equivalent function

## diagonal

IsErrorArray.diagonal (offset $=0$, axis $1=0$, axis $2=1$ )
Return specified diagonals. In NumPy 1.9 the returned array is a read-only view instead of a copy as in previous NumPy versions. In a future version the read-only restriction will be removed.

Refer to numpy. diagonal () for full documentation.
numpy.diagonal : equivalent function
dot

IsErrorArray.dot ( $b$, out=None)
Dot product of two arrays.
Refer to numpy.dot for full documentation.
numpy.dot : equivalent function

```
>>> a = np.eye(2)
>>> b = np.ones((2, 2)) * 2
>>> a.dot(b)
array([[ 2., 2.],
    [ 2., 2.]])
```

This array method can be conveniently chained:

```
>>> a.dot(b).dot(b)
array([[ 8., 8.],
    [ 8., 8.]])
```


## dump

IsErrorArray. dump (file)
Dump a pickle of the array to the specified file. The array can be read back with pickle.load or numpy.load.
file [str] A string naming the dump file.

## dumps

## IsErrorArray.dumps()

Returns the pickle of the array as a string. pickle.loads or numpy.loads will convert the string back to an array.

None
fill

IsErrorArray.fill (value)
Fill the array with a scalar value.
value [scalar] All elements of $a$ will be assigned this value.

```
>>> a = np.array([1, 2])
>>> a.fill(0)
>>> a
array([0, 0])
>>> a = np.empty(2)
>>> a.fill(1)
>>> a
array([ 1., 1.])
```


## flatten

IsErrorArray.flatten (order='C')
Return a copy of the array collapsed into one dimension.
order [\{ ' C ', ' F ', ' $A$ ', ' K '\}, optional] ' C ' means to flatten in row-major ( C -style) order. ' F ' means to flatten in column-major (Fortran- style) order. 'A' means to flatten in column-major order if $a$ is Fortran contiguous in memory, row-major order otherwise. ' K ' means to flatten $a$ in the order the elements occur in memory. The default is ' C '.
$\mathbf{y}$ [ndarray] A copy of the input array, flattened to one dimension.
ravel : Return a flattened array. flat: A 1-D flat iterator over the array.

```
>>> a = np.array([[1,2], [3,4]])
>>> a.flatten()
array([1, 2, 3, 4])
>>> a.flatten('F')
array([1, 3, 2, 4])
```


## getfield

## IsErrorArray.getfield (dtype, offset=0)

Returns a field of the given array as a certain type.
A field is a view of the array data with a given data-type. The values in the view are determined by the given type and the offset into the current array in bytes. The offset needs to be such that the view dtype fits in the array dtype; for example an array of dtype complex 128 has 16-byte elements. If taking a view with a 32-bit integer ( 4 bytes), the offset needs to be between 0 and 12 bytes.
dtype [str or dtype] The data type of the view. The dtype size of the view can not be larger than that of the array itself.
offset [int] Number of bytes to skip before beginning the element view.

```
>>> x = np.diag([1.+1.j]*2)
>>> x[1, 1] = 2 + 4.j
>>> x
array([[ 1.+1.j, 0.+0.j],
    [ 0.+0.j, 2.+4.j]])
>>> x.getfield(np.float64)
array([[ 1., 0.],
    [ 0., 2.]])
```

By choosing an offset of 8 bytes we can select the complex part of the array for our view:

```
>>> x.getfield(np.float64, offset=8)
array([[ 1., 0.],
    [ 0., 4.]])
```


## item

## IsErrorArray.item (*args)

Copy an element of an array to a standard Python scalar and return it.
*args : Arguments (variable number and type)

- none: in this case, the method only works for arrays with one element (a.size $==1$ ), which element is copied into a standard Python scalar object and returned.
- int_type: this argument is interpreted as a flat index into the array, specifying which element to copy and return.
- tuple of int_types: functions as does a single int_type argument, except that the argument is interpreted as an nd-index into the array.
z [Standard Python scalar object] A copy of the specified element of the array as a suitable Python scalar
When the data type of $a$ is longdouble or clongdouble, item() returns a scalar array object because there is no available Python scalar that would not lose information. Void arrays return a buffer object for item(), unless fields are defined, in which case a tuple is returned.
item is very similar to a[args], except, instead of an array scalar, a standard Python scalar is returned. This can be useful for speeding up access to elements of the array and doing arithmetic on elements of the array using Python's optimized math.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.item(3)
2
>>> x.item(7)
5
>>> x.item((0, 1))
1
>>> x.item((2, 2))
3
```


## itemset

## IsErrorArray.itemset (*args)

Insert scalar into an array (scalar is cast to array's dtype, if possible)
There must be at least 1 argument, and define the last argument as item. Then, a.itemset (*args) is equivalent to but faster than a [args] = item. The item should be a scalar value and args must select a single item in the array $a$.

1*args [Arguments] If one argument: a scalar, only used in case $a$ is of size 1 . If two arguments: the last argument is the value to be set and must be a scalar, the first argument specifies a single array element location. It is either an int or a tuple.

Compared to indexing syntax, itemset provides some speed increase for placing a scalar into a particular location in an ndarray, if you must do this. However, generally this is discouraged: among other problems, it complicates the appearance of the code. Also, when using itemset (and item) inside a loop, be sure to assign the methods to a local variable to avoid the attribute look-up at each loop iteration.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.itemset (4, 0)
>>> x.itemset ((2, 2), 9)
>>> x
array([[3, 1, 7],
    [2, 0, 3],
    [8, 5, 9]])
```

max

IsErrorArray.max (axis=None, out=None)
Return the maximum along a given axis.
Refer to numpy.amax for full documentation.
numpy.amax : equivalent function
mean

IsErrorArray .mean (axis=None, dtype=None, out=None, , eepdims=False)
Returns the average of the array elements along given axis.
Refer to numpy.mean for full documentation.
numpy.mean : equivalent function
min

IsErrorArray.min (axis=None, out=None, keepdims=False)
Return the minimum along a given axis.
Refer to numpy.amin for full documentation.
numpy.amin : equivalent function
newbyteorder

IsErrorArray. newbyteorder (new_order='S')
Return the array with the same data viewed with a different byte order.
Equivalent to:
arr.view(arr.dtype.newbytorder(new_order))
Changes are also made in all fields and sub-arrays of the array data type.
new_order [string, optional] Byte order to force; a value from the byte order specifications below. new_order codes can be any of:

- 'S' - swap dtype from current to opposite endian
- \{ '<', 'L'\} - little endian
- \{'>', 'B'\} - big endian
- $\{$ ' $=$ ', ' N ' $\}$ - native order
- \{ ' 1 ', 'I' $\}$ - ignore (no change to byte order)

The default value ('S') results in swapping the current byte order. The code does a case-insensitive check on the first letter of new_order for the alternatives above. For example, any of ' B ' or ' $b$ ' or 'biggish' are valid to specify big-endian.
new_arr [array] New array object with the dtype reflecting given change to the byte order.

## nonzero

## IsErrorArray.nonzero ()

Return the indices of the elements that are non-zero.
Refer to numpy.nonzero for full documentation.
numpy.nonzero : equivalent function

## partition

IsErrorArray.partition (kth, axis=-1, kind='introselect', order=None)
Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. All elements smaller than the kth element are moved before this element and all equal or greater are moved behind it. The ordering of the elements in the two partitions is undefined.

New in version 1.8.0.
$\mathbf{k t h}$ [int or sequence of ints] Element index to partition by. The kth element value will be in its final sorted position and all smaller elements will be moved before it and all equal or greater elements behind it. The order all elements in the partitions is undefined. If provided with a sequence of kth it will partition all elements indexed by kth of them into their sorted position at once.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'introselect'\}, optional] Selection algorithm. Default is 'introselect'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.partition : Return a parititioned copy of an array. argpartition : Indirect partition. sort : Full sort.
See np. partition for notes on the different algorithms.

```
>>> a = np.array([3, 4, 2, 1])
>>> a.partition(a, 3)
>>> a
array([2, 1, 3, 4])
```

```
>>> a.partition((1, 3))
array([1, 2, 3, 4])
```

prod

IsErrorArray $\cdot \operatorname{prod}($ axis=None, dtype=None, out=None, keepdims=False $)$
Return the product of the array elements over the given axis
Refer to numpy.prod for full documentation.
numpy.prod : equivalent function
ptp

IsErrorArray.ptp (axis=None, out=None)
Peak to peak (maximum - minimum) value along a given axis.
Refer to numpy.ptp for full documentation.
numpy.ptp : equivalent function
put

IsErrorArray.put (indices, values, mode='raise')
Set a.flat $[\mathrm{n}]=$ values $[\mathrm{n}]$ for all $n$ in indices.
Refer to numpy.put for full documentation.
numpy.put : equivalent function
ravel

IsErrorArray. ravel ([order])
Return a flattened array.
Refer to numpy.ravel for full documentation.
numpy.ravel : equivalent function
ndarray.flat : a flat iterator on the array.
repeat

IsErrorArray.repeat (repeats, axis=None)
Repeat elements of an array.
Refer to numpy.repeat for full documentation.
numpy.repeat : equivalent function

## reshape

IsErrorArray. reshape (shape, order='C')
Returns an array containing the same data with a new shape.
Refer to numpy.reshape for full documentation.
numpy.reshape : equivalent function
resize

IsErrorArray. resize (new_shape, refcheck=True)
Change shape and size of array in-place.
new_shape [tuple of ints, or $n$ ints] Shape of resized array.
refcheck [bool, optional] If False, reference count will not be checked. Default is True.
None
ValueError If $a$ does not own its own data or references or views to it exist, and the data memory must be changed.

SystemError If the order keyword argument is specified. This behaviour is a bug in NumPy.
resize : Return a new array with the specified shape.
This reallocates space for the data area if necessary.
Only contiguous arrays (data elements consecutive in memory) can be resized.
The purpose of the reference count check is to make sure you do not use this array as a buffer for another Python object and then reallocate the memory. However, reference counts can increase in other ways so if you are sure that you have not shared the memory for this array with another Python object, then you may safely set refcheck to False.

Shrinking an array: array is flattened (in the order that the data are stored in memory), resized, and reshaped:

```
>>> a = np.array([[0, 1], [2, 3]], order='C')
>>> a.resize((2, 1))
>>> a
array([[0],
    [1]])
```

```
>>> a = np.array([[0, 1], [2, 3]], order='F')
>>> a.resize((2, 1))
>>> a
array([[0],
    [2]])
```

Enlarging an array: as above, but missing entries are filled with zeros:

```
>>> b = np.array([[0, 1], [2, 3]])
>>> b.resize(2, 3) # new_shape parameter doesn't have to be a tuple
>>> b
array([[0, 1, 2],
    [3, 0, 0]])
```

Referencing an array prevents resizing...

```
>>> C = a
>>> a.resize((1, 1))
Traceback (most recent call last):
ValueError: cannot resize an array that has been referenced ...
```

Unless refcheck is False:

```
>>> a.resize((1, 1), refcheck=False)
>>> a
array([[0]])
>>> c
array([[0]])
```

round

IsErrorArray. round (decimals=0, out=None)
Return $a$ with each element rounded to the given number of decimals.
Refer to numpy.around for full documentation.
numpy.around : equivalent function

## searchsorted

IsErrorArray.searchsorted ( $v$, side='left', sorter=None)
Find indices where elements of v should be inserted in a to maintain order.
For full documentation, see numpy.searchsorted
numpy.searchsorted : equivalent function

## setfield

IsErrorArray.setfield (val, dtype, offset=0)
Put a value into a specified place in a field defined by a data-type.
Place val into $a$ 's field defined by dtype and beginning offset bytes into the field.
val [object] Value to be placed in field.
dtype [dtype object] Data-type of the field in which to place val.
offset [int, optional] The number of bytes into the field at which to place val.
None
getfield

```
>>> x = np.eye(3)
>>> x.getfield(np.float64)
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
>>> x.setfield(3, np.int32)
>>> x.getfield(np.int32)
```

```
array([[3, 3, 3],
    [3, 3, 3],
    [3, 3, 3]])
>>> x
array([[ 1.00000000e+000, 1.48219694e-323, 1.48219694e-323],
    [ 1.48219694e-323, 1.00000000e+000, 1.48219694e-323],
    [ 1.48219694e-323, 1.48219694e-323, 1.00000000e+000]])
>>> x.setfield(np.eye(3), np.int32)
>>> x
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
```


## setflags

IsErrorArray.setflags (write $=$ None, align $=$ None, uic $=$ None)
Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively.
These Boolean-valued flags affect how numpy interprets the memory area used by $a$ (see Notes below). The ALIGNED flag can only be set to True if the data is actually aligned according to the type. The UPDATEIFCOPY flag can never be set to True. The flag WRITEABLE can only be set to True if the array owns its own memory, or the ultimate owner of the memory exposes a writeable buffer interface, or is a string. (The exception for string is made so that unpickling can be done without copying memory.)
write [bool, optional] Describes whether or not $a$ can be written to.
align [bool, optional] Describes whether or not $a$ is aligned properly for its type.
uic [bool, optional] Describes whether or not $a$ is a copy of another "base" array.
Array flags provide information about how the memory area used for the array is to be interpreted. There are 6 Boolean flags in use, only three of which can be changed by the user: UPDATEIFCOPY, WRITEABLE, and ALIGNED.

WRITEABLE (W) the data area can be written to;
ALIGNED (A) the data and strides are aligned appropriately for the hardware (as determined by the compiler);

UPDATEIFCOPY (U) this array is a copy of some other array (referenced by .base). When this array is deallocated, the base array will be updated with the contents of this array.

All flags can be accessed using their first (upper case) letter as well as the full name.

```
>>> y
array([[3, 1, 7],
    [2, 0, 0],
    [8, 5, 9]])
>>> y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : True
    ALIGNED : True
    UPDATEIFCOPY : False
>>> y.setflags(write=0, align=0)
>>> y.flags
```

```
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : False
    ALIGNED : False
    UPDATEIFCOPY : False
>>> y.setflags(uic=1)
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: cannot set UPDATEIFCOPY flag to True
```


## sort

IsErrorArray.sort (axis=-1, kind='quicksort', order=None)
Sort an array, in-place.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'quicksort', 'mergesort', 'heapsort'\}, optional] Sorting algorithm. Default is 'quicksort'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.sort : Return a sorted copy of an array. argsort : Indirect sort. lexsort : Indirect stable sort on multiple keys. searchsorted : Find elements in sorted array. partition: Partial sort.

See sort for notes on the different sorting algorithms.

```
>>> a = np.array([[1,4], [3,1]])
>>> a.sort(axis=1)
>>> a
array([[1, 4],
    [1, 3]])
>>> a.sort(axis=0)
>>> a
array([[1, 3],
    [1, 4]])
```

Use the order keyword to specify a field to use when sorting a structured array:

```
>>> a = np.array([('a', 2), ('c', 1)], dtype=[('x', 'S1'), ('y', int)])
>>> a.sort (order='y')
>>> a
array([('c', 1), ('a', 2)],
    dtype=[('x', '|S1'), ('y', '<i4')])
```


## squeeze

## IsErrorArray.squeeze (axis=None)

Remove single-dimensional entries from the shape of $a$.
Refer to numpy.squeeze for full documentation.
numpy.squeeze : equivalent function
std
IsErrorArray.std (axis=None, dtype=None, out=None, ddof=0, keepdims=False)
Returns the standard deviation of the array elements along given axis.
Refer to numpy.std for full documentation.
numpy.std : equivalent function
sum
IsErrorArray. sum (axis=None, dtype=None, out=None, keepdims=False)
Return the sum of the array elements over the given axis.
Refer to numpy.sum for full documentation.
numpy.sum : equivalent function
swapaxes
IsErrorArray.swapaxes (axisl, axis2)
Return a view of the array with axis1 and axis2 interchanged.
Refer to numpy.swapaxes for full documentation.
numpy.swapaxes : equivalent function

## take

IsErrorArray.take (indices, axis=None, out=None, mode='raise')
Return an array formed from the elements of $a$ at the given indices.
Refer to numpy.take for full documentation.
numpy.take : equivalent function

## tobytes

IsErrorArray.tobytes (order='C')
Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C -order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.
New in version 1.9.0.
order [\{ ' C ', ' F ', None \}, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01\x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01 \x00\x00\x00\x03\x00\x00\x00'
```

tofile

IsErrorArray.tofile (fid, sep="", format="\%s")
Write array to a file as text or binary (default).
Data is always written in ' C ' order, independent of the order of $a$. The data produced by this method can be recovered using the function fromfile().
fid [file or str] An open file object, or a string containing a filename.
sep [str] Separator between array items for text output. If "" (empty), a binary file is written, equivalent to file.write(a.tobytes()).
format [str] Format string for text file output. Each entry in the array is formatted to text by first converting it to the closest Python type, and then using "format" \% item.

This is a convenience function for quick storage of array data. Information on endianness and precision is lost, so this method is not a good choice for files intended to archive data or transport data between machines with different endianness. Some of these problems can be overcome by outputting the data as text files, at the expense of speed and file size.

## tolist

IsErrorArray.tolist()
Return the array as a (possibly nested) list.
Return a copy of the array data as a (nested) Python list. Data items are converted to the nearest compatible Python type.
none
$\mathbf{y}$ [list] The possibly nested list of array elements.
The array may be recreated, $a=n p . a r r a y(a . t o l i s t())$.

```
>>> a = np.array([1, 2])
>>> a.tolist()
[1, 2]
>>> a = np.array([[1, 2], [3, 4]])
>>> list(a)
[array([1, 2]), array([3, 4])]
>>> a.tolist()
[[1, 2], [3, 4]]
```

tostring

IsErrorArray.tostring (order='C')
Construct Python bytes containing the raw data bytes in the array.

Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.
This function is a compatibility alias for tobytes. Despite its name it returns bytes not strings.
order [ $\{$ ' $C$ ', ' $F$ ', None \}, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01 \x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01\x00\x00\x00\x03\x00\x00\x00'
```

trace

IsErrorArray.trace (offset=0, axis $1=0$, axis $2=1$, dtype $=$ None, out $=$ None)
Return the sum along diagonals of the array.
Refer to numpy.trace for full documentation.
numpy.trace : equivalent function

## transpose

## IsErrorArray.transpose (*axes)

Returns a view of the array with axes transposed.
For a 1-D array, this has no effect. (To change between column and row vectors, first cast the 1-D array into a matrix object.) For a 2-D array, this is the usual matrix transpose. For an n-D array, if axes are given, their order indicates how the axes are permuted (see Examples). If axes are not provided anda.shape $=(i[0], i[1], \ldots i[n-2], i[n-1])$, then a.transpose(). shape $=$ (i[n-1], i[n-2], ... i[1], i[0]).
axes: None, tuple of ints, or $n$ ints

- None or no argument: reverses the order of the axes.
- tuple of ints: $i$ in the $j$-th place in the tuple means $a$ 's $i$-th axis becomes a.transpose()'s $j$-th axis.
- $n$ ints: same as an $n$-tuple of the same ints (this form is intended simply as a "convenience" alternative to the tuple form)
out [ndarray] View of $a$, with axes suitably permuted.
ndarray.T : Array property returning the array transposed.

```
>>> a = np.array([[1, 2], [3, 4]])
>>> a
array([[1, 2],
    [3, 4]])
```

```
>>> a.transpose()
array([[1, 3],
    [2, 4]])
>>> a.transpose((1, 0))
array([[1, 3],
    [2, 4]])
>>> a.transpose(1, 0)
array([[1, 3],
    [2, 4]])
```

var

IsErrorArray.var (axis=None, dtype=None, out=None, ddof=0, , eeepdims=False )
Returns the variance of the array elements, along given axis.
Refer to numpy.var for full documentation.
numpy.var : equivalent function

## view

IsErrorArray.view (dtype=None, type=None)
New view of array with the same data.
dtype [data-type or ndarray sub-class, optional] Data-type descriptor of the returned view, e.g., float32 or int16. The default, None, results in the view having the same data-type as $a$. This argument can also be specified as an ndarray sub-class, which then specifies the type of the returned object (this is equivalent to setting the type parameter).
type [Python type, optional] Type of the returned view, e.g., ndarray or matrix. Again, the default None results in type preservation.
a. view () is used two different ways:
a.view (some_dtype) or a.view (dtype=some_dtype) constructs a view of the array's memory with a different data-type. This can cause a reinterpretation of the bytes of memory.
a.view(ndarray_subclass) or a.view(type=ndarray_subclass) just returns an instance of ndarray_subclass that looks at the same array (same shape, dtype, etc.) This does not cause a reinterpretation of the memory.

For a.view (some_dtype), if some_dtype has a different number of bytes per entry than the previous dtype (for example, converting a regular array to a structured array), then the behavior of the view cannot be predicted just from the superficial appearance of a (shown by print (a) ). It also depends on exactly how a is stored in memory. Therefore if a is C-ordered versus fortran-ordered, versus defined as a slice or transpose, etc., the view may give different results.

```
>>> x = np.array([(1, 2)], dtype=[('a', np.int8), ('b', np.int8)])
```

Viewing array data using a different type and dtype:

```
>>> y = x.view(dtype=np.int16, type=np.matrix)
>>> y
matrix([[513]], dtype=int16)
```

```
>>> print(type(y))
<class 'numpy.matrixlib.defmatrix.matrix'>
```

Creating a view on a structured array so it can be used in calculations

```
>>> x = np.array([(1, 2), (3,4)], dtype=[('a', np.int8), ('b', np.int8)])
>>> xv = x.view(dtype=np.int8).reshape(-1,2)
>>> xv
array([[1, 2],
    [3, 4]], dtype=int8)
>>> xv.mean(0)
array([ 2., 3.])
```

Making changes to the view changes the underlying array

```
>>> xv[0,1] = 20
>>> print(x)
[(1, 20) (3, 4)]
```

Using a view to convert an array to a recarray:

```
>>> z = x.view(np.recarray)
>>> z.a
array([1], dtype=int8)
```

Views share data:

```
>>> x[0] = (9, 10)
>>> z[0]
(9, 10)
```

Views that change the dtype size (bytes per entry) should normally be avoided on arrays defined by slices, transposes, fortran-ordering, etc.:

```
>>> x = np.array([[1,2,3],[4,5,6]], dtype=np.int16)
>>> y = x[:, 0:2]
>>> y
array([[1, 2],
    [4, 5]], dtype=int16)
>>> y.view(dtype=[('width', np.int16), ('length', np.int16)])
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: new type not compatible with array.
>>> z = y.copy()
>>> z.view(dtype=[('width', np.int16), ('length', np.int16)])
array([[(1, 2)],
    [(4, 5)]], dtype=[('width', '<i2'), ('length', '<i2')])
```

_init__()

Initialize self. See help(type(self)) for accurate signature.

## Attributes

| T | Same as self.transpose(), except that self is returned <br> if self.ndim < 2. |
| :--- | :--- |
| base | Base object if memory is from some other object. |
| ctypes | An object to simplify the interaction of the array with <br> the ctypes module. |
| data | Python buffer object pointing to the start of the ar- <br> ray's data. |
| dtype | Data-type of the array's elements. |
| flags | Information about the memory layout of the array. |
| flat | A 1-D iterator over the array. |
| imag | The imaginary part of the array. |
| itemsize | Length of one array element in bytes. |
| nbytes | Total bytes consumed by the elements of the array. |
| ndim | Number of array dimensions. |
| real | The real part of the array. |
| shape | Tuple of array dimensions. |
| size | Number of elements in the array. |
| strides | Tuple of bytes to step in each dimension when <br> traversing an array. |

T

IsErrorArray.T
Same as self.transpose(), except that self is returned if self.ndim $<2$.

```
>>> x = np.array([[1.,2.],[3.,4.]])
>>> x
array([[ 1., 2.],
    [ 3., 4.]])
>>> x.T
array([[ 1., 3.],
    [ 2., 4.]])
>>> x = np.array([1.,2.,3.,4.])
>>> x
array([ 1., 2., 3., 4.])
>>> x.T
array([ 1., 2., 3., 4.])
```

base

IsErrorArray.base
Base object if memory is from some other object.
The base of an array that owns its memory is None:

```
>>> x = np.array([1,2,3,4])
>>> x.base is None
True
```

Slicing creates a view, whose memory is shared with x :

```
>>> y = x[2:]
>>> y.base is x
True
```


## ctypes

## IsErrorArray.ctypes

An object to simplify the interaction of the array with the ctypes module.
This attribute creates an object that makes it easier to use arrays when calling shared libraries with the ctypes module. The returned object has, among others, data, shape, and strides attributes (see Notes below) which themselves return ctypes objects that can be used as arguments to a shared library.

## None

c [Python object] Possessing attributes data, shape, strides, etc.
numpy.ctypeslib
Below are the public attributes of this object which were documented in "Guide to NumPy" (we have omitted undocumented public attributes, as well as documented private attributes):

- data: A pointer to the memory area of the array as a Python integer. This memory area may contain data that is not aligned, or not in correct byte-order. The memory area may not even be writeable. The array flags and data-type of this array should be respected when passing this attribute to arbitrary C-code to avoid trouble that can include Python crashing. User Beware! The value of this attribute is exactly the same as self._array_interface_['data'][0].
- shape (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the C-integer corresponding to dtype( ' p ') on this platform. This base-type could be c_int, c_long, or c_longlong depending on the platform. The c_intp type is defined accordingly in numpy.ctypeslib. The ctypes array contains the shape of the underlying array.
- strides (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the same as for the shape attribute. This ctypes array contains the strides information from the underlying array. This strides information is important for showing how many bytes must be jumped to get to the next element in the array.
- data_as(obj): Return the data pointer cast to a particular c-types object. For example, calling self._as_parameter_ is equivalent to self.data_as(ctypes.c_void_p). Perhaps you want to use the data as a pointer to a ctypes array of floating-point data: self.data_as(ctypes.POINTER(ctypes.c_double)).
- shape_as(obj): Return the shape tuple as an array of some other c-types type. For example: self.shape_as(ctypes.c_short).
- strides_as(obj): Return the strides tuple as an array of some other c-types type. For example: self.strides_as(ctypes.c_longlong).

Be careful using the ctypes attribute - especially on temporary arrays or arrays constructed on the fly. For example, calling $(a+b)$. ctypes.data_as (ctypes.c_void_p) returns a pointer to memory that is invalid because the array created as $(a+b)$ is deallocated before the next Python statement. You can avoid this problem using either $c=a+b$ or $c t=(a+b) . c t y p e s$. In the latter case, ct will hold a reference to the array until ct is deleted or re-assigned.

If the ctypes module is not available, then the ctypes attribute of array objects still returns something useful, but ctypes objects are not returned and errors may be raised instead. In particular, the object will still have the as parameter attribute which will return an integer equal to the data attribute.

```
>>> import ctypes
>>> x
array([[0, 1],
    [2, 3]])
>>> x.ctypes.data
30439712
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long))
<ctypes.LP_c_long object at 0x01F01300>
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long)).contents
c_long(0)
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_longlong)).contents
c_longlong(4294967296L)
>>> x.ctypes.shape
<numpy.core._internal.c_long_Array_2 object at 0x01FFD580>
>>> x.ctypes.shape_as(ctypes.c_long)
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides_as(ctypes.c_longlong)
<numpy.core._internal.c_longlong_Array_2 object at 0x01F01300>
```

data

IsErrorArray.data
Python buffer object pointing to the start of the array's data.

## dtype

IsErrorArray.dtype
Data-type of the array's elements.
None
d: numpy dtype object
numpy.dtype

```
>>> x
array([[0, 1],
    [2, 3]])
>>> x.dtype
dtype('int32')
>>> type(x.dtype)
<type 'numpy.dtype'>
```


## flags

## IsErrorArray.flags

Information about the memory layout of the array.
C_CONTIGUOUS (C) The data is in a single, C-style contiguous segment.
F_CONTIGUOUS (F) The data is in a single, Fortran-style contiguous segment.
OWNDATA (O) The array owns the memory it uses or borrows it from another object.

WRITEABLE (W) The data area can be written to. Setting this to False locks the data, making it readonly. A view (slice, etc.) inherits WRITEABLE from its base array at creation time, but a view of a writeable array may be subsequently locked while the base array remains writeable. (The opposite is not true, in that a view of a locked array may not be made writeable. However, currently, locking a base object does not lock any views that already reference it, so under that circumstance it is possible to alter the contents of a locked array via a previously created writeable view onto it.) Attempting to change a non-writeable array raises a RuntimeError exception.

ALIGNED (A) The data and all elements are aligned appropriately for the hardware.
UPDATEIFCOPY ( $\mathbf{U}$ ) This array is a copy of some other array. When this array is deallocated, the base array will be updated with the contents of this array.

FNC F_CONTIGUOUS and not C_CONTIGUOUS.
FORC F_CONTIGUOUS or C_CONTIGUOUS (one-segment test).

## BEHAVED (B) ALIGNED and WRITEABLE.

CARRAY (CA) BEHAVED and C_CONTIGUOUS.
FARRAY (FA) BEHAVED and F_CONTIGUOUS and not C_CONTIGUOUS.
The flags object can be accessed dictionary-like (as in a.flags ['WRITEABLE']), or by using lowercased attribute names (as in a.flags.writeable). Short flag names are only supported in dictionary access.

Only the UPDATEIFCOPY, WRITEABLE, and ALIGNED flags can be changed by the user, via direct assignment to the attribute or dictionary entry, or by calling ndarray.setflags.

The array flags cannot be set arbitrarily:

- UPDATEIFCOPY can only be set False.
- ALIGNED can only be set True if the data is truly aligned.
- WRITEABLE can only be set True if the array owns its own memory or the ultimate owner of the memory exposes a writeable buffer interface or is a string.

Arrays can be both C-style and Fortran-style contiguous simultaneously. This is clear for 1-dimensional arrays, but can also be true for higher dimensional arrays.

Even for contiguous arrays a stride for a given dimension arr.strides [dim] may be arbitrary if arr.shape[dim] == 1 or the array has no elements. It does not generally hold that self. strides[-1] == self.itemsize for C-style contiguous arrays or self.strides[0] == self.itemsize for Fortran-style contiguous arrays is true.

## flat

## IsErrorArray.flat

A 1-D iterator over the array.
This is a numpy.flatiter instance, which acts similarly to, but is not a subclass of, Python's built-in iterator object.
flatten : Return a copy of the array collapsed into one dimension.
flatiter

```
>>> x = np.arange(1, 7).reshape(2, 3)
>>> x
array([[1, 2, 3],
    [4, 5, 6]])
>>> x.flat[3]
4
>>> x.T
array([[1, 4],
    [2, 5],
    [3, 6]])
>>> x.T.flat[3]
5
>>> type(x.flat)
<type 'numpy.flatiter'>
```

An assignment example:

```
>>> x.flat = 3; x
array([[3, 3, 3],
    [3, 3, 3]])
>>> x.flat[[1,4]] = 1; x
array([[3, 1, 3],
    [3, 1, 3]])
```

imag

## IsErrorArray.imag

The imaginary part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.imag
array([ 0. , 0.70710678])
>>> x.imag.dtype
dtype('float64')
```


## itemsize

IsErrorArray.itemsize
Length of one array element in bytes.

```
>>> x = np.array([1,2,3], dtype=np.float64)
>>> x.itemsize
8
>>> x = np.array([1,2,3], dtype=np.complex128)
>>> x.itemsize
16
```


## nbytes

## IsErrorArray.nbytes

Total bytes consumed by the elements of the array.
Does not include memory consumed by non-element attributes of the array object.

```
>>> x = np.zeros((3,5,2), dtype=np.complex128)
>>> x.nbytes
480
>>> np.prod(x.shape) * x.itemsize
480
```

ndim

IsErrorArray.ndim
Number of array dimensions.

```
>>> x = np.array([1, 2, 3])
>>> x.ndim
1
>>> y = np.zeros((2, 3, 4))
>>> y.ndim
3
```

real

IsErrorArray.real
The real part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.real
array([ 1. , 0.70710678])
>>> x.real.dtype
dtype('float64')
```

numpy.real : equivalent function
shape

IsErrorArray.shape
Tuple of array dimensions.
May be used to "reshape" the array, as long as this would not require a change in the total number of elements

```
>>> x = np.array([1, 2, 3, 4])
>>> x.shape
(4,)
>>> y = np.zeros((2, 3, 4))
>>> y.shape
(2, 3, 4)
>>> y.shape = (3, 8)
>>> y
array([[ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.]])
>>> y.shape = (3, 6)
Traceback (most recent call last):
```

```
    File "<stdin>", line 1, in <module>
ValueError: total size of new array must be unchanged
```


## size

IsErrorArray.size
Number of elements in the array.
Equivalent to np.prod (a.shape), i.e., the product of the array's dimensions.

```
>>> x = np.zeros((3, 5, 2), dtype=np.complex128)
>>> x.size
30
>>> np.prod(x.shape)
30
```


## strides

## IsErrorArray.strides

Tuple of bytes to step in each dimension when traversing an array.
The byte offset of element (i[0], i[1], ..., i[n]) in an array $a$ is:
offset $=$ sum(np.array(i) * a.strides)

A more detailed explanation of strides can be found in the "ndarray.rst" file in the NumPy reference guide.
Imagine an array of 32-bit integers (each 4 bytes):
$x=n p \cdot \operatorname{array}([[0,1,2,3,4]$,
$[5,6,7,8,9]]$, dtype=np.int 32)
This array is stored in memory as 40 bytes, one after the other (known as a contiguous block of memory). The strides of an array tell us how many bytes we have to skip in memory to move to the next position along a certain axis. For example, we have to skip 4 bytes ( 1 value) to move to the next column, but 20 bytes ( 5 values) to get to the same position in the next row. As such, the strides for the array $x$ will be $(20,4)$.
numpy.lib.stride_tricks.as_strided

```
>>> y = np.reshape(np.arange (2*3*4), (2, 3, 4))
>>> y
array([[[ 0, 1, 2, 3],
    [4, 5, 6, 7],
    [ 8, 9, 10, 11]],
    [[12, 13, 14, 15],
    [16, 17, 18, 19],
    [20, 21, 22, 23]]])
>>> y.strides
(48, 16, 4)
>>> y[1,1,1]
17
>>> offset=sum(y.strides * np.array((1,1,1)))
```

```
>>> offset/y.itemsize
```

17

```
>>> x = np.reshape(np.arange(5*6*7*8), (5, 6, 7, 8)).transpose (2, 3, 1,0)
>>> x.strides
(32, 4, 224, 1344)
>>> i = np.array ([3,5,2,2])
>>> offset = sum(i * x.strides)
>>> x[3,5,2,2]
813
>>> offset / x.itemsize
813
```

logic

Python equivalents of logical Excel functions.

## Functions

| solve_cycle |
| :--- |
| xif |
| xiferror |
| solve_cycle |
| solve_cycle (*args) |
| xif |
| xif (condition, x=True, y=False) |
| xiferror |
| xiferror (val, val_if_error) |
| xiferror_otype |
| xiferror_otype (val, val_if_error) |
| xifes |


| IfArray |
| :--- |
| IfErrorArray |

## If Array

class IfArray

Methods

| all | Returns True if all elements evaluate to True. |
| :---: | :---: |
| any | Returns True if any of the elements of $a$ evaluate to True. |
| argmax | Return indices of the maximum values along the given axis. |
| argmin | Return indices of the minimum values along the given axis of $a$. |
| argpartition | Returns the indices that would partition this array. |
| argsort | Returns the indices that would sort this array. |
| astype | Copy of the array, cast to a specified type. |
| byteswap | Swap the bytes of the array elements |
| choose | Use an index array to construct a new array from a set of choices. |
| clip | Return an array whose values are limited to [min, max]. |
| collapse |  |
| compress | Return selected slices of this array along given axis. |
| conj | Complex-conjugate all elements. |
| conjugate | Return the complex conjugate, element-wise. |
| copy | Return a copy of the array. |
| cumprod | Return the cumulative product of the elements along the given axis. |
| cumsum | Return the cumulative sum of the elements along the given axis. |
| diagonal | Return specified diagonals. |
| dot | Dot product of two arrays. |
| dump | Dump a pickle of the array to the specified file. |
| dumps | Returns the pickle of the array as a string. |
| fill | Fill the array with a scalar value. |
| flatten | Return a copy of the array collapsed into one dimension. |
| getfield | Returns a field of the given array as a certain type. |
| item | Copy an element of an array to a standard Python scalar and return it. |
| itemset | Insert scalar into an array (scalar is cast to array's dtype, if possible) |
| max | Return the maximum along a given axis. |
| mean | Returns the average of the array elements along given axis. |
| min | Return the minimum along a given axis. |
| newbyteorder | Return the array with the same data viewed with a different byte order. |
| nonzero | Return the indices of the elements that are non-zero. |

Continued on next page

Table 53 - continued from previous page

| partition | Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. |
| :---: | :---: |
| prod | Return the product of the array elements over the given axis |
| ptp | Peak to peak (maximum - minimum) value along a given axis. |
| put | Set a.flat $[\mathrm{n}]=$ values $[\mathrm{n}]$ for all $n$ in indices. |
| ravel | Return a flattened array. |
| repeat | Repeat elements of an array. |
| reshape | Returns an array containing the same data with a new shape. |
| resize | Change shape and size of array in-place. |
| round | Return $a$ with each element rounded to the given number of decimals. |
| searchsorted | Find indices where elements of v should be inserted in a to maintain order. |
| setfield | Put a value into a specified place in a field defined by a data-type. |
| setflags | Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively. |
| sort | Sort an array, in-place. |
| squeeze | Remove single-dimensional entries from the shape of $a$. |
| std | Returns the standard deviation of the array elements along given axis. |
| sum | Return the sum of the array elements over the given axis. |
| swapaxes | Return a view of the array with axis1 and axis2 interchanged. |
| take | Return an array formed from the elements of $a$ at the given indices. |
| tobytes | Construct Python bytes containing the raw data bytes in the array. |
| tofile | Write array to a file as text or binary (default). |
| tolist | Return the array as a (possibly nested) list. |
| tostring | Construct Python bytes containing the raw data bytes in the array. |
| trace | Return the sum along diagonals of the array. |
| transpose | Returns a view of the array with axes transposed. |
| var | Returns the variance of the array elements, along given axis. |
| view | New view of array with the same data. |

all

IfArray.all (axis=None, out=None, keepdims=False)
Returns True if all elements evaluate to True.
Refer to numpy.all for full documentation.
numpy.all : equivalent function

## any

IfArray. any (axis=None, out=None, keepdims=False)
Returns True if any of the elements of $a$ evaluate to True.
Refer to numpy.any for full documentation.
numpy.any : equivalent function

## argmax

IfArray. argmax (axis=None, out=None)
Return indices of the maximum values along the given axis.
Refer to numpy.argmax for full documentation.
numpy.argmax : equivalent function
argmin

IfArray.argmin (axis=None, out=None)
Return indices of the minimum values along the given axis of $a$.
Refer to numpy.argmin for detailed documentation.
numpy.argmin : equivalent function

## argpartition

IfArray. argpartition (kth, axis=-1, kind='introselect', order=None)
Returns the indices that would partition this array.
Refer to numpy.argpartition for full documentation.
New in version 1.8.0.
numpy.argpartition : equivalent function
argsort

IfArray. argsort (axis=-1, kind='quicksort', order=None)
Returns the indices that would sort this array.
Refer to numpy.argsort for full documentation.
numpy.argsort : equivalent function
astype

IfArray. astype (dtype, order=' $K^{\prime}$, casting='unsafe', subok=True, copy=True)
Copy of the array, cast to a specified type.
dtype [str or dtype] Typecode or data-type to which the array is cast.
order [\{ ' C ', ' F ', ' A ', ' K '\}, optional] Controls the memory layout order of the result. ' C ' means C order, ' $F$ ' means Fortran order, ' $A$ ' means ' $F$ ' order if all the arrays are Fortran contiguous, ' $C$ ' order otherwise, and ' K ' means as close to the order the array elements appear in memory as possible. Default is ' $K$ '.
casting [\{'no', 'equiv', 'safe', 'same_kind', 'unsafe' \}, optional] Controls what kind of data casting may occur. Defaults to 'unsafe' for backwards compatibility.

- 'no' means the data types should not be cast at all.
- 'equiv' means only byte-order changes are allowed.
- 'safe' means only casts which can preserve values are allowed.
- 'same_kind' means only safe casts or casts within a kind, like float64 to float32, are allowed.
- 'unsafe' means any data conversions may be done.
subok [bool, optional] If True, then sub-classes will be passed-through (default), otherwise the returned array will be forced to be a base-class array.
copy [bool, optional] By default, astype always returns a newly allocated array. If this is set to false, and the dtype, order, and subok requirements are satisfied, the input array is returned instead of a copy.
arr_t [ndarray] Unless copy is False and the other conditions for returning the input array are satisfied (see description for copy input parameter), arr_t is a new array of the same shape as the input array, with dtype, order given by dtype, order.

Starting in NumPy 1.9, astype method now returns an error if the string dtype to cast to is not long enough in 'safe' casting mode to hold the max value of integer/float array that is being casted. Previously the casting was allowed even if the result was truncated.
ComplexWarning When casting from complex to float or int. To avoid this, one should use a. real. astype(t).

```
>>> x = np.array([1, 2, 2.5])
>>> x
array([ 1. , 2. , 2.5])
```

```
>>> x.astype(int)
array([1, 2, 2])
```


## byteswap

## IfArray.byteswap (inplace)

Swap the bytes of the array elements
Toggle between low-endian and big-endian data representation by returning a byteswapped array, optionally swapped in-place.
inplace [bool, optional] If True, swap bytes in-place, default is False.
out [ndarray] The byteswapped array. If inplace is True, this is a view to self.

```
>>> A = np.array([1, 256, 8755], dtype=np.int16)
>>> map(hex, A)
['0x1', '0x100', '0x2233']
```

(continues on next page)

```
>>> A.byteswap (True)
array([ 256, 1, 13090], dtype=int16)
>>> map(hex, A)
['0x100', '0x1', '0x3322']
```

Arrays of strings are not swapped

```
>>> A = np.array(['ceg', 'fac'])
>>> A.byteswap()
array(['ceg', 'fac'],
    dtype='|S3')
```

choose
IfArray. choose (choices, out=None, mode='raise')
Use an index array to construct a new array from a set of choices.

Refer to numpy.choose for full documentation.
numpy.choose : equivalent function
clip

IfArray.clip (min=None, max=None, out=None)
Return an array whose values are limited to [min, max]. One of max or min must be given.
Refer to numpy.clip for full documentation.
numpy.clip : equivalent function
collapse

IfArray.collapse (shape)
compress

IfArray. compress (condition, axis=None, out=None)
Return selected slices of this array along given axis.
Refer to numpy.compress for full documentation.
numpy.compress : equivalent function
conj

IfArray.conj()
Complex-conjugate all elements.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function

## conjugate

IfArray. conjugate ()
Return the complex conjugate, element-wise.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function
copy

IfArray. copy (order='C')
Return a copy of the array.
order [ $\{$ ' C ', ' F ', ' A ', ' K '\}, optional] Controls the memory layout of the copy. ' C ' means C -order, ' F ' means F -order, ' A ' means ' F ' if $a$ is Fortran contiguous, ' C ' otherwise. ' K ' means match the layout of $a$ as closely as possible. (Note that this function and :func:numpy.copy are very similar, but have different default values for their order= arguments.)
numpy.copy numpy.copyto

```
>>> x = np.array([[1,2,3],[4,5,6]], order='F')
```

```
>>> y = x.copy()
```

```
>>> x.fill(0)
```

```
>>> x
array([[0, 0, 0],
    [0, 0, 0]])
```

```
>>> y
array([[1, 2, 3],
    [4, 5, 6]])
```

>>> y.flags['C_CONTIGUOUS']
True
cumprod

IfArray. cumprod (axis=None, dtype=None, out=None)
Return the cumulative product of the elements along the given axis.
Refer to numpy.cumprod for full documentation.
numpy.cumprod : equivalent function

## cumsum

IfArray. cumsum (axis=None, dtype=None, out=None)
Return the cumulative sum of the elements along the given axis.
Refer to numpy.cumsum for full documentation.
numpy.cumsum : equivalent function

## diagonal

IfArray.diagonal (offset=0, axis $1=0$, axis $2=1$ )
Return specified diagonals. In NumPy 1.9 the returned array is a read-only view instead of a copy as in previous NumPy versions. In a future version the read-only restriction will be removed.

Refer to numpy. diagonal () for full documentation.
numpy.diagonal : equivalent function
dot

IfArray. dot ( $b$, out=None)
Dot product of two arrays.
Refer to numpy.dot for full documentation.
numpy.dot : equivalent function

```
>>> a = np.eye(2)
>>> b = np.ones((2, 2)) * 2
>>> a.dot(b)
array([[ 2., 2.],
    [ 2., 2.]])
```

This array method can be conveniently chained:

```
>>> a.dot(b).dot(b)
array([[ 8., 8.],
    [ 8., 8.]])
```


## dump

IfArray. dump (file)
Dump a pickle of the array to the specified file. The array can be read back with pickle.load or numpy.load.
file [str] A string naming the dump file.

## dumps

```
IfArray.dumps()
Returns the pickle of the array as a string. pickle.loads or numpy.loads will convert the string back to an array.
None
```

fill

IfArray.fill (value)
Fill the array with a scalar value.
value [scalar] All elements of $a$ will be assigned this value.

```
>>> a = np.array([1, 2])
>>> a.fill(0)
>>> a
array([0, 0])
>>> a = np.empty(2)
>>> a.fill(1)
>>> a
array([ 1., 1.])
```


## flatten

IfArray.flatten (order='C')
Return a copy of the array collapsed into one dimension.
order [ $\{$ ' C ', ' F ', ' $A$ ', ' K '\}, optional] ' C ' means to flatten in row-major ( C -style) order. ' F ' means to flatten in column-major (Fortran- style) order. 'A' means to flatten in column-major order if $a$ is Fortran contiguous in memory, row-major order otherwise. ' $K$ ' means to flatten $a$ in the order the elements occur in memory. The default is ' C '.
$\mathbf{y}$ [ndarray] A copy of the input array, flattened to one dimension.
ravel : Return a flattened array. flat: A 1-D flat iterator over the array.

```
>>> a = np.array([[1,2], [3,4]])
>>> a.flatten()
array([1, 2, 3, 4])
>>> a.flatten('F')
array([1, 3, 2, 4])
```


## getfield

IfArray.getfield (dtype, offset=0)
Returns a field of the given array as a certain type.
A field is a view of the array data with a given data-type. The values in the view are determined by the given type and the offset into the current array in bytes. The offset needs to be such that the view dtype fits in the array dtype; for example an array of dtype complex 128 has 16-byte elements. If taking a view with a 32-bit integer ( 4 bytes), the offset needs to be between 0 and 12 bytes.
dtype [str or dtype] The data type of the view. The dtype size of the view can not be larger than that of the array itself.
offset [int] Number of bytes to skip before beginning the element view.

```
>>> x = np.diag([1.+1.j]*2)
>>> x[1, 1] = 2 + 4.j
>>> x
array([[ 1.+1.j, 0.+0.j],
    [ 0.+0.j, 2.+4.j]])
>>> x.getfield(np.float64)
array([[ 1., 0.],
    [ 0., 2.]])
```

By choosing an offset of 8 bytes we can select the complex part of the array for our view:

```
>>> x.getfield(np.float64, offset=8)
array([[ 1., 0.],
    [ 0., 4.]])
```

item

IfArray.item (*args)
Copy an element of an array to a standard Python scalar and return it.
*args : Arguments (variable number and type)

- none: in this case, the method only works for arrays with one element (a.size $==1$ ), which element is copied into a standard Python scalar object and returned.
- int_type: this argument is interpreted as a flat index into the array, specifying which element to copy and return.
- tuple of int_types: functions as does a single int_type argument, except that the argument is interpreted as an nd-index into the array.
$\mathbf{z}$ [Standard Python scalar object] A copy of the specified element of the array as a suitable Python scalar
When the data type of $a$ is longdouble or clongdouble, item() returns a scalar array object because there is no available Python scalar that would not lose information. Void arrays return a buffer object for item(), unless fields are defined, in which case a tuple is returned.
item is very similar to a[args], except, instead of an array scalar, a standard Python scalar is returned. This can be useful for speeding up access to elements of the array and doing arithmetic on elements of the array using Python's optimized math.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.item(3)
2
>>> x.item(7)
5
>>> x.item((0, 1))
1
>>> x.item((2, 2))
3
```


## itemset

IfArray.itemset (*args)
Insert scalar into an array (scalar is cast to array's dtype, if possible)
There must be at least 1 argument, and define the last argument as item. Then, a.itemset (*args) is equivalent to but faster than a [args] = item. The item should be a scalar value and args must select a single item in the array $a$.

1*args [Arguments] If one argument: a scalar, only used in case $a$ is of size 1 . If two arguments: the last argument is the value to be set and must be a scalar, the first argument specifies a single array element location. It is either an int or a tuple.

Compared to indexing syntax, itemset provides some speed increase for placing a scalar into a particular location in an ndarray, if you must do this. However, generally this is discouraged: among other problems, it complicates the appearance of the code. Also, when using itemset (and item) inside a loop, be sure to assign the methods to a local variable to avoid the attribute look-up at each loop iteration.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.itemset (4, 0)
>>> x.itemset ((2, 2), 9)
>>> x
array([[3, 1, 7],
    [2, 0, 3],
    [8, 5, 9]])
```

max

IfArray.max (axis=None, out=None)
Return the maximum along a given axis.
Refer to numpy.amax for full documentation.
numpy.amax : equivalent function
mean

IfArray.mean (axis=None, dtype=None, out=None, keepdims=False)
Returns the average of the array elements along given axis.
Refer to numpy.mean for full documentation.
numpy.mean : equivalent function
min

IfArray.min $($ axis=None, out=None, keepdims $=$ False $)$
Return the minimum along a given axis.
Refer to numpy.amin for full documentation.
numpy.amin : equivalent function
newbyteorder

IfArray. newbyteorder (new_order='S')
Return the array with the same data viewed with a different byte order.
Equivalent to:

```
arr.view(arr.dtype.newbytorder(new_order))
```

Changes are also made in all fields and sub-arrays of the array data type.
new_order [string, optional] Byte order to force; a value from the byte order specifications below. new_order codes can be any of:

- 'S' - swap dtype from current to opposite endian
- \{ '<', 'L'\} - little endian
- \{'>', 'B'\} - big endian
- \{ '=', 'N' $\}$ - native order
- \{ ' 1 ', 'I' $\}$ - ignore (no change to byte order)

The default value ('S') results in swapping the current byte order. The code does a case-insensitive check on the first letter of new_order for the alternatives above. For example, any of ' B ' or ' $b$ ' or 'biggish' are valid to specify big-endian.
new_arr [array] New array object with the dtype reflecting given change to the byte order.

## nonzero

## IfArray.nonzero()

Return the indices of the elements that are non-zero.
Refer to numpy.nonzero for full documentation.
numpy.nonzero : equivalent function

## partition

IfArray.partition (kth, axis=-1, kind='introselect', order=None)
Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. All elements smaller than the kth element are moved before this element and all equal or greater are moved behind it. The ordering of the elements in the two partitions is undefined.

New in version 1.8.0.
kth [int or sequence of ints] Element index to partition by. The kth element value will be in its final sorted position and all smaller elements will be moved before it and all equal or greater elements behind it. The order all elements in the partitions is undefined. If provided with a sequence of kth it will partition all elements indexed by kth of them into their sorted position at once.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'introselect'\}, optional] Selection algorithm. Default is 'introselect'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.partition : Return a parititioned copy of an array. argpartition : Indirect partition. sort : Full sort.
See np. partition for notes on the different algorithms.

```
>>> a = np.array([3, 4, 2, 1])
>>> a.partition(a, 3)
>>> a
array([2, 1, 3, 4])
```

```
>>> a.partition((1, 3))
```

array ([1, 2, 3, 4])
prod

IfArray.prod (axis=None, dtype=None, out=None, keepdims=False)
Return the product of the array elements over the given axis
Refer to numpy.prod for full documentation.
numpy.prod : equivalent function
ptp

IfArray.ptp (axis=None, out=None)
Peak to peak (maximum - minimum) value along a given axis.
Refer to numpy.ptp for full documentation.
numpy.ptp : equivalent function
put

IfArray.put (indices, values, mode='raise')
Seta.flat $[\mathrm{n}]=$ values $[\mathrm{n}]$ for all $n$ in indices.
Refer to numpy.put for full documentation.
numpy.put : equivalent function
ravel

IfArray. ravel ([order])
Return a flattened array.
Refer to numpy.ravel for full documentation.
numpy.ravel : equivalent function
ndarray.flat : a flat iterator on the array.
repeat

IfArray.repeat (repeats, axis=None)
Repeat elements of an array.
Refer to numpy.repeat for full documentation.
numpy.repeat : equivalent function

## reshape

IfArray. reshape (shape, order='C')
Returns an array containing the same data with a new shape.
Refer to numpy.reshape for full documentation.
numpy.reshape : equivalent function
resize

IfArray. resize (new_shape, refcheck=True)
Change shape and size of array in-place.
new_shape [tuple of ints, or $n$ ints] Shape of resized array.
refcheck [bool, optional] If False, reference count will not be checked. Default is True.
None
ValueError If $a$ does not own its own data or references or views to it exist, and the data memory must be changed.

SystemError If the order keyword argument is specified. This behaviour is a bug in NumPy.
resize : Return a new array with the specified shape.
This reallocates space for the data area if necessary.
Only contiguous arrays (data elements consecutive in memory) can be resized.
The purpose of the reference count check is to make sure you do not use this array as a buffer for another Python object and then reallocate the memory. However, reference counts can increase in other ways so if you are sure that you have not shared the memory for this array with another Python object, then you may safely set refcheck to False.

Shrinking an array: array is flattened (in the order that the data are stored in memory), resized, and reshaped:

```
>>> a = np.array([[0, 1], [2, 3]], order='C')
>>> a.resize((2, 1))
>>> a
array([[0],
    [1]])
```

```
>>> a = np.array([[0, 1], [2, 3]], order='F')
>>> a.resize((2, 1))
>>> a
array([[0],
    [2]])
```

Enlarging an array: as above, but missing entries are filled with zeros:

```
>>> b = np.array([[0, 1], [2, 3]])
>>> b.resize(2, 3) # new_shape parameter doesn't have to be a tuple
>>> b
array([[0, 1, 2],
    [3, 0, 0]])
```

Referencing an array prevents resizing...

```
>>> C = a
>>> a.resize((1, 1))
Traceback (most recent call last):
ValueError: cannot resize an array that has been referenced ...
```

Unless refcheck is False:

```
>>> a.resize((1, 1), refcheck=False)
>>> a
array([[0]])
>>> c
array([[0]])
```

round

IfArray. round (decimals=0, out=None)
Return $a$ with each element rounded to the given number of decimals.
Refer to numpy.around for full documentation.
numpy.around : equivalent function

## searchsorted

IfArray.searchsorted ( $v$, side $=$ 'left', sorter $=$ None )
Find indices where elements of v should be inserted in a to maintain order.
For full documentation, see numpy.searchsorted
numpy.searchsorted : equivalent function

## setfield

IfArray.setfield(val, dtype, offset=0)
Put a value into a specified place in a field defined by a data-type.
Place val into $a$ 's field defined by dtype and beginning offset bytes into the field.
val [object] Value to be placed in field.
dtype [dtype object] Data-type of the field in which to place val.
offset [int, optional] The number of bytes into the field at which to place val.
None
getfield

```
>>> x = np.eye(3)
>>> x.getfield(np.float64)
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
>>> x.setfield(3, np.int32)
>>> x.getfield(np.int32)
```

```
array([[3, 3, 3],
    [3, 3, 3],
    [3, 3, 3]])
>>> x
array([[ 1.00000000e+000, 1.48219694e-323, 1.48219694e-323],
    [ 1.48219694e-323, 1.00000000e+000, 1.48219694e-323],
    [ 1.48219694e-323, 1.48219694e-323, 1.00000000e+000]])
>>> x.setfield(np.eye(3), np.int32)
>>> x
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
```


## setflags

IfArray.setflags (write=None, align=None, uic=None)
Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively.
These Boolean-valued flags affect how numpy interprets the memory area used by $a$ (see Notes below). The ALIGNED flag can only be set to True if the data is actually aligned according to the type. The UPDATEIFCOPY flag can never be set to True. The flag WRITEABLE can only be set to True if the array owns its own memory, or the ultimate owner of the memory exposes a writeable buffer interface, or is a string. (The exception for string is made so that unpickling can be done without copying memory.)
write [bool, optional] Describes whether or not $a$ can be written to.
align [bool, optional] Describes whether or not $a$ is aligned properly for its type.
uic [bool, optional] Describes whether or not $a$ is a copy of another "base" array.
Array flags provide information about how the memory area used for the array is to be interpreted. There are 6 Boolean flags in use, only three of which can be changed by the user: UPDATEIFCOPY, WRITEABLE, and ALIGNED.

WRITEABLE (W) the data area can be written to;
ALIGNED (A) the data and strides are aligned appropriately for the hardware (as determined by the compiler);

UPDATEIFCOPY (U) this array is a copy of some other array (referenced by .base). When this array is deallocated, the base array will be updated with the contents of this array.

All flags can be accessed using their first (upper case) letter as well as the full name.

```
>>> y
array([[3, 1, 7],
    [2, 0, 0],
    [8, 5, 9]])
>>> y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : True
    ALIGNED : True
    UPDATEIFCOPY : False
>>> y.setflags(write=0, align=0)
>>> y.flags
```

(continued from previous page)

```
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : False
    ALIGNED : False
    UPDATEIFCOPY : False
>>> y.setflags(uic=1)
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: cannot set UPDATEIFCOPY flag to True
```


## sort

```
IfArray.sort (axis=-1, kind='quicksort',order=None)
```

Sort an array, in-place.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'quicksort', 'mergesort', 'heapsort'\}, optional] Sorting algorithm. Default is 'quicksort'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.sort : Return a sorted copy of an array. argsort : Indirect sort. lexsort : Indirect stable sort on multiple keys. searchsorted : Find elements in sorted array. partition: Partial sort.

See sort for notes on the different sorting algorithms.

```
>>> a = np.array([[1,4], [3,1]])
>>> a.sort(axis=1)
>>> a
array([[1, 4],
    [1, 3]])
>>> a.sort(axis=0)
>>> a
array([[1, 3],
    [1, 4]])
```

Use the order keyword to specify a field to use when sorting a structured array:

```
>>> a = np.array([('a', 2), ('c', 1)], dtype=[('x', 'S1'), ('y', int)])
>>> a.sort(order='y')
>>> a
array([('c', 1), ('a', 2)],
    dtype=[('x', '|S1'), ('y', '<i4')])
```


## squeeze

## IfArray.squeeze (axis=None)

Remove single-dimensional entries from the shape of $a$.
Refer to numpy.squeeze for full documentation.
numpy.squeeze : equivalent function
std

IfArray.std (axis=None, dtype=None, out=None, ddof=0, keepdims=False)
Returns the standard deviation of the array elements along given axis.
Refer to numpy.std for full documentation.
numpy.std : equivalent function
sum

IfArray. sum (axis=None, dtype=None, out=None, keepdims=False)
Return the sum of the array elements over the given axis.
Refer to numpy.sum for full documentation.
numpy.sum : equivalent function

## swapaxes

IfArray.swapaxes (axis1, axis2)
Return a view of the array with axis1 and axis2 interchanged.
Refer to numpy.swapaxes for full documentation.
numpy.swapaxes : equivalent function
take

IfArray.take (indices, axis=None, out=None, mode='raise')
Return an array formed from the elements of $a$ at the given indices.
Refer to numpy.take for full documentation.
numpy.take : equivalent function

## tobytes

IfArray.tobytes (order=' ${ }^{\prime}$ ')
Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.

New in version 1.9.0.
order [ $\{$ ' $C$ ', ' $F$ ', None \}, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
s [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01 \x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01 \x00\x00\x00\x03\x00\x00\x00'
```


## tofile

IfArray.tofile (fid, sep="", format="\%s")
Write array to a file as text or binary (default).
Data is always written in ' C ' order, independent of the order of $a$. The data produced by this method can be recovered using the function fromfile().
fid [file or str] An open file object, or a string containing a filename.
sep [str] Separator between array items for text output. If "" (empty), a binary file is written, equivalent to file.write(a.tobytes()).
format [str] Format string for text file output. Each entry in the array is formatted to text by first converting it to the closest Python type, and then using "format" \% item.

This is a convenience function for quick storage of array data. Information on endianness and precision is lost, so this method is not a good choice for files intended to archive data or transport data between machines with different endianness. Some of these problems can be overcome by outputting the data as text files, at the expense of speed and file size.

## tolist

IfArray.tolist()
Return the array as a (possibly nested) list.
Return a copy of the array data as a (nested) Python list. Data items are converted to the nearest compatible Python type.
none
$\mathbf{y}$ [list] The possibly nested list of array elements.
The array may be recreated, $a=n p . a r r a y(a . t o l i s t())$.

```
>>> a = np.array([1, 2])
>>> a.tolist()
[1, 2]
>>> a = np.array([[1, 2], [3, 4]])
>>> list(a)
[array([1, 2]), array([3, 4])]
>>> a.tolist()
[[1, 2], [3, 4]]
```

tostring

IfArray.tostring (order='C')
Construct Python bytes containing the raw data bytes in the array.

Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.
This function is a compatibility alias for tobytes. Despite its name it returns bytes not strings.
order [ $\{$ ' $C$ ', ' $F$ ', None $\}$, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01 \x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01\x00\x00\x00\x03\x00\x00\x00'
```

trace

```
IfArray.trace (offset=0, axis 1=0, axis 2=1,dtype=None, out=None)
```

Return the sum along diagonals of the array.
Refer to numpy.trace for full documentation.
numpy.trace : equivalent function

## transpose

IfArray.transpose (*axes)
Returns a view of the array with axes transposed.
For a 1-D array, this has no effect. (To change between column and row vectors, first cast the 1-D array into a matrix object.) For a 2-D array, this is the usual matrix transpose. For an n-D array, if axes are given, their order indicates how the axes are permuted (see Examples). If axes are not provided anda.shape $=(i[0], i[1], \ldots i[n-2], i[n-1])$, then a.transpose(). shape $=$ (i[n-1], i[n-2], ... i[1], i[0]).
axes: None, tuple of ints, or $n$ ints

- None or no argument: reverses the order of the axes.
- tuple of ints: $i$ in the $j$-th place in the tuple means $a$ 's $i$-th axis becomes a.transpose()'s $j$-th axis.
- $n$ ints: same as an n-tuple of the same ints (this form is intended simply as a "convenience" alternative to the tuple form)
out [ndarray] View of $a$, with axes suitably permuted.
ndarray.T : Array property returning the array transposed.

```
>>> a = np.array([[1, 2], [3, 4]])
>>> a
array([[1, 2],
    [3, 4]])
```

```
>>> a.transpose()
array([[1, 3],
    [2, 4]])
>>> a.transpose((1, 0))
array([[1, 3],
    [2, 4]])
>>> a.transpose(1, 0)
array([[1, 3],
    [2, 4]])
```

var

IfArray.var (axis=None, dtype=None, out=None, ddof=0, keepdims=False)
Returns the variance of the array elements, along given axis.
Refer to numpy.var for full documentation.
numpy.var : equivalent function

## view

IfArray.view (dtype=None, type=None)
New view of array with the same data.
dtype [data-type or ndarray sub-class, optional] Data-type descriptor of the returned view, e.g., float32 or int16. The default, None, results in the view having the same data-type as $a$. This argument can also be specified as an ndarray sub-class, which then specifies the type of the returned object (this is equivalent to setting the type parameter).
type [Python type, optional] Type of the returned view, e.g., ndarray or matrix. Again, the default None results in type preservation.
a.view () is used two different ways:
a.view (some_dtype) or a.view (dtype=some_dtype) constructs a view of the array's memory with a different data-type. This can cause a reinterpretation of the bytes of memory.
a.view(ndarray_subclass) or a.view(type=ndarray_subclass) just returns an instance of ndarray_subclass that looks at the same array (same shape, dtype, etc.) This does not cause a reinterpretation of the memory.
For a.view (some_dtype), if some_dtype has a different number of bytes per entry than the previous dtype (for example, converting a regular array to a structured array), then the behavior of the view cannot be predicted just from the superficial appearance of a (shown by print (a)). It also depends on exactly how a is stored in memory. Therefore if a is C-ordered versus fortran-ordered, versus defined as a slice or transpose, etc., the view may give different results.

```
>>> x = np.array([(1, 2)], dtype=[('a', np.int8), ('b', np.int8)])
```

Viewing array data using a different type and dtype:

```
>>> y = x.view(dtype=np.int16, type=np.matrix)
>>> Y
matrix([[513]], dtype=int16)
```

```
>>> print(type(y))
<class 'numpy.matrixlib.defmatrix.matrix'>
```

Creating a view on a structured array so it can be used in calculations

```
>>> x = np.array([(1, 2), (3,4)], dtype=[('a', np.int8), ('b', np.int8)])
>>> xv = x.view(dtype=np.int8).reshape(-1,2)
>>> xv
array([[1, 2],
    [3, 4]], dtype=int8)
>>> xv.mean(0)
array([ 2., 3.])
```

Making changes to the view changes the underlying array

```
>>> xv[0,1] = 20
>>> print(x)
[(1, 20) (3, 4)]
```

Using a view to convert an array to a recarray:

```
>>> z = x.view(np.recarray)
>>> z.a
array([1], dtype=int8)
```

Views share data:

```
>>> x[0] = (9, 10)
>>> z[0]
(9, 10)
```

Views that change the dtype size (bytes per entry) should normally be avoided on arrays defined by slices, transposes, fortran-ordering, etc.:

```
>>> x = np.array([[1,2,3],[4,5,6]], dtype=np.int16)
>>> y = x[:, 0:2]
>>> y
array([[1, 2],
    [4, 5]], dtype=int16)
>>> y.view(dtype=[('width', np.int16), ('length', np.int16)])
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: new type not compatible with array.
>>> z = y.copy()
>>> z.view(dtype=[('width', np.int16), ('length', np.int16)])
array([[(1, 2)],
    [(4, 5)]], dtype=[('width', '<i2'), ('length', '<i2')])
```

Initialize self. See help(type(self)) for accurate signature.

## Attributes

| T | Same as self.transpose(), except that self is returned <br> if self.ndim < 2. |
| :--- | :--- |
| base | Base object if memory is from some other object. |
| ctypes | An object to simplify the interaction of the array with <br> the ctypes module. |
| data | Python buffer object pointing to the start of the ar- <br> ray's data. |
| dtype | Data-type of the array's elements. |
| flags | Information about the memory layout of the array. |
| flat | A 1-D iterator over the array. |
| imag | The imaginary part of the array. |
| itemsize | Length of one array element in bytes. |
| nbytes | Total bytes consumed by the elements of the array. |
| ndim | Number of array dimensions. |
| real | The real part of the array. |
| shape | Tuple of array dimensions. |
| size | Number of elements in the array. |
| strides | Tuple of bytes to step in each dimension when <br> traversing an array. |

T

IfArray. T
Same as self.transpose(), except that self is returned if self.ndim $<2$.

```
>>> x = np.array([[1.,2.],[3.,4.]])
>>> x
array([[ 1., 2.],
    [ 3., 4.]])
>>> x.T
array([[ 1., 3.],
    [ 2., 4.]])
>>> x = np.array([1.,2.,3.,4.])
>>> x
array([ 1., 2., 3., 4.])
>>> x.T
array([ 1., 2., 3., 4.])
```

base

IfArray.base
Base object if memory is from some other object.
The base of an array that owns its memory is None:

```
>>> x = np.array([1,2,3,4])
>>> x.base is None
True
```

Slicing creates a view, whose memory is shared with x :

```
>>> y = x[2:]
>>> y.base is x
True
```

ctypes

## IfArray.ctypes

An object to simplify the interaction of the array with the ctypes module.
This attribute creates an object that makes it easier to use arrays when calling shared libraries with the ctypes module. The returned object has, among others, data, shape, and strides attributes (see Notes below) which themselves return ctypes objects that can be used as arguments to a shared library.

## None

c [Python object] Possessing attributes data, shape, strides, etc.
numpy.ctypeslib
Below are the public attributes of this object which were documented in "Guide to NumPy" (we have omitted undocumented public attributes, as well as documented private attributes):

- data: A pointer to the memory area of the array as a Python integer. This memory area may contain data that is not aligned, or not in correct byte-order. The memory area may not even be writeable. The array flags and data-type of this array should be respected when passing this attribute to arbitrary C-code to avoid trouble that can include Python crashing. User Beware! The value of this attribute is exactly the same as self._array_interface_['data'][0].
- shape (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the C-integer corresponding to dtype( ' p ') on this platform. This base-type could be c_int, c_long, or c_longlong depending on the platform. The c_intp type is defined accordingly in numpy.ctypeslib. The ctypes array contains the shape of the underlying array.
- strides (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the same as for the shape attribute. This ctypes array contains the strides information from the underlying array. This strides information is important for showing how many bytes must be jumped to get to the next element in the array.
- data_as(obj): Return the data pointer cast to a particular c-types object. For example, calling self._as_parameter_ is equivalent to self.data_as(ctypes.c_void_p). Perhaps you want to use the data as a pointer to a ctypes array of floating-point data: self.data_as(ctypes.POINTER(ctypes.c_double)).
- shape_as(obj): Return the shape tuple as an array of some other c-types type. For example: self.shape_as(ctypes.c_short).
- strides_as(obj): Return the strides tuple as an array of some other c-types type. For example: self.strides_as(ctypes.c_longlong).

Be careful using the ctypes attribute - especially on temporary arrays or arrays constructed on the fly. For example, calling ( $a+b$ ) . ctypes.data_as (ctypes.c_void_p) returns a pointer to memory that is invalid because the array created as $(a+b)$ is deallocated before the next Python statement. You can avoid this problem using either $c=a+b$ or $c t=(a+b) . c t y p e s$. In the latter case, ct will hold a reference to the array until ct is deleted or re-assigned.

If the ctypes module is not available, then the ctypes attribute of array objects still returns something useful, but ctypes objects are not returned and errors may be raised instead. In particular, the object will still have the as parameter attribute which will return an integer equal to the data attribute.

```
>>> import ctypes
>>> x
array([[0, 1],
    [2, 3]])
>>> x.ctypes.data
30439712
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long))
<ctypes.LP_c_long object at 0x01F01300>
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long)).contents
c_long(0)
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_longlong)).contents
c_longlong(4294967296L)
>>> x.ctypes.shape
<numpy.core._internal.c_long_Array_2 object at 0x01FFD580>
>>> x.ctypes.shape_as(ctypes.c_long)
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides_as(ctypes.c_longlong)
<numpy.core._internal.c_longlong_Array_2 object at 0x01F01300>
```

data

## IfArray.data

Python buffer object pointing to the start of the array's data.
dtype

## IfArray.dtype

Data-type of the array's elements.
None
d: numpy dtype object
numpy.dtype

```
>>> x
array([[0, 1],
    [2, 3]])
>>> x.dtype
dtype('int32')
>>> type(x.dtype)
<type 'numpy.dtype'>
```


## flags

## IfArray.flags

Information about the memory layout of the array.
C_CONTIGUOUS (C) The data is in a single, C-style contiguous segment.
F_CONTIGUOUS (F) The data is in a single, Fortran-style contiguous segment.
OWNDATA (O) The array owns the memory it uses or borrows it from another object.

WRITEABLE (W) The data area can be written to. Setting this to False locks the data, making it readonly. A view (slice, etc.) inherits WRITEABLE from its base array at creation time, but a view of a writeable array may be subsequently locked while the base array remains writeable. (The opposite is not true, in that a view of a locked array may not be made writeable. However, currently, locking a base object does not lock any views that already reference it, so under that circumstance it is possible to alter the contents of a locked array via a previously created writeable view onto it.) Attempting to change a non-writeable array raises a RuntimeError exception.

ALIGNED (A) The data and all elements are aligned appropriately for the hardware.
UPDATEIFCOPY (U) This array is a copy of some other array. When this array is deallocated, the base array will be updated with the contents of this array.

FNC F_CONTIGUOUS and not C_CONTIGUOUS.
FORC F_CONTIGUOUS or C_CONTIGUOUS (one-segment test).

## BEHAVED (B) ALIGNED and WRITEABLE.

CARRAY (CA) BEHAVED and C_CONTIGUOUS.
FARRAY (FA) BEHAVED and F_CONTIGUOUS and not C_CONTIGUOUS.
The flags object can be accessed dictionary-like (as in a.flags ['WRITEABLE']), or by using lowercased attribute names (as in a.flags.writeable). Short flag names are only supported in dictionary access.

Only the UPDATEIFCOPY, WRITEABLE, and ALIGNED flags can be changed by the user, via direct assignment to the attribute or dictionary entry, or by calling ndarray.setflags.

The array flags cannot be set arbitrarily:

- UPDATEIFCOPY can only be set False.
- ALIGNED can only be set True if the data is truly aligned.
- WRITEABLE can only be set True if the array owns its own memory or the ultimate owner of the memory exposes a writeable buffer interface or is a string.

Arrays can be both C-style and Fortran-style contiguous simultaneously. This is clear for 1-dimensional arrays, but can also be true for higher dimensional arrays.

Even for contiguous arrays a stride for a given dimension arr.strides [dim] may be arbitrary if arr.shape[dim] == 1 or the array has no elements. It does not generally hold that self. strides [-1] == self.itemsize for C-style contiguous arrays or self.strides[0] == self.itemsize for Fortran-style contiguous arrays is true.
flat

## IfArray.flat

A 1-D iterator over the array.
This is a numpy.flatiter instance, which acts similarly to, but is not a subclass of, Python's built-in iterator object.
flatten : Return a copy of the array collapsed into one dimension.
flatiter

```
>>> x = np.arange(1, 7).reshape(2, 3)
>>> x
array([[1, 2, 3],
    [4, 5, 6]])
>>> x.flat[3]
4
>>> x.T
array([[1, 4],
    [2, 5],
    [3, 6]])
>>> x.T.flat[3]
5
>>> type(x.flat)
<type 'numpy.flatiter'>
```

An assignment example:

```
>>> x.flat = 3; x
array([[3, 3, 3],
    [3, 3, 3]])
>>> x.flat[[1,4]] = 1; x
array([[3, 1, 3],
    [3, 1, 3]])
```

imag

## IfArray.imag

The imaginary part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.imag
array([ 0. , 0.70710678])
>>> x.imag.dtype
dtype('float64')
```


## itemsize

IfArray.itemsize
Length of one array element in bytes.

```
>>> x = np.array([1,2,3], dtype=np.float64)
>>> x.itemsize
8
>>> x = np.array([1,2,3], dtype=np.complex128)
>>> x.itemsize
16
```


## nbytes

IfArray. nbytes
Total bytes consumed by the elements of the array.
Does not include memory consumed by non-element attributes of the array object.

```
>>> x = np.zeros((3,5,2), dtype=np.complex128)
>>> x.nbytes
480
>>> np.prod(x.shape) * x.itemsize
480
```

ndim

IfArray.ndim
Number of array dimensions.

```
>>> x = np.array([1, 2, 3])
>>> x.ndim
1
>>> y = np.zeros((2, 3, 4))
>>> y.ndim
3
```

real

IfArray.real
The real part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.real
array([ 1. , 0.70710678])
>>> x.real.dtype
dtype('float64')
```

numpy.real : equivalent function
shape

IfArray.shape
Tuple of array dimensions.
May be used to "reshape" the array, as long as this would not require a change in the total number of elements

```
>>> x = np.array([1, 2, 3, 4])
>>> x.shape
(4,)
>>> y = np.zeros((2, 3, 4))
>>> y.shape
(2, 3, 4)
>>> y.shape = (3, 8)
>>> y
array([[[ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.]])
>>> y.shape = (3, 6)
Traceback (most recent call last):
```

```
    File "<stdin>", line 1, in <module>
ValueError: total size of new array must be unchanged
```


## size

IfArray.size
Number of elements in the array.
Equivalent to np.prod (a.shape), i.e., the product of the array's dimensions.

```
>>> x = np.zeros((3, 5, 2), dtype=np.complex128)
>>> x.size
30
>>> np.prod(x.shape)
30
```


## strides

## IfArray.strides

Tuple of bytes to step in each dimension when traversing an array.
The byte offset of element (i[0], $i[1], \ldots, i[n]$ ) in an array $a$ is:

```
offset = sum(np.array(i) * a.strides)
```

A more detailed explanation of strides can be found in the "ndarray.rst" file in the NumPy reference guide.
Imagine an array of 32-bit integers (each 4 bytes):
$x=n p \cdot \operatorname{array}([[0,1,2,3,4]$,
$[5,6,7,8,9]]$, dtype=np.int 32)
This array is stored in memory as 40 bytes, one after the other (known as a contiguous block of memory). The strides of an array tell us how many bytes we have to skip in memory to move to the next position along a certain axis. For example, we have to skip 4 bytes ( 1 value) to move to the next column, but 20 bytes ( 5 values) to get to the same position in the next row. As such, the strides for the array $x$ will be $(20,4)$.
numpy.lib.stride_tricks.as_strided

```
>>> y = np.reshape(np.arange (2*3*4), (2,3,4))
>>> y
array([[[ 0, 1, 2, 3],
    [4, 5, 6, 7],
    [ 8, 9, 10, 11]],
    [[12, 13, 14, 15],
    [16, 17, 18, 19],
    [20, 21, 22, 23]]])
>>> y.strides
(48, 16, 4)
>>> y[1,1,1]
17
>>> offset=sum(y.strides * np.array((1,1,1)))
```

```
>>> offset/y.itemsize
```

17

```
>>> x = np.reshape(np.arange(5*6*7*8), (5, 6,7,8)).transpose (2, 3, 1,0)
>>> x.strides
(32, 4, 224, 1344)
>>> i = np.array ([3,5,2,2])
>>> offset = sum(i * x.strides)
>>> x[3,5,2,2]
813
>>> offset / x.itemsize
813
```


## IfErrorArray

class IfErrorArray

## Methods

| all | Returns True if all elements evaluate to True. |
| :---: | :---: |
| any | Returns True if any of the elements of $a$ evaluate to True. |
| argmax | Return indices of the maximum values along the given axis. |
| argmin | Return indices of the minimum values along the given axis of $a$. |
| argpartition | Returns the indices that would partition this array. |
| argsort | Returns the indices that would sort this array. |
| astype | Copy of the array, cast to a specified type. |
| byteswap | Swap the bytes of the array elements |
| choose | Use an index array to construct a new array from a set of choices. |
| clip | Return an array whose values are limited to [min, max]. |
| collapse |  |
| compress | Return selected slices of this array along given axis. |
| conj | Complex-conjugate all elements. |
| conjugate | Return the complex conjugate, element-wise. |
| copy | Return a copy of the array. |
| cumprod | Return the cumulative product of the elements along the given axis. |
| cumsum | Return the cumulative sum of the elements along the given axis. |
| diagonal | Return specified diagonals. |
| dot | Dot product of two arrays. |
| dump | Dump a pickle of the array to the specified file. |
| dumps | Returns the pickle of the array as a string. |
| fill | Fill the array with a scalar value. |

Continued on next page

Table 55 - continued from previous page

| flatten | Return a copy of the array collapsed into one dimension. |
| :---: | :---: |
| getfield | Returns a field of the given array as a certain type. |
| item | Copy an element of an array to a standard Python scalar and return it. |
| itemset | Insert scalar into an array (scalar is cast to array's dtype, if possible) |
| max | Return the maximum along a given axis. |
| mean | Returns the average of the array elements along given axis. |
| min | Return the minimum along a given axis. |
| newbyteorder | Return the array with the same data viewed with a different byte order. |
| nonzero | Return the indices of the elements that are non-zero. |
| partition | Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. |
| prod | Return the product of the array elements over the given axis |
| ptp | Peak to peak (maximum - minimum) value along a given axis. |
| put | Set a.flat $[\mathrm{n}]=$ values $[\mathrm{n}]$ for all $n$ in indices. |
| ravel | Return a flattened array. |
| repeat | Repeat elements of an array. |
| reshape | Returns an array containing the same data with a new shape. |
| resize | Change shape and size of array in-place. |
| round | Return $a$ with each element rounded to the given number of decimals. |
| searchsorted | Find indices where elements of v should be inserted in a to maintain order. |
| setfield | Put a value into a specified place in a field defined by a data-type. |
| setflags | Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively. |
| sort | Sort an array, in-place. |
| squeeze | Remove single-dimensional entries from the shape of $a$. |
| std | Returns the standard deviation of the array elements along given axis. |
| sum | Return the sum of the array elements over the given axis. |
| swapaxes | Return a view of the array with axis1 and axis2 interchanged. |
| take | Return an array formed from the elements of $a$ at the given indices. |
| tobytes | Construct Python bytes containing the raw data bytes in the array. |
| tofile | Write array to a file as text or binary (default). |
| tolist | Return the array as a (possibly nested) list. |

Table 55 - continued from previous page

| tostring | Construct Python bytes containing the raw data bytes <br> in the array. |
| :--- | :--- |
| trace | Return the sum along diagonals of the array. |
| transpose | Returns a view of the array with axes transposed. |
| var | Returns the variance of the array elements, along <br> given axis. |
| view | New view of array with the same data. |

all
IfErrorArray.all (axis=None, out=None, keepdims=False)
Returns True if all elements evaluate to True.
Refer to numpy.all for full documentation.
numpy.all : equivalent function
any
IfErrorArray.any (axis=None, out=None, keepdims=False)
Returns True if any of the elements of $a$ evaluate to True.
Refer to numpy.any for full documentation.
numpy.any : equivalent function
argmax

IfErrorArray.argmax (axis=None, out=None)
Return indices of the maximum values along the given axis.
Refer to numpy.argmax for full documentation.
numpy.argmax : equivalent function

## argmin

IfErrorArray.argmin (axis=None, out=None)
Return indices of the minimum values along the given axis of $a$.
Refer to numpy.argmin for detailed documentation.
numpy.argmin : equivalent function

## argpartition

IfErrorArray.argpartition (kth, axis=-1, kind='introselect', order=None)
Returns the indices that would partition this array.
Refer to numpy.argpartition for full documentation.
New in version 1.8.0.
numpy.argpartition : equivalent function
argsort

IfErrorArray. argsort (axis=-1, kind='quicksort', order=None)
Returns the indices that would sort this array.
Refer to numpy.argsort for full documentation.
numpy.argsort : equivalent function
astype

IfErrorArray. astype (dtype, order='K', casting='unsafe', subok=True, copy=True)
Copy of the array, cast to a specified type.
dtype [str or dtype] Typecode or data-type to which the array is cast.
order [\{ 'C', ' F ', ' A ', ' K '\}, optional] Controls the memory layout order of the result. ' C ' means C order, ' F ' means Fortran order, ' A ' means ' F ' order if all the arrays are Fortran contiguous, ' C ' order otherwise, and ' K ' means as close to the order the array elements appear in memory as possible. Default is ' K '.
casting [ $\{$ 'no', 'equiv', 'safe', 'same_kind', 'unsafe' \}, optional] Controls what kind of data casting may occur. Defaults to 'unsafe' for backwards compatibility.

- 'no' means the data types should not be cast at all.
- 'equiv' means only byte-order changes are allowed.
- 'safe' means only casts which can preserve values are allowed.
- 'same_kind' means only safe casts or casts within a kind, like float64 to float32, are allowed.
- 'unsafe' means any data conversions may be done.
subok [bool, optional] If True, then sub-classes will be passed-through (default), otherwise the returned array will be forced to be a base-class array.
copy [bool, optional] By default, astype always returns a newly allocated array. If this is set to false, and the dtype, order, and subok requirements are satisfied, the input array is returned instead of a copy.
arr_t [ndarray] Unless copy is False and the other conditions for returning the input array are satisfied (see description for copy input parameter), arr_t is a new array of the same shape as the input array, with dtype, order given by dtype, order.

Starting in NumPy 1.9, astype method now returns an error if the string dtype to cast to is not long enough in 'safe' casting mode to hold the max value of integer/float array that is being casted. Previously the casting was allowed even if the result was truncated.
ComplexWarning When casting from complex to float or int. To avoid this, one should use a.real. astype ( t ).

```
>>> x = np.array([1, 2, 2.5])
>>> x
array([ 1., 2. , 2.5])
```

```
>>> x.astype(int)
array([1, 2, 2])
```


## byteswap

## IfErrorArray.byteswap (inplace)

Swap the bytes of the array elements
Toggle between low-endian and big-endian data representation by returning a byteswapped array, optionally swapped in-place.
inplace [bool, optional] If True, swap bytes in-place, default is False.
out [ndarray] The byteswapped array. If inplace is True, this is a view to self.

```
>>> A = np.array([1, 256, 8755], dtype=np.int16)
>>> map(hex, A)
['0x1', '0x100', '0x2233']
>>> A.byteswap(True)
array([ 256, 1, 13090], dtype=int16)
>>> map (hex, A)
['0x100', '0x1', '0x3322']
```

Arrays of strings are not swapped

```
>>> A = np.array(['ceg', 'fac'])
>>> A.byteswap()
array(['ceg', 'fac'],
    dtype='|S3')
```

choose
IfErrorArray. choose (choices, out=None, mode='raise')

Use an index array to construct a new array from a set of choices.
Refer to numpy.choose for full documentation.
numpy.choose : equivalent function
clip

IfErrorArray.clip (min=None, max=None, out=None)
Return an array whose values are limited to [min, max]. One of max or min must be given.
Refer to numpy.clip for full documentation.
numpy.clip : equivalent function
collapse

IfErrorArray.collapse (shape)
compress

IfErrorArray. compress (condition, axis=None, out=None)
Return selected slices of this array along given axis.

Refer to numpy.compress for full documentation.
numpy.compress : equivalent function
conj

IfErrorArray.conj()
Complex-conjugate all elements.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function

## conjugate

IfErrorArray.conjugate()
Return the complex conjugate, element-wise.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function
copy

IfErrorArray.copy (order='C')
Return a copy of the array.
order [\{ 'C', ' F ', ' A ', ' K '\}, optional] Controls the memory layout of the copy. ' C ' means C -order, ' F ' means F -order, ' A ' means ' F ' if $a$ is Fortran contiguous, ' C ' otherwise. ' K ' means match the layout of $a$ as closely as possible. (Note that this function and :func:numpy.copy are very similar, but have different default values for their order= arguments.)
numpy.copy numpy.copyto
$\ggg x=n p . \operatorname{array}([[1,2,3],[4,5,6]]$, order='F')
>>> y $=x . \operatorname{copy}()$

```
>>> x.fill(0)
```

```
>>> x
array([[0, 0, 0],
    [0, 0, 0]])
```

```
>>> y
array([[1, 2, 3],
    [4, 5, 6]])
```

>>> y.flags['C_CONTIGUOUS']
True
cumprod

IfErrorArray. Cumprod (axis=None, dtype=None, out=None)
Return the cumulative product of the elements along the given axis.
Refer to numpy.cumprod for full documentation.
numpy.cumprod : equivalent function

## cumsum

IfErrorArray. cumsum (axis=None, dtype=None, out=None)
Return the cumulative sum of the elements along the given axis.
Refer to numpy.cumsum for full documentation.
numpy.cumsum : equivalent function

## diagonal

IfErrorArray.diagonal (offset $=0$, axis $1=0$, axis $2=1$ )
Return specified diagonals. In NumPy 1.9 the returned array is a read-only view instead of a copy as in previous NumPy versions. In a future version the read-only restriction will be removed.
Refer to numpy. diagonal () for full documentation.
numpy.diagonal : equivalent function
dot

IfErrorArray.dot ( $b$, out=None)
Dot product of two arrays.
Refer to numpy.dot for full documentation.
numpy.dot : equivalent function

```
>>> a = np.eye(2)
>>> b = np.ones((2, 2)) * 2
>>> a.dot(b)
array([[ 2., 2.],
    [ 2., 2.]])
```

This array method can be conveniently chained:

```
>>> a.dot(b).dot(b)
array([[ 8., 8.],
    [ 8., 8.]])
```

dump

IfErrorArray. dump (file)
Dump a pickle of the array to the specified file. The array can be read back with pickle.load or numpy.load.
file [ str ] A string naming the dump file.

## dumps

IfErrorArray.dumps()
Returns the pickle of the array as a string. pickle.loads or numpy.loads will convert the string back to an array.

None
fill

IfErrorArray.fill (value)
Fill the array with a scalar value.
value [scalar] All elements of $a$ will be assigned this value.

```
>>> a = np.array([1, 2])
>>> a.fill(0)
>>> a
array([0, 0])
>>> a = np.empty(2)
>>> a.fill(1)
>>> a
array([ 1., 1.])
```

flatten

IfErrorArray.flatten (order $=$ ' $C^{\prime}$ )
Return a copy of the array collapsed into one dimension.
order [ $\{$ ' C ', ' F ', ' $A$ ', ' K '\}, optional] ' C ' means to flatten in row-major ( C -style) order. ' F ' means to flatten in column-major (Fortran- style) order. 'A' means to flatten in column-major order if $a$ is Fortran contiguous in memory, row-major order otherwise. ' K ' means to flatten $a$ in the order the elements occur in memory. The default is ' C '.
$\mathbf{y}$ [ndarray] A copy of the input array, flattened to one dimension.
ravel : Return a flattened array. flat : A 1-D flat iterator over the array.

```
>>> a = np.array([[1,2], [3,4]])
>>> a.flatten()
array([1, 2, 3, 4])
>>> a.flatten('F')
array([1, 3, 2, 4])
```


## getfield

IfErrorArray.getfield (dtype, offset=0)
Returns a field of the given array as a certain type.
A field is a view of the array data with a given data-type. The values in the view are determined by the given type and the offset into the current array in bytes. The offset needs to be such that the view dtype fits in the array dtype; for example an array of dtype complex 128 has 16-byte elements. If taking a view with a 32-bit integer ( 4 bytes), the offset needs to be between 0 and 12 bytes.
dtype [str or dtype] The data type of the view. The dtype size of the view can not be larger than that of the array itself.
offset [int] Number of bytes to skip before beginning the element view.

```
>>> x = np.diag([1.+1.j]*2)
>>> x[1, 1] = 2 + 4.j
>>> x
array([[ 1.+1.j, 0.+0.j],
    [ 0.+0.j, 2.+4.j]])
>>> x.getfield(np.float64)
array([[ 1., 0.],
    [0., 2.]])
```

By choosing an offset of 8 bytes we can select the complex part of the array for our view:

```
>>> x.getfield(np.float64, offset=8)
array([[ 1., 0.],
    [ 0., 4.]])
```


## item

IfErrorArray.item (*args)
Copy an element of an array to a standard Python scalar and return it.
*args : Arguments (variable number and type)

- none: in this case, the method only works for arrays with one element (a.size $==1$ ), which element is copied into a standard Python scalar object and returned.
- int_type: this argument is interpreted as a flat index into the array, specifying which element to copy and return.
- tuple of int_types: functions as does a single int_type argument, except that the argument is interpreted as an nd-index into the array.
$\mathbf{z}$ [Standard Python scalar object] A copy of the specified element of the array as a suitable Python scalar
When the data type of $a$ is longdouble or clongdouble, item() returns a scalar array object because there is no available Python scalar that would not lose information. Void arrays return a buffer object for item(), unless fields are defined, in which case a tuple is returned.
item is very similar to a[args], except, instead of an array scalar, a standard Python scalar is returned. This can be useful for speeding up access to elements of the array and doing arithmetic on elements of the array using Python's optimized math.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.item(3)
2
>>> x.item(7)
5
>>> x.item((0, 1))
1
```

```
>>> x.item((2, 2))
```

3

## itemset

## IfErrorArray.itemset (*args)

Insert scalar into an array (scalar is cast to array's dtype, if possible)
There must be at least 1 argument, and define the last argument as item. Then, a.itemset (*args) is equivalent to but faster than a [args] = item. The item should be a scalar value and args must select a single item in the array $a$.

1*args [Arguments] If one argument: a scalar, only used in case $a$ is of size 1. If two arguments: the last argument is the value to be set and must be a scalar, the first argument specifies a single array element location. It is either an int or a tuple.

Compared to indexing syntax, itemset provides some speed increase for placing a scalar into a particular location in an ndarray, if you must do this. However, generally this is discouraged: among other problems, it complicates the appearance of the code. Also, when using itemset (and item) inside a loop, be sure to assign the methods to a local variable to avoid the attribute look-up at each loop iteration.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.itemset (4, 0)
>>> x.itemset((2, 2), 9)
>>> x
array([[3, 1, 7],
    [2, 0, 3],
    [8, 5, 9]])
```

max

IfErrorArray.max (axis=None, out=None)
Return the maximum along a given axis.
Refer to numpy.amax for full documentation.
numpy.amax : equivalent function
mean

IfErrorArray.mean (axis=None, dtype=None, out=None, keepdims=False)
Returns the average of the array elements along given axis.
Refer to numpy.mean for full documentation.
numpy.mean : equivalent function
$\min$
IfErrorArray.min (axis=None, out=None, keepdims=False)
Return the minimum along a given axis.
Refer to numpy.amin for full documentation.
numpy.amin : equivalent function

## newbyteorder

IfErrorArray. newbyteorder (new_order='S')
Return the array with the same data viewed with a different byte order.
Equivalent to:

```
arr.view(arr.dtype.newbytorder(new_order))
```

Changes are also made in all fields and sub-arrays of the array data type.
new_order [string, optional] Byte order to force; a value from the byte order specifications below. new_order codes can be any of:

- 'S' - swap dtype from current to opposite endian
- \{‘‘’, 'L’\} - little endian
- $\{‘>’, ~ ‘ B ’\}$ - big endian
- $\{$ ' $=$ ', ' N ' $\}$ - native order
- $\{$ ' 1 ', ' I ' $\}$ - ignore (no change to byte order)

The default value ('S') results in swapping the current byte order. The code does a case-insensitive check on the first letter of new_order for the alternatives above. For example, any of ' $B$ ' or ' $b$ ' or 'biggish' are valid to specify big-endian.
new_arr [array] New array object with the dtype reflecting given change to the byte order.

## nonzero

IfErrorArray.nonzero()
Return the indices of the elements that are non-zero.
Refer to numpy.nonzero for full documentation.
numpy.nonzero : equivalent function
partition

IfErrorArray.partition (kth, axis=-1, kind='introselect', order=None)
Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. All elements smaller than the kth element are moved before this element and all equal or greater are moved behind it. The ordering of the elements in the two partitions is undefined.

New in version 1.8.0.
$\mathbf{k t h}$ [int or sequence of ints] Element index to partition by. The kth element value will be in its final sorted position and all smaller elements will be moved before it and all equal or greater elements behind it. The order all elements in the partitions is undefined. If provided with a sequence of kth it will partition all elements indexed by kth of them into their sorted position at once.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'introselect'\}, optional] Selection algorithm. Default is 'introselect'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.partition : Return a parititioned copy of an array. argpartition : Indirect partition. sort : Full sort.
See np.partition for notes on the different algorithms.

```
>>> a = np.array([3, 4, 2, 1])
>>> a.partition(a, 3)
>>> a
array([2, 1, 3, 4])
```

```
>>> a.partition((1, 3))
array([1, 2, 3, 4])
```

prod

IfErrorArray.prod (axis=None, dtype=None, out=None, keepdims=False)
Return the product of the array elements over the given axis
Refer to numpy.prod for full documentation.
numpy.prod : equivalent function
ptp

IfErrorArray.ptp (axis=None, out=None)
Peak to peak (maximum - minimum) value along a given axis.
Refer to numpy.ptp for full documentation.
numpy.ptp : equivalent function
put

IfErrorArray.put (indices, values, mode='raise')
Set a.flat $[\mathrm{n}]=$ values [ n$]$ for all $n$ in indices.
Refer to numpy.put for full documentation.
numpy.put : equivalent function
ravel

IfErrorArray. ravel ([order])
Return a flattened array.
Refer to numpy.ravel for full documentation.
numpy.ravel : equivalent function
ndarray.flat : a flat iterator on the array.
repeat

IfErrorArray.repeat (repeats, axis=None)
Repeat elements of an array.
Refer to numpy.repeat for full documentation.
numpy.repeat : equivalent function
reshape

IfErrorArray. reshape (shape, order='C')
Returns an array containing the same data with a new shape.
Refer to numpy.reshape for full documentation.
numpy.reshape : equivalent function
resize

IfErrorArray.resize (new_shape, refcheck=True)
Change shape and size of array in-place.
new_shape [tuple of ints, or $n$ ints] Shape of resized array.
refcheck [bool, optional] If False, reference count will not be checked. Default is True.
None
ValueError If $a$ does not own its own data or references or views to it exist, and the data memory must be changed.

SystemError If the order keyword argument is specified. This behaviour is a bug in NumPy.
resize : Return a new array with the specified shape.
This reallocates space for the data area if necessary.
Only contiguous arrays (data elements consecutive in memory) can be resized.
The purpose of the reference count check is to make sure you do not use this array as a buffer for another Python object and then reallocate the memory. However, reference counts can increase in other ways so if you are sure that you have not shared the memory for this array with another Python object, then you may safely set refcheck to False.
Shrinking an array: array is flattened (in the order that the data are stored in memory), resized, and reshaped:

```
>>> a = np.array([[0, 1], [2, 3]], order='C')
>>> a.resize((2, 1))
>>> a
array([[0],
    [1]])
```

```
>>> a = np.array([[0, 1], [2, 3]], order='F')
>>> a.resize((2, 1))
>>> a
array([[0],
    [2]])
```

Enlarging an array: as above, but missing entries are filled with zeros:

```
>>> b = np.array([[0, 1], [2, 3]])
>>> b.resize(2, 3) # new_shape parameter doesn't have to be a tuple
>>> b
array([[0, 1, 2],
    [3, 0, 0]])
```

Referencing an array prevents resizing...

```
>>> c = a
>>> a.resize((1, 1))
Traceback (most recent call last):
ValueError: cannot resize an array that has been referenced ...
```

Unless refcheck is False:

```
>>> a.resize((1, 1), refcheck=False)
>>> a
array([[0]])
>>> c
array([[0]])
```

round

IfErrorArray.round (decimals=0, out=None)
Return $a$ with each element rounded to the given number of decimals.
Refer to numpy.around for full documentation.
numpy.around : equivalent function
searchsorted

IfErrorArray. searchsorted ( $v$, side $=$ 'left', sorter $=$ None)
Find indices where elements of v should be inserted in a to maintain order.
For full documentation, see numpy.searchsorted
numpy.searchsorted : equivalent function

## setfield

IfErrorArray.setfield(val, dtype, offset=0)
Put a value into a specified place in a field defined by a data-type.
Place val into $a$ 's field defined by dtype and beginning offset bytes into the field.
val [object] Value to be placed in field.
dtype [dtype object] Data-type of the field in which to place val.
offset [int, optional] The number of bytes into the field at which to place val.
None
getfield

```
>>> x = np.eye(3)
>>> x.getfield(np.float64)
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
>>> x.setfield(3, np.int32)
>>> x.getfield(np.int32)
array([[3, 3, 3],
    [3, 3, 3],
    [3, 3, 3]])
>>> x
array([[ 1.00000000e+000, 1.48219694e-323, 1.48219694e-323],
    [ 1.48219694e-323, 1.00000000e+000, 1.48219694e-323],
    [ 1.48219694e-323, 1.48219694e-323, 1.00000000e+000]])
>>> x.setfield(np.eye(3), np.int32)
>>> x
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
```


## setflags

IfErrorArray.setflags (write $=$ None, align=None, uic $=$ None)
Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively.
These Boolean-valued flags affect how numpy interprets the memory area used by $a$ (see Notes below). The ALIGNED flag can only be set to True if the data is actually aligned according to the type. The UPDATEIFCOPY flag can never be set to True. The flag WRITEABLE can only be set to True if the array owns its own memory, or the ultimate owner of the memory exposes a writeable buffer interface, or is a string. (The exception for string is made so that unpickling can be done without copying memory.)
write [bool, optional] Describes whether or not $a$ can be written to.
align [bool, optional] Describes whether or not $a$ is aligned properly for its type.
uic [bool, optional] Describes whether or not $a$ is a copy of another "base" array.
Array flags provide information about how the memory area used for the array is to be interpreted. There are 6 Boolean flags in use, only three of which can be changed by the user: UPDATEIFCOPY, WRITEABLE, and ALIGNED.

WRITEABLE (W) the data area can be written to;

ALIGNED (A) the data and strides are aligned appropriately for the hardware (as determined by the compiler);
UPDATEIFCOPY (U) this array is a copy of some other array (referenced by base). When this array is deallocated, the base array will be updated with the contents of this array.

All flags can be accessed using their first (upper case) letter as well as the full name.

```
>>> y
array([[3, 1, 7],
    [2, 0, 0],
    [8, 5, 9]])
>>> y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : True
    ALIGNED : True
    UPDATEIFCOPY : False
>>> y.setflags(write=0, align=0)
>>> y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : False
    ALIGNED : False
    UPDATEIFCOPY : False
>>> Y.setflags(uic=1)
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: cannot set UPDATEIFCOPY flag to True
```

sort

IfErrorArray.sort (axis=-1, kind='quicksort', order=None)
Sort an array, in-place.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [ \{ 'quicksort', 'mergesort', 'heapsort'\}, optional] Sorting algorithm. Default is 'quicksort'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.sort : Return a sorted copy of an array. argsort : Indirect sort. lexsort : Indirect stable sort on multiple keys. searchsorted : Find elements in sorted array. partition: Partial sort.

See sort for notes on the different sorting algorithms.

```
>>> a = np.array([[1,4], [3,1]])
>>> a.sort(axis=1)
>>> a
array([[1, 4],
    [1, 3]])
>>> a.sort(axis=0)
>>> a
```

(continued from previous page)

```
array([[1, 3],
```

    \([1,4]])\)
    Use the order keyword to specify a field to use when sorting a structured array:

```
>>> a = np.array([('a', 2), ('c', 1)], dtype=[('x', 'S1'), ('y', int)])
>>> a.sort(order='y')
>>> a
array([('c', 1), ('a', 2)],
    dtype=[('x', '|S1'), ('y', '<i4')])
```


## squeeze

## IfErrorArray.squeeze (axis=None)

Remove single-dimensional entries from the shape of $a$.
Refer to numpy.squeeze for full documentation.
numpy.squeeze : equivalent function
std

IfErrorArray.std (axis=None, dtype=None, out=None, ddof=0, keepdims=False)
Returns the standard deviation of the array elements along given axis.
Refer to numpy.std for full documentation.
numpy.std : equivalent function
sum

IfErrorArray.sum (axis=None, dtype=None, out=None, keepdims=False)
Return the sum of the array elements over the given axis.
Refer to numpy.sum for full documentation.
numpy.sum : equivalent function
swapaxes

IfErrorArray.swapaxes (axis1, axis2)
Return a view of the array with axis1 and axis2 interchanged.
Refer to numpy.swapaxes for full documentation.
numpy.swapaxes : equivalent function
take

IfErrorArray.take (indices, axis=None, out=None, mode='raise')
Return an array formed from the elements of $a$ at the given indices.
Refer to numpy.take for full documentation.
numpy.take : equivalent function

## tobytes

IfErrorArray.tobytes (order='C')
Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.

New in version 1.9.0.
order [\{ 'C', 'F', None\}, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01\x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01\x00\x00\x00\x03\x00\x00\x00'
```

tofile

IfErrorArray.tofile (fid, sep="", format="\%s")
Write array to a file as text or binary (default).
Data is always written in ' $C$ ' order, independent of the order of $a$. The data produced by this method can be recovered using the function fromfile().
fid [file or str] An open file object, or a string containing a filename.
sep [str] Separator between array items for text output. If "" (empty), a binary file is written, equivalent to file.write(a.tobytes()).
format [str] Format string for text file output. Each entry in the array is formatted to text by first converting it to the closest Python type, and then using "format" \% item.
This is a convenience function for quick storage of array data. Information on endianness and precision is lost, so this method is not a good choice for files intended to archive data or transport data between machines with different endianness. Some of these problems can be overcome by outputting the data as text files, at the expense of speed and file size.

## tolist

## IfErrorArray.tolist()

Return the array as a (possibly nested) list.
Return a copy of the array data as a (nested) Python list. Data items are converted to the nearest compatible Python type.
none
$\mathbf{y}$ [list] The possibly nested list of array elements.
The array may be recreated, $a=n p . a r r a y(a . t o l i s t())$.

```
>>> a = np.array([1, 2])
>>> a.tolist()
[1, 2]
>>> a = np.array([[1, 2], [3, 4]])
>>> list(a)
[array([1, 2]), array([3, 4])]
>>> a.tolist()
[[1, 2], [3, 4]]
```


## tostring

IfErrorArray.tostring (order='C')
Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.

This function is a compatibility alias for tobytes. Despite its name it returns bytes not strings.
order [ $\{$ ' C ', ' F ', None \}, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01 \x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01 \x00\x00\x00\x03\x00\x00\x00'
```


## trace

IfErrorArray.trace (offset=0, axis $1=0$, axis $2=1$, dtype $=$ None, out $=$ None)
Return the sum along diagonals of the array.
Refer to numpy.trace for full documentation.
numpy.trace : equivalent function

## transpose

IfErrorArray.transpose (*axes)
Returns a view of the array with axes transposed.
For a 1-D array, this has no effect. (To change between column and row vectors, first cast the 1-D array into a matrix object.) For a 2-D array, this is the usual matrix transpose. For an n-D array, if axes are given, their order indicates how the axes are permuted (see Examples). If axes are not provided

```
and a.shape = (i[0], i[1], ... i[n-2], i[n-1]),then a.transpose().shape =
```

(i[n-1], i[n-2], ... i[1], i[0]).
axes : None, tuple of ints, or $n$ ints

- None or no argument: reverses the order of the axes.
- tuple of ints: $i$ in the $j$-th place in the tuple means $a$ 's $i$-th axis becomes a.transpose()'s $j$-th axis.
- $n$ ints: same as an n-tuple of the same ints (this form is intended simply as a "convenience" alternative to the tuple form)
out [ndarray] View of $a$, with axes suitably permuted.
ndarray.T : Array property returning the array transposed.

```
>>> a = np.array([[1, 2], [3, 4]])
>>> a
array([[1, 2],
    [3, 4]])
>>> a.transpose()
array([[1, 3],
    [2, 4]])
>>> a.transpose((1, 0))
array([[1, 3],
    [2, 4]])
>>> a.transpose(1, 0)
array([[1, 3],
    [2, 4]])
```

var

IfErrorArray.var (axis=None, dtype=None, out=None, ddof=0, , eeepdims=False )
Returns the variance of the array elements, along given axis.
Refer to numpy.var for full documentation.
numpy.var : equivalent function
view

IfErrorArray.view (dtype=None, type=None)
New view of array with the same data.
dtype [data-type or ndarray sub-class, optional] Data-type descriptor of the returned view, e.g., float32 or int16. The default, None, results in the view having the same data-type as $a$. This argument can also be specified as an ndarray sub-class, which then specifies the type of the returned object (this is equivalent to setting the type parameter).
type [Python type, optional] Type of the returned view, e.g., ndarray or matrix. Again, the default None results in type preservation.
a.view () is used two different ways:
a.view (some_dtype) or a.view (dtype=some_dtype) constructs a view of the array's memory with a different data-type. This can cause a reinterpretation of the bytes of memory.
a.view(ndarray_subclass) or a.view(type=ndarray_subclass) just returns an instance of ndarray_subclass that looks at the same array (same shape, dtype, etc.) This does not cause a reinterpretation of the memory.

For a.view (some_dtype), if some_dtype has a different number of bytes per entry than the previous dtype (for example, converting a regular array to a structured array), then the behavior of the view cannot be predicted just from the superficial appearance of a (shown by print (a)). It also depends on exactly how a is stored in memory. Therefore if a is C-ordered versus fortran-ordered, versus defined as a slice or transpose, etc., the view may give different results.

```
>>> x = np.array([(1, 2)], dtype=[('a'', np.int8), ('b', np.int8)])
```

Viewing array data using a different type and dtype:

```
>>> y = x.view(dtype=np.int16, type=np.matrix)
>>> Y
matrix([[513]], dtype=int16)
>>> print(type(y))
<class 'numpy.matrixlib.defmatrix.matrix'>
```

Creating a view on a structured array so it can be used in calculations

```
>>> x = np.array([(1, 2), (3,4)], dtype=[('a', np.int8), ('b', np.int8)])
>>> xv = x.view(dtype=np.int8).reshape(-1,2)
>>> xv
array([[1, 2],
    [3, 4]], dtype=int8)
>>> xv.mean(0)
array([ 2., 3.])
```

Making changes to the view changes the underlying array

```
>>> xv[0,1] = 20
>>> print(x)
[(1, 20) (3, 4)]
```

Using a view to convert an array to a recarray:

```
>>> z = x.view(np.recarray)
>>> z.a
array([1], dtype=int8)
```

Views share data:

```
>>> x[0] = (9, 10)
>>> z[0]
(9, 10)
```

Views that change the dtype size (bytes per entry) should normally be avoided on arrays defined by slices, transposes, fortran-ordering, etc.:

```
>>> x = np.array([[1,2,3],[4,5,6]], dtype=np.int16)
>>> y = x[:, 0:2]
>>> y
array([[1, 2],
    [4, 5]], dtype=int16)
>>> y.view(dtype=[('width', np.int16), ('length', np.int16)])
```

```
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: new type not compatible with array.
>>> z = y.copy()
>>> z.view(dtype=[('width', np.int16), ('length', np.int16)])
array([[(1, 2)],
        [(4, 5)]], dtype=[('width', '<i2'), ('length', '<i2')])
```

    _init__()
    Initialize self. See help(type(self)) for accurate signature.

## Attributes

| T | Same as self.transpose(), except that self is returned <br> if self.ndim <2. |
| :--- | :--- |
| base | Base object if memory is from some other object. |
| ctypes | An object to simplify the interaction of the array with <br> the ctypes module. |
| data | Python buffer object pointing to the start of the ar- <br> ray's data. |
| dtype | Data-type of the array's elements. |
| flags | Information about the memory layout of the array. |
| flat | A 1-D iterator over the array. |
| imag | The imaginary part of the array. |
| itemsize | Length of one array element in bytes. |
| nbytes | Total bytes consumed by the elements of the array. |
| ndim | Number of array dimensions. |
| real | The real part of the array. |
| shape | Tuple of array dimensions. |
| size | Number of elements in the array. |
| strides | Tuple of bytes to step in each dimension when <br> traversing an array. |

## T

## IfErrorArray.T

Same as self.transpose(), except that self is returned if self.ndim $<2$.

```
>>> x = np.array([[1.,2.],[3.,4.]])
>>> x
array([[ 1., 2.],
    [ 3., 4.]])
>>> x.T
array([[ 1., 3.],
    [ 2., 4.]])
>>> x = np.array([1.,2.,3.,4.])
>>> x
array([ 1., 2., 3., 4.])
>>> x.T
array([ 1., 2., 3., 4.])
```


## base

## IfErrorArray.base

Base object if memory is from some other object.
The base of an array that owns its memory is None:

```
>>> x = np.array([1,2,3,4])
>>> x.base is None
True
```

Slicing creates a view, whose memory is shared with x :

```
>>> y = x[2:]
>>> y.base is x
True
```

ctypes

## IfErrorArray.ctypes

An object to simplify the interaction of the array with the ctypes module.
This attribute creates an object that makes it easier to use arrays when calling shared libraries with the ctypes module. The returned object has, among others, data, shape, and strides attributes (see Notes below) which themselves return ctypes objects that can be used as arguments to a shared library.
None
c [Python object] Possessing attributes data, shape, strides, etc.
numpy.ctypeslib
Below are the public attributes of this object which were documented in "Guide to NumPy" (we have omitted undocumented public attributes, as well as documented private attributes):

- data: A pointer to the memory area of the array as a Python integer. This memory area may contain data that is not aligned, or not in correct byte-order. The memory area may not even be writeable. The array flags and data-type of this array should be respected when passing this attribute to arbitrary C-code to avoid trouble that can include Python crashing. User Beware! The value of this attribute is exactly the same as self._array_interface_['data’][0].
- shape (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the C-integer corresponding to dtype(' p ') on this platform. This base-type could be c_int, c_long, or c_longlong depending on the platform. The c_intp type is defined accordingly in numpy.ctypeslib. The ctypes array contains the shape of the underlying array.
- strides (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the same as for the shape attribute. This ctypes array contains the strides information from the underlying array. This strides information is important for showing how many bytes must be jumped to get to the next element in the array.
- data_as(obj): Return the data pointer cast to a particular c-types object. For example, calling self._as_parameter_ is equivalent to self.data_as(ctypes.c_void_p). Perhaps you want to use the data as a pointer to a ctypes array of floating-point data: self.data_as(ctypes.POINTER(ctypes.c_double)).
- shape_as(obj): Return the shape tuple as an array of some other c-types type. For example: self.shape_as(ctypes.c_short).
- strides_as(obj): Return the strides tuple as an array of some other c-types type. For example: self.strides_as(ctypes.c_longlong).
Be careful using the ctypes attribute - especially on temporary arrays or arrays constructed on the fly. For example, calling ( $a+b$ ) . ctypes.data_as (ctypes.c_void_p) returns a pointer to memory that is invalid because the array created as $(a+b)$ is deallocated before the next Python statement. You can avoid this problem using either $c=a+b$ or $c t=(a+b) . c t y p e s$. In the latter case, ct will hold a reference to the array until ct is deleted or re-assigned.

If the ctypes module is not available, then the ctypes attribute of array objects still returns something useful, but ctypes objects are not returned and errors may be raised instead. In particular, the object will still have the as parameter attribute which will return an integer equal to the data attribute.

```
>>> import ctypes
>>> x
array([[0, 1],
        [2, 3]])
>>> x.ctypes.data
30439712
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long))
<ctypes.LP_c_long object at 0x01F01300>
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long)).contents
c_long(0)
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_longlong)).contents
c_longlong(4294967296L)
>>> x.ctypes.shape
<numpy.core._internal.c_long_Array_2 object at 0x01FFD580>
>>> x.ctypes.shape_as(ctypes.c_long)
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides_as(ctypes.c_longlong)
<numpy.core._internal.c_longlong_Array_2 object at 0x01F01300>
```

data

IfErrorArray.data
Python buffer object pointing to the start of the array's data.

## dtype

IfErrorArray.dtype
Data-type of the array's elements.
None
d : numpy dtype object
numpy.dtype

```
>>> x
array([[0, 1],
    [2, 3]])
>>> x.dtype
dtype('int32')
```

```
>>> type(x.dtype)
```

<type 'numpy.dtype'>

## flags

## IfErrorArray.flags

Information about the memory layout of the array.
C_CONTIGUOUS (C) The data is in a single, C-style contiguous segment.
F_CONTIGUOUS (F) The data is in a single, Fortran-style contiguous segment.
OWNDATA ( $\mathbf{O}$ ) The array owns the memory it uses or borrows it from another object.
WRITEABLE (W) The data area can be written to. Setting this to False locks the data, making it readonly. A view (slice, etc.) inherits WRITEABLE from its base array at creation time, but a view of a writeable array may be subsequently locked while the base array remains writeable. (The opposite is not true, in that a view of a locked array may not be made writeable. However, currently, locking a base object does not lock any views that already reference it, so under that circumstance it is possible to alter the contents of a locked array via a previously created writeable view onto it.) Attempting to change a non-writeable array raises a RuntimeError exception.

ALIGNED (A) The data and all elements are aligned appropriately for the hardware.
UPDATEIFCOPY (U) This array is a copy of some other array. When this array is deallocated, the base array will be updated with the contents of this array.

FNC F_CONTIGUOUS and not C_CONTIGUOUS.
FORC F_CONTIGUOUS or C_CONTIGUOUS (one-segment test).
BEHAVED (B) ALIGNED and WRITEABLE.
CARRAY (CA) BEHAVED and C_CONTIGUOUS.
FARRAY (FA) BEHAVED and F_CONTIGUOUS and not C_CONTIGUOUS.
The flags object can be accessed dictionary-like (as in a.flags ['WRITEABLE' ]), or by using lowercased attribute names (as in a.flags.writeable). Short flag names are only supported in dictionary access.

Only the UPDATEIFCOPY, WRITEABLE, and ALIGNED flags can be changed by the user, via direct assignment to the attribute or dictionary entry, or by calling ndarray.setflags.
The array flags cannot be set arbitrarily:

- UPDATEIFCOPY can only be set False.
- ALIGNED can only be set True if the data is truly aligned.
- WRITEABLE can only be set True if the array owns its own memory or the ultimate owner of the memory exposes a writeable buffer interface or is a string.

Arrays can be both C-style and Fortran-style contiguous simultaneously. This is clear for 1-dimensional arrays, but can also be true for higher dimensional arrays.

Even for contiguous arrays a stride for a given dimension arr.strides [dim] may be arbitrary if arr.shape[dim] == 1 or the array has no elements. It does not generally hold that self. strides[-1] == self.itemsize for C-style contiguous arrays or self.strides[0] == self.itemsize for Fortran-style contiguous arrays is true.
flat

## IfErrorArray.flat

A 1-D iterator over the array.
This is a numpy.flatiter instance, which acts similarly to, but is not a subclass of, Python's built-in iterator object.
flatten : Return a copy of the array collapsed into one dimension.
flatiter

```
>>> x = np.arange(1, 7).reshape(2, 3)
>>> x
array([[1, 2, 3],
    [4, 5, 6]])
>>> x.flat[3]
4
>>> x.T
array([[1, 4],
    [2, 5],
    [3, 6]])
>>> x.T.flat[3]
5
>>> type(x.flat)
<type 'numpy.flatiter'>
```

An assignment example:

```
>>> x.flat = 3; x
array([[3, 3, 3],
    [3, 3, 3]])
>>> x.flat[[1,4]] = 1; x
array([[3, 1, 3],
    [3, 1, 3]])
```

imag

IfErrorArray.imag
The imaginary part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.imag
array([ 0. , 0.70710678])
>>> x.imag.dtype
dtype('float64')
```


## itemsize

## IfErrorArray.itemsize

Length of one array element in bytes.
>>> $x=n p$.array $([1,2,3]$, dtype=np.float64)
>>> x.itemsize

```
8
>>> x = np.array([1,2,3], dtype=np.complex128)
>>> x.itemsize
16
```


## nbytes

IfErrorArray.nbytes
Total bytes consumed by the elements of the array.
Does not include memory consumed by non-element attributes of the array object.

```
>>> x = np.zeros((3,5,2), dtype=np.complex128)
>>> x.nbytes
480
>>> np.prod(x.shape) * x.itemsize
480
```

ndim

IfErrorArray.ndim
Number of array dimensions.

```
>>> x = np.array([1, 2, 3])
>>> x.ndim
1
>>> y = np.zeros((2, 3, 4))
>>> y.ndim
3
```

real

## IfErrorArray.real

The real part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.real
array([ 1. , 0.70710678])
>>> x.real.dtype
dtype('float64')
```

numpy.real : equivalent function

## shape

IfErrorArray.shape
Tuple of array dimensions.
May be used to "reshape" the array, as long as this would not require a change in the total number of elements

```
>>> x = np.array([1, 2, 3, 4])
>>> x.shape
(4,)
>>> y = np.zeros((2, 3, 4))
>>> y.shape
(2, 3, 4)
>>> y.shape = (3, 8)
>>> Y
array([[ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.]])
>>> y.shape = (3, 6)
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: total size of new array must be unchanged
```

size

IfErrorArray.size
Number of elements in the array.
Equivalent to np.prod (a.shape), i.e., the product of the array's dimensions.

```
>>> x = np.zeros((3, 5, 2), dtype=np.complex128)
>>> x.size
30
>>> np.prod(x.shape)
30
```


## strides

IfErrorArray.strides
Tuple of bytes to step in each dimension when traversing an array.
The byte offset of element (i[0], i[1], ..., $i[n]$ ) in an array $a$ is:

```
offset = sum(np.array(i) * a.strides)
```

A more detailed explanation of strides can be found in the "ndarray.rst" file in the NumPy reference guide.
Imagine an array of 32-bit integers (each 4 bytes):

```
x = np.array([[0, 1, 2, 3, 4],
    [5, 6, 7, 8, 9]], dtype=np.int32)
```

This array is stored in memory as 40 bytes, one after the other (known as a contiguous block of memory). The strides of an array tell us how many bytes we have to skip in memory to move to the next position along a certain axis. For example, we have to skip 4 bytes ( 1 value) to move to the next column, but 20 bytes ( 5 values) to get to the same position in the next row. As such, the strides for the array $x$ will be $(20,4)$.
numpy.lib.stride_tricks.as_strided

```
>>> y = np.reshape(np.arange(2*3*4), (2,3,4))
>>> y
array([[[ 0, 1, 2, 3],
    [ 4, 5, 6, 7],
    [ 8, 9, 10, 11]],
    [[12, 13, 14, 15],
    [16, 17, 18, 19],
    [20, 21, 22, 23]]])
>>> y.strides
(48, 16, 4)
>>> y[1,1,1]
17
>>> offset=sum(y.strides * np.array((1,1,1)))
>>> offset/y.itemsize
1 7
```

```
>>> x = np.reshape(np.arange(5*6*7*8), (5,6,7,8)).transpose(2,3,1,0)
>>> x.strides
(32, 4, 224, 1344)
>>> i = np.array([3,5,2,2])
>>> offset = sum(i * x.strides)
>>> x[3,5,2,2]
813
>>> offset / x.itemsize
813
```

look

Python equivalents of lookup and reference Excel functions.

## Functions

| xcolumn |
| :--- |
| xlookup |
| xratch |
| xcolumn |
| xcolumn (cell=None, ref=None) |
| xlookup |
| xlookup (lookup_val, lookup_vec, result_vec=None, match_type=1) |
| xmatch |
| xmatch (lookup_value, lookup_array, match_type=1) |

```
xrow
xrow (cell=None, ref=None)
math
```

Python equivalents of math and trigonometry Excel functions.

Functions

| xarabic |
| :--- |
| xarctan2 |
| xceiling |
| xceiling_math |
| xcot |
| xdecimal |
| xeven |
| xfact |
| xfactdouble |
| xmod |
| xmround |
| xodd |
| xpower |
| xrandbetween |
| xroman |
| xround |
| xsrqtpi |
| xsum |
| xsumproduct |

## xarabic

xarabic (text)
xarctan2
xarctan2 $(x, y)$
xceiling
xceiling (num, sig, ceil=<built-in function ceil>, $d f=0$ )
xceiling_math
xceiling_math (num, sig=None, mode $=0$, ceil $=<$ built-in function ceil $>$ )

```
xcot
xcot (x, func=<ufunc 'tan'>)
xdecimal
xdecimal(text, radix)
xeven
xeven (x)
xfact
xfact (number, fact=<built-in function factorial>, limit=0)
xfactdouble
xfactdouble(number)
xmod
xmod (x,y)
xmround
xmround(*args)
xodd
xodd (x)
xpower
xpower(number, power)
xrandbetween
xrandbetween (bottom, top)
xroman
xroman(num, form=0)
```

```
xround
xround ( }x,d,func=<built-in function round> )
xsrqtpi
xsrqtpi(number)
xsum
xsum(*args)
xsumproduct
xsumproduct (*args)
operators
```

Python equivalents of Excel operators.

## Functions

| logic_input_parser |  |
| :--- | :--- |
| logic_wrap | Helps call a numpy universal function (ufunc). |
| numeric_wrap | Helps call a numpy universal function (ufunc). |

logic_input_parser

```
logic_input_parser (x,y)
```

logic_wrap
logic_wrap (func, *, input_parser=<function logic_input_parser>, check_error=<function get_error>,
args_parser $=<$ function $<$ lambda>>, otype $=<$ function $<$ lambda>>, ranges $=$ False,$* * k w$ )

Helps call a numpy universal function (ufunc).

```
numeric_wrap
```

numeric_wrap (func, input_parser=<function <lambda>>, check_error=<function get_error>,
args_parser $=<$ function <lambda>>, *, otype $=<$ function <lambda>>, ranges=False,
***w)

Helps call a numpy universal function (ufunc).

## Classes

```
OperatorArray
OperatorArray
class OperatorArray
```


## Methods

| all | Returns True if all elements evaluate to True. |
| :---: | :---: |
| any | Returns True if any of the elements of $a$ evaluate to True. |
| argmax | Return indices of the maximum values along the given axis. |
| argmin | Return indices of the minimum values along the given axis of $a$. |
| argpartition | Returns the indices that would partition this array. |
| argsort | Returns the indices that would sort this array. |
| astype | Copy of the array, cast to a specified type. |
| byteswap | Swap the bytes of the array elements |
| choose | Use an index array to construct a new array from a set of choices. |
| clip | Return an array whose values are limited to [min, max]. |
| collapse |  |
| compress | Return selected slices of this array along given axis. |
| conj | Complex-conjugate all elements. |
| conjugate | Return the complex conjugate, element-wise. |
| copy | Return a copy of the array. |
| cumprod | Return the cumulative product of the elements along the given axis. |
| cumsum | Return the cumulative sum of the elements along the given axis. |
| diagonal | Return specified diagonals. |
| dot | Dot product of two arrays. |
| dump | Dump a pickle of the array to the specified file. |
| dumps | Returns the pickle of the array as a string. |
| fill | Fill the array with a scalar value. |
| flatten | Return a copy of the array collapsed into one dimension. |
| getfield | Returns a field of the given array as a certain type. |
| item | Copy an element of an array to a standard Python scalar and return it. |
| itemset | Insert scalar into an array (scalar is cast to array's dtype, if possible) |
| $\max$ | Return the maximum along a given axis. |
| mean | Returns the average of the array elements along given axis. |

Continued on next page

Table 61 - continued from previous page

| min | Return the minimum along a given axis. |
| :---: | :---: |
| newbyteorder | Return the array with the same data viewed with a different byte order. |
| nonzero | Return the indices of the elements that are non-zero. |
| partition | Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. |
| prod | Return the product of the array elements over the given axis |
| ptp | Peak to peak (maximum - minimum) value along a given axis. |
| put | Set a.flat $[\mathrm{n}]=$ values $[\mathrm{n}]$ for all $n$ in indices. |
| ravel | Return a flattened array. |
| repeat | Repeat elements of an array. |
| reshape | Returns an array containing the same data with a new shape. |
| resize | Change shape and size of array in-place. |
| round | Return $a$ with each element rounded to the given number of decimals. |
| searchsorted | Find indices where elements of v should be inserted in a to maintain order. |
| setfield | Put a value into a specified place in a field defined by a data-type. |
| setflags | Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively. |
| sort | Sort an array, in-place. |
| squeeze | Remove single-dimensional entries from the shape of $a$. |
| std | Returns the standard deviation of the array elements along given axis. |
| sum | Return the sum of the array elements over the given axis. |
| swapaxes | Return a view of the array with axis1 and axis2 interchanged. |
| take | Return an array formed from the elements of $a$ at the given indices. |
| tobytes | Construct Python bytes containing the raw data bytes in the array. |
| tofile | Write array to a file as text or binary (default). |
| tolist | Return the array as a (possibly nested) list. |
| tostring | Construct Python bytes containing the raw data bytes in the array. |
| trace | Return the sum along diagonals of the array. |
| transpose | Returns a view of the array with axes transposed. |
| var | Returns the variance of the array elements, along given axis. |
| view | New view of array with the same data. |

## all

OperatorArray.all (axis=None, out=None, keepdims=False)
Returns True if all elements evaluate to True.
Refer to numpy.all for full documentation.
numpy.all : equivalent function
any

OperatorArray. any (axis=None, out=None, keepdims=False)
Returns True if any of the elements of $a$ evaluate to True.
Refer to numpy.any for full documentation.
numpy.any : equivalent function
$\operatorname{argmax}$

OperatorArray.argmax (axis=None, out=None)
Return indices of the maximum values along the given axis.
Refer to numpy.argmax for full documentation.
numpy.argmax : equivalent function

## argmin

OperatorArray.argmin (axis=None, out=None)
Return indices of the minimum values along the given axis of $a$.
Refer to numpy.argmin for detailed documentation.
numpy.argmin : equivalent function

## argpartition

OperatorArray.argpartition (kth, axis=-1, kind='introselect', order=None)
Returns the indices that would partition this array.
Refer to numpy.argpartition for full documentation.
New in version 1.8.0.
numpy.argpartition : equivalent function
argsort

OperatorArray.argsort (axis=-1, kind='quicksort', order=None)
Returns the indices that would sort this array.
Refer to numpy.argsort for full documentation.
numpy.argsort : equivalent function

## astype

OperatorArray. astype (dtype, order='K', casting='unsafe', subok=True, copy=True)
Copy of the array, cast to a specified type.
dtype [str or dtype] Typecode or data-type to which the array is cast.
order [ $\{$ ' $C$ ', ' $F$ ', ' $A$ ', ' $K$ ' \}, optional] Controls the memory layout order of the result. ' $C$ ' means $C$ order, ' $F$ ' means Fortran order, 'A' means ' $F$ ' order if all the arrays are Fortran contiguous, 'C' order otherwise, and ' K ' means as close to the order the array elements appear in memory as possible. Default is ' K '.
casting [\{ 'no', 'equiv', 'safe', 'same_kind', 'unsafe'\}, optional] Controls what kind of data casting may occur. Defaults to 'unsafe' for backwards compatibility.

- 'no' means the data types should not be cast at all.
- 'equiv’ means only byte-order changes are allowed.
- 'safe' means only casts which can preserve values are allowed.
- 'same_kind' means only safe casts or casts within a kind, like float64 to float32, are allowed.
- 'unsafe' means any data conversions may be done.
subok [bool, optional] If True, then sub-classes will be passed-through (default), otherwise the returned array will be forced to be a base-class array.
copy [bool, optional] By default, astype always returns a newly allocated array. If this is set to false, and the dtype, order, and subok requirements are satisfied, the input array is returned instead of a copy.
arr_t [ndarray] Unless copy is False and the other conditions for returning the input array are satisfied (see description for copy input parameter), arr $\_$is a new array of the same shape as the input array, with dtype, order given by dtype, order.

Starting in NumPy 1.9, astype method now returns an error if the string dtype to cast to is not long enough in 'safe' casting mode to hold the max value of integer/float array that is being casted. Previously the casting was allowed even if the result was truncated.
ComplexWarning When casting from complex to float or int. To avoid this, one should use a.real. astype ( t ).

```
>>> x = np.array([1, 2, 2.5])
>>> X
array([ 1., 2. , 2.5])
```

```
>>> x.astype(int)
array([1, 2, 2])
```


## byteswap

OperatorArray.byteswap (inplace)
Swap the bytes of the array elements
Toggle between low-endian and big-endian data representation by returning a byteswapped array, optionally swapped in-place.
inplace [bool, optional] If True, swap bytes in-place, default is False.
out [ndarray] The byteswapped array. If inplace is True, this is a view to self.

```
>>> A = np.array([1, 256, 8755], dtype=np.int16)
>>> map(hex, A)
['0x1', '0x100', '0x2233']
>>> A.byteswap(True)
array([ 256, 1, 13090], dtype=int16)
>>> map(hex, A)
['0x100', '0x1', '0x3322']
```

Arrays of strings are not swapped

```
>>> A = np.array(['ceg', 'fac'])
>>> A.byteswap()
array(['ceg', 'fac'],
    dtype='|S3')
```


## choose

OperatorArray. choose (choices, out=None, mode='raise')
Use an index array to construct a new array from a set of choices.
Refer to numpy.choose for full documentation.
numpy.choose : equivalent function
clip

OperatorArray.clip ( min=None, max=None, out=None)
Return an array whose values are limited to [min, max]. One of max or min must be given.
Refer to numpy.clip for full documentation.
numpy.clip : equivalent function
collapse

OperatorArray.collapse (shape)
compress

OperatorArray.compress (condition, axis=None, out=None)
Return selected slices of this array along given axis.
Refer to numpy.compress for full documentation.
numpy.compress : equivalent function
conj

OperatorArray.conj()
Complex-conjugate all elements.

Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function

## conjugate

OperatorArray.conjugate ()
Return the complex conjugate, element-wise.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function
copy

OperatorArray. copy (order='C')
Return a copy of the array.
order [ $\{$ ' C ', ' F ', ' A ', ' K '\}, optional] Controls the memory layout of the copy. ' C ' means C -order, ' F ' means $F$-order, ' $A$ ' means ' F ' if $a$ is Fortran contiguous, ' C ' otherwise. ' K ' means match the layout of $a$ as closely as possible. (Note that this function and :func:numpy.copy are very similar, but have different default values for their order= arguments.)
numpy.copy numpy.copyto

```
>>> x = np.array([[1,2,3],[4,5,6]], order='F')
```

```
>>> y = x.copy()
```

>>> x.fill(0)

```
>>> x
array([[0, 0, 0],
    [0, 0, 0]])
```

```
>>> y
array([[1, 2, 3],
    [4, 5, 6]])
```

```
>>> y.flags['C_CONTIGUOUS']
True
```

cumprod

OperatorArray. cumprod (axis=None, dtype=None, out=None)
Return the cumulative product of the elements along the given axis.
Refer to numpy.cumprod for full documentation.
numpy.cumprod : equivalent function

## cumsum

OperatorArray. cumsum (axis=None, dtype=None, out=None)
Return the cumulative sum of the elements along the given axis.
Refer to numpy.cumsum for full documentation.
numpy.cumsum : equivalent function

## diagonal

OperatorArray.diagonal (offset $=0$, axis $1=0$, axis $2=1$ )
Return specified diagonals. In NumPy 1.9 the returned array is a read-only view instead of a copy as in previous NumPy versions. In a future version the read-only restriction will be removed.

Refer to numpy.diagonal () for full documentation.
numpy.diagonal : equivalent function
dot

OperatorArray.dot ( $b$, out=None)
Dot product of two arrays.
Refer to numpy.dot for full documentation.
numpy.dot : equivalent function

```
>>> a = np.eye(2)
>>> b = np.ones((2, 2)) * 2
>>> a.dot(b)
array([[ 2., 2.],
    [ 2., 2.]])
```

This array method can be conveniently chained:

```
>>> a.dot(b).dot(b)
array([[ 8., 8.],
    [ 8., 8.]])
```

dump

OperatorArray.dump (file)
Dump a pickle of the array to the specified file. The array can be read back with pickle.load or numpy.load.
file [str] A string naming the dump file.

## dumps

## OperatorArray.dumps()

Returns the pickle of the array as a string. pickle.loads or numpy.loads will convert the string back to an array.

None
fill

OperatorArray.fill (value)
Fill the array with a scalar value.
value [scalar] All elements of $a$ will be assigned this value.

```
>>> a = np.array([1, 2])
>>> a.fill(0)
>>> a
array([0, 0])
>>> a = np.empty(2)
>>> a.fill(1)
>>> a
array([ 1., 1.])
```


## flatten

OperatorArray.flatten (order='C')
Return a copy of the array collapsed into one dimension.
order [\{ 'C', ' F ', ' A ', ' K '\}, optional] ' C ' means to flatten in row-major (C-style) order. ' F ' means to flatten in column-major (Fortran- style) order. 'A' means to flatten in column-major order if $a$ is Fortran contiguous in memory, row-major order otherwise. ' K ' means to flatten $a$ in the order the elements occur in memory. The default is ' C '.
$\mathbf{y}$ [ndarray] A copy of the input array, flattened to one dimension.
ravel : Return a flattened array. flat : A 1-D flat iterator over the array.

```
>>> a = np.array([[1,2], [3,4]])
>>> a.flatten()
array([1, 2, 3, 4])
>>> a.flatten('F')
array([1, 3, 2, 4])
```


## getfield

OperatorArray.getfield (dtype, offset=0)
Returns a field of the given array as a certain type.
A field is a view of the array data with a given data-type. The values in the view are determined by the given type and the offset into the current array in bytes. The offset needs to be such that the view dtype fits in the array dtype; for example an array of dtype complex 128 has 16-byte elements. If taking a view with a 32-bit integer ( 4 bytes), the offset needs to be between 0 and 12 bytes.
dtype [str or dtype] The data type of the view. The dtype size of the view can not be larger than that of the array itself.
offset [int] Number of bytes to skip before beginning the element view.

```
>>> x = np.diag([1.+1.j]*2)
>>> x[1, 1] = 2 + 4.j
>>> x
```

```
array([[ 1.+1.j, 0.+0.j],
    [ 0.+0.j, 2.+4.j]])
>>> x.getfield(np.float64)
array([[ 1., 0.],
    [ 0., 2.]])
```

By choosing an offset of 8 bytes we can select the complex part of the array for our view:

```
>>> x.getfield(np.float64, offset=8)
array([[ 1., 0.],
    [ 0., 4.]])
```

item

## OperatorArray.item (*args)

Copy an element of an array to a standard Python scalar and return it.
*args : Arguments (variable number and type)

- none: in this case, the method only works for arrays with one element (a.size $==1$ ), which element is copied into a standard Python scalar object and returned.
- int_type: this argument is interpreted as a flat index into the array, specifying which element to copy and return.
- tuple of int_types: functions as does a single int_type argument, except that the argument is interpreted as an nd-index into the array.
$\mathbf{z}$ [Standard Python scalar object] A copy of the specified element of the array as a suitable Python scalar
When the data type of $a$ is longdouble or clongdouble, item() returns a scalar array object because there is no available Python scalar that would not lose information. Void arrays return a buffer object for item(), unless fields are defined, in which case a tuple is returned.
item is very similar to a[args], except, instead of an array scalar, a standard Python scalar is returned. This can be useful for speeding up access to elements of the array and doing arithmetic on elements of the array using Python's optimized math.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.item(3)
2
>>> x.item(7)
5
>>> x.item((0, 1))
1
>>> x.item((2, 2))
3
```


## itemset

OperatorArray.itemset (*args)
Insert scalar into an array (scalar is cast to array's dtype, if possible)
There must be at least 1 argument, and define the last argument as item. Then, a.itemset (*args) is equivalent to but faster than a [args] = item. The item should be a scalar value and args must select a single item in the array $a$.

1*args [Arguments] If one argument: a scalar, only used in case $a$ is of size 1 . If two arguments: the last argument is the value to be set and must be a scalar, the first argument specifies a single array element location. It is either an int or a tuple.

Compared to indexing syntax, itemset provides some speed increase for placing a scalar into a particular location in an ndarray, if you must do this. However, generally this is discouraged: among other problems, it complicates the appearance of the code. Also, when using itemset (and item) inside a loop, be sure to assign the methods to a local variable to avoid the attribute look-up at each loop iteration.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.itemset (4, 0)
>>> x.itemset ((2, 2), 9)
>>> x
array([[3, 1, 7],
    [2, 0, 3],
    [8, 5, 9]])
```

max

OperatorArray.max (axis=None, out=None)
Return the maximum along a given axis.
Refer to numpy.amax for full documentation.
numpy.amax : equivalent function
mean

OperatorArray.mean (axis=None, dtype=None, out=None, keepdims=False)
Returns the average of the array elements along given axis.
Refer to numpy.mean for full documentation.
numpy.mean : equivalent function
min

OperatorArray.min (axis=None, out=None, keepdims=False)
Return the minimum along a given axis.
Refer to numpy.amin for full documentation.
numpy.amin : equivalent function

## newbyteorder

OperatorArray. newbyteorder (new_order='S')
Return the array with the same data viewed with a different byte order.
Equivalent to:
arr.view(arr.dtype.newbytorder(new_order))

Changes are also made in all fields and sub-arrays of the array data type.
new_order [string, optional] Byte order to force; a value from the byte order specifications below. new_order codes can be any of:

- 'S' - swap dtype from current to opposite endian
- \{ '<', 'L'\} - little endian
- \{'>', 'B'\} - big endian
- $\{$ ' $=$ ', ' $N$ ' $\}$ - native order
- \{'l', 'I' \} - ignore (no change to byte order)

The default value (' $S$ ') results in swapping the current byte order. The code does a case-insensitive check on the first letter of new_order for the alternatives above. For example, any of ' B ' or ' $b$ ' or 'biggish' are valid to specify big-endian.
new_arr [array] New array object with the dtype reflecting given change to the byte order.

## nonzero

OperatorArray.nonzero()
Return the indices of the elements that are non-zero.
Refer to numpy.nonzero for full documentation.
numpy.nonzero : equivalent function

## partition

OperatorArray.partition (kth, axis=-1, kind='introselect', order=None)
Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. All elements smaller than the kth element are moved before this element and all equal or greater are moved behind it. The ordering of the elements in the two partitions is undefined.

New in version 1.8.0.
kth [int or sequence of ints] Element index to partition by. The kth element value will be in its final sorted position and all smaller elements will be moved before it and all equal or greater elements behind it. The order all elements in the partitions is undefined. If provided with a sequence of kth it will partition all elements indexed by kth of them into their sorted position at once.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'introselect'\}, optional] Selection algorithm. Default is 'introselect'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.partition : Return a parititioned copy of an array. argpartition : Indirect partition. sort : Full sort.
See np. partition for notes on the different algorithms.

```
>>> a = np.array([3, 4, 2, 1])
>>> a.partition(a, 3)
>>> a
array([2, 1, 3, 4])
```

```
>>> a.partition((1, 3))
```

$\operatorname{array}([1,2,3,4])$
prod

OperatorArray.prod (axis=None, dtype=None, out=None, keepdims=False)
Return the product of the array elements over the given axis
Refer to numpy.prod for full documentation.
numpy.prod : equivalent function
ptp

OperatorArray.ptp (axis=None, out=None)
Peak to peak (maximum - minimum) value along a given axis.
Refer to numpy.ptp for full documentation.
numpy.ptp : equivalent function
put

OperatorArray.put (indices, values, mode='raise')
Seta.flat $[\mathrm{n}]=$ values [ n$]$ for all $n$ in indices.
Refer to numpy.put for full documentation.
numpy.put : equivalent function
ravel

OperatorArray.ravel ([order])
Return a flattened array.
Refer to numpy.ravel for full documentation.
numpy.ravel : equivalent function
ndarray.flat : a flat iterator on the array.

## repeat

OperatorArray.repeat (repeats, axis=None)
Repeat elements of an array.
Refer to numpy.repeat for full documentation.
numpy.repeat : equivalent function

## reshape

OperatorArray. reshape (shape, order='C')
Returns an array containing the same data with a new shape.
Refer to numpy.reshape for full documentation.
numpy.reshape : equivalent function
resize

OperatorArray.resize (new_shape, refcheck=True)
Change shape and size of array in-place.
new_shape [tuple of ints, or $n$ ints] Shape of resized array.
refcheck [bool, optional] If False, reference count will not be checked. Default is True.
None
ValueError If $a$ does not own its own data or references or views to it exist, and the data memory must be changed.

SystemError If the order keyword argument is specified. This behaviour is a bug in NumPy.
resize : Return a new array with the specified shape.
This reallocates space for the data area if necessary.
Only contiguous arrays (data elements consecutive in memory) can be resized.
The purpose of the reference count check is to make sure you do not use this array as a buffer for another Python object and then reallocate the memory. However, reference counts can increase in other ways so if you are sure that you have not shared the memory for this array with another Python object, then you may safely set refcheck to False.

Shrinking an array: array is flattened (in the order that the data are stored in memory), resized, and reshaped:

```
>>> a = np.array([[0, 1], [2, 3]], order='C')
>>> a.resize((2, 1))
>>> a
array([[0],
    [1]])
```

```
>>> a = np.array([[0, 1], [2, 3]], order='F')
>>> a.resize((2, 1))
>>> a
array([[0],
    [2]])
```

Enlarging an array: as above, but missing entries are filled with zeros:

```
>>> b = np.array([[0, 1], [2, 3]])
>>> b.resize(2, 3) # new_shape parameter doesn't have to be a tuple
>>> b
array([[0, 1, 2],
    [3, 0, 0]])
```

Referencing an array prevents resizing...

```
>>> c=a
>>> a.resize((1, 1))
Traceback (most recent call last):
..
ValueError: cannot resize an array that has been referenced ...
```

Unless refcheck is False:

```
>>> a.resize((1, 1), refcheck=False)
>>> a
array([[0]])
>>> c
array([[0]])
```

round

OperatorArray.round (decimals=0, out=None)
Return $a$ with each element rounded to the given number of decimals.
Refer to numpy.around for full documentation.
numpy.around : equivalent function

## searchsorted

OperatorArray. searchsorted ( $v$, side $=$ 'left', sorter $=$ None)
Find indices where elements of v should be inserted in a to maintain order.
For full documentation, see numpy.searchsorted
numpy.searchsorted : equivalent function

## setfield

OperatorArray.setfield (val, dtype, offset=0)
Put a value into a specified place in a field defined by a data-type.
Place val into $a$ 's field defined by dtype and beginning offset bytes into the field.
val [object] Value to be placed in field.
dtype [dtype object] Data-type of the field in which to place val.
offset [int, optional] The number of bytes into the field at which to place val.

None
getfield

```
>>> x = np.eye(3)
>>> x.getfield(np.float64)
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
>>> x.setfield(3, np.int32)
>>> x.getfield(np.int32)
array([[3, 3, 3],
    [3, 3, 3],
    [3, 3, 3]])
>>> x
array([[ 1.00000000e+000, 1.48219694e-323, 1.48219694e-323],
    [ 1.48219694e-323, 1.00000000e+000, 1.48219694e-323],
    [ 1.48219694e-323, 1.48219694e-323, 1.00000000e+000]])
>>> x.setfield(np.eye(3), np.int32)
>>> x
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
```


## setflags

OperatorArray.setflags (write=None, align=None, uic=None)
Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively.
These Boolean-valued flags affect how numpy interprets the memory area used by $a$ (see Notes below). The ALIGNED flag can only be set to True if the data is actually aligned according to the type. The UPDATEIFCOPY flag can never be set to True. The flag WRITEABLE can only be set to True if the array owns its own memory, or the ultimate owner of the memory exposes a writeable buffer interface, or is a string. (The exception for string is made so that unpickling can be done without copying memory.)
write [bool, optional] Describes whether or not $a$ can be written to.
align [bool, optional] Describes whether or not $a$ is aligned properly for its type.
uic [bool, optional] Describes whether or not $a$ is a copy of another "base" array.
Array flags provide information about how the memory area used for the array is to be interpreted. There are 6 Boolean flags in use, only three of which can be changed by the user: UPDATEIFCOPY, WRITEABLE, and ALIGNED.

WRITEABLE (W) the data area can be written to;
ALIGNED (A) the data and strides are aligned appropriately for the hardware (as determined by the compiler);

UPDATEIFCOPY (U) this array is a copy of some other array (referenced by .base). When this array is deallocated, the base array will be updated with the contents of this array.

All flags can be accessed using their first (upper case) letter as well as the full name.

```
>>> Y
array([[3, 1, 7],
    [2, 0, 0],
    [8, 5, 9]])
```

```
>>> y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : True
    ALIGNED : True
    UPDATEIFCOPY : False
>>> y.setflags(write=0, align=0)
>>> y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : False
    ALIGNED : False
    UPDATEIFCOPY : False
>>> y.setflags(uic=1)
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: cannot set UPDATEIFCOPY flag to True
```


## sort

OperatorArray.sort (axis=-1, kind='quicksort', order=None)
Sort an array, in-place.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'quicksort', 'mergesort', 'heapsort'\}, optional] Sorting algorithm. Default is 'quicksort'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.sort : Return a sorted copy of an array. argsort : Indirect sort. lexsort : Indirect stable sort on multiple keys. searchsorted : Find elements in sorted array. partition: Partial sort.

See sort for notes on the different sorting algorithms.

```
>>> a = np.array([[1,4], [3,1]])
>>> a.sort(axis=1)
>>> a
array([[1, 4],
    [1, 3]])
>>> a.sort(axis=0)
>>> a
array([[1, 3],
        [1, 4]])
```

Use the order keyword to specify a field to use when sorting a structured array:

```
>>> a = np.array([('a', 2), ('c', 1)], dtype=[('x', 'S1'), ('y', int)])
>>> a.sort (order='y')
>>> a
array([('c', 1), ('a', 2)],
    dtype=[('x', '|S1'), ('y', '<i4')])
```


## squeeze

OperatorArray.squeeze (axis=None)
Remove single-dimensional entries from the shape of $a$.
Refer to numpy.squeeze for full documentation.
numpy.squeeze : equivalent function
std

OperatorArray.std (axis=None, dtype=None, out=None, ddof=0, keepdims=False)
Returns the standard deviation of the array elements along given axis.
Refer to numpy.std for full documentation.
numpy.std : equivalent function
sum

OperatorArray.sum (axis=None, dtype=None, out=None, keepdims=False)
Return the sum of the array elements over the given axis.
Refer to numpy.sum for full documentation.
numpy.sum : equivalent function
swapaxes

OperatorArray.swapaxes (axis1, axis2)
Return a view of the array with axis1 and axis 2 interchanged.
Refer to numpy.swapaxes for full documentation.
numpy.swapaxes : equivalent function

## take

OperatorArray.take (indices, axis=None, out=None, mode='raise')
Return an array formed from the elements of $a$ at the given indices.
Refer to numpy.take for full documentation.
numpy.take : equivalent function

## tobytes

OperatorArray.tobytes (order='C')
Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.

New in version 1.9.0.
order [\{ 'C', 'F', None \}, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01\x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b}\mp@subsup{b}{}{\prime}\x00\x00\x00\x00\x02\x00\x00\x00\x0\\x00\x00\x00\x03\x00\x00\x00'
```

tofile

OperatorArray.tofile (fid, sep="", format="\%s")
Write array to a file as text or binary (default).
Data is always written in ' $C$ ' order, independent of the order of $a$. The data produced by this method can be recovered using the function fromfile().
fid [file or str] An open file object, or a string containing a filename.
sep [str] Separator between array items for text output. If "" (empty), a binary file is written, equivalent to file.write(a.tobytes()).
format [str] Format string for text file output. Each entry in the array is formatted to text by first converting it to the closest Python type, and then using "format" \% item.

This is a convenience function for quick storage of array data. Information on endianness and precision is lost, so this method is not a good choice for files intended to archive data or transport data between machines with different endianness. Some of these problems can be overcome by outputting the data as text files, at the expense of speed and file size.

## tolist

OperatorArray.tolist()
Return the array as a (possibly nested) list.
Return a copy of the array data as a (nested) Python list. Data items are converted to the nearest compatible Python type.
none
y [list] The possibly nested list of array elements.
The array may be recreated, $a=n p . \operatorname{array}(a . t o l i s t())$.

```
>>> a = np.array([1, 2])
>>> a.tolist()
[1, 2]
>>> a = np.array([[1, 2], [3, 4]])
>>> list(a)
[array([1, 2]), array([3, 4])]
>>> a.tolist()
[[1, 2], [3, 4]]
```


## tostring

OperatorArray.tostring (order='C')
Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either 'C' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C -order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.

This function is a compatibility alias for tobytes. Despite its name it returns bytes not strings.
order [\{ 'C', ' F ', None \}, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01\x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01\x00\x00\x00\x03\x00\x00\x00'
```

trace

OperatorArray.trace (offset $=0$, axis $1=0$, axis $2=1$, dtype $=$ None, out $=$ None)
Return the sum along diagonals of the array.
Refer to numpy.trace for full documentation.
numpy.trace : equivalent function

## transpose

OperatorArray.transpose (*axes)
Returns a view of the array with axes transposed.
For a 1-D array, this has no effect. (To change between column and row vectors, first cast the 1-D array into a matrix object.) For a 2-D array, this is the usual matrix transpose. For an n-D array, if axes are given, their order indicates how the axes are permuted (see Examples). If axes are not provided and a.shape $=(i[0], i[1], \ldots i[n-2], i[n-1])$, then a.transpose(). shape $=$ (i[n-1], i[n-2], ... i[1], i[0]).
axes : None, tuple of ints, or $n$ ints

- None or no argument: reverses the order of the axes.
- tuple of ints: $i$ in the $j$-th place in the tuple means $a$ 's $i$-th axis becomes a.transpose()'s $j$-th axis.
- $n$ ints: same as an $n$-tuple of the same ints (this form is intended simply as a "convenience" alternative to the tuple form)
out [ndarray] View of $a$, with axes suitably permuted.
ndarray. T : Array property returning the array transposed.

```
>>> a = np.array([[1, 2], [3, 4]])
>>> a
array([[1, 2],
    [3, 4]])
>>> a.transpose()
array([[1, 3],
    [2, 4]])
>>> a.transpose((1, 0))
array([[1, 3],
    [2, 4]])
>>> a.transpose(1, 0)
array([[1, 3],
    [2, 4]])
```

var

OperatorArray. $\operatorname{var}($ axis $=$ None, dtype $=$ None, out $=$ None, , ddof $=0$, keepdims $=$ False )
Returns the variance of the array elements, along given axis.
Refer to numpy.var for full documentation.
numpy.var : equivalent function

## view

OperatorArray.view (dtype=None, type=None)
New view of array with the same data.
dtype [data-type or ndarray sub-class, optional] Data-type descriptor of the returned view, e.g., float32 or int16. The default, None, results in the view having the same data-type as $a$. This argument can also be specified as an ndarray sub-class, which then specifies the type of the returned object (this is equivalent to setting the type parameter).
type [Python type, optional] Type of the returned view, e.g., ndarray or matrix. Again, the default None results in type preservation.
a.view () is used two different ways:
a.view (some_dtype) or a.view (dtype=some_dtype) constructs a view of the array's memory with a different data-type. This can cause a reinterpretation of the bytes of memory.
a.view (ndarray_subclass) or a.view (type=ndarray_subclass) just returns an instance of ndarray_subclass that looks at the same array (same shape, dtype, etc.) This does not cause a reinterpretation of the memory.

For a.view (some_dtype), if some_dtype has a different number of bytes per entry than the previous dtype (for example, converting a regular array to a structured array), then the behavior of the view cannot be predicted just from the superficial appearance of a (shown by print (a)). It also depends on exactly how a is stored in memory. Therefore if a is C-ordered versus fortran-ordered, versus defined as a slice or transpose, etc., the view may give different results.

```
>>> x = np.array([(1, 2)], dtype=[('a', np.int8), ('b', np.int8)])
```

Viewing array data using a different type and dtype:

```
>>> y = x.view(dtype=np.int16, type=np.matrix)
>>> y
matrix([[513]], dtype=int16)
>>> print(type(y))
<class 'numpy.matrixlib.defmatrix.matrix'>
```

Creating a view on a structured array so it can be used in calculations

```
>>> x = np.array([(1, 2), (3,4)], dtype=[('a', np.int8), ('b', np.int8)])
>>> xv = x.view(dtype=np.int8).reshape(-1,2)
>>> xv
array([[1, 2],
    [3, 4]], dtype=int8)
>>> xv.mean(0)
array([ 2., 3.])
```

Making changes to the view changes the underlying array

```
>>> xv[0,1] = 20
>>> print(x)
[(1, 20) (3, 4)]
```

Using a view to convert an array to a recarray:

```
>>> z = x.view(np.recarray)
>>> z.a
array([1], dtype=int8)
```

Views share data:

```
>>> x[0] = (9, 10)
>>> z[0]
(9, 10)
```

Views that change the dtype size (bytes per entry) should normally be avoided on arrays defined by slices, transposes, fortran-ordering, etc.:

```
>>> x = np.array([[1,2,3],[4,5,6]], dtype=np.int16)
>>> y = x[:, 0:2]
>>> Y
array([[1, 2],
    [4, 5]], dtype=int16)
>>> y.view(dtype=[('width', np.int16), ('length', np.int16)])
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: new type not compatible with array.
>>> z = y.copy()
>>> z.view(dtype=[('width', np.int16), ('length', np.int16)])
array([[(1, 2)],
    [(4, 5)]], dtype=[('width', '<i2'), ('length', '<i2')])
```

_init__()

Initialize self. See help(type(self)) for accurate signature.

## Attributes

| T | Same as self.transpose(), except that self is returned <br> if self.ndim < 2. |
| :--- | :--- |
| base | Base object if memory is from some other object. |
| ctypes | An object to simplify the interaction of the array with <br> the ctypes module. |
| data | Python buffer object pointing to the start of the ar- <br> ray's data. |
| dtype | Data-type of the array's elements. |
| flags | Information about the memory layout of the array. |
| flat | A 1-D iterator over the array. |
| imag | The imaginary part of the array. |
| itemsize | Length of one array element in bytes. |
| nbytes | Total bytes consumed by the elements of the array. |
| ndim | Number of array dimensions. |
| real | The real part of the array. |
| shape | Tuple of array dimensions. |
| size | Number of elements in the array. |
| strides | Tuple of bytes to step in each dimension when <br> traversing an array. |

T

OperatorArray.T
Same as self.transpose(), except that self is returned if self.ndim $<2$.

```
>>> x = np.array([[1.,2.],[3.,4.]])
>>> x
array([[ 1., 2.],
    [ 3., 4.]])
>>> x.T
array([[ 1., 3.],
    [ 2., 4.]])
>>> x = np.array([1.,2.,3.,4.])
>>> x
array([ 1., 2., 3., 4.])
>>> x.T
array([ 1., 2., 3., 4.])
```

base

OperatorArray.base
Base object if memory is from some other object.
The base of an array that owns its memory is None:

```
>>> x = np.array([1,2,3,4])
>>> x.base is None
True
```

Slicing creates a view, whose memory is shared with x :

```
>>> y = x[2:]
>>> y.base is x
True
```

ctypes

## OperatorArray.ctypes

An object to simplify the interaction of the array with the ctypes module.
This attribute creates an object that makes it easier to use arrays when calling shared libraries with the ctypes module. The returned object has, among others, data, shape, and strides attributes (see Notes below) which themselves return ctypes objects that can be used as arguments to a shared library.

## None

c [Python object] Possessing attributes data, shape, strides, etc.
numpy.ctypeslib
Below are the public attributes of this object which were documented in "Guide to NumPy" (we have omitted undocumented public attributes, as well as documented private attributes):

- data: A pointer to the memory area of the array as a Python integer. This memory area may contain data that is not aligned, or not in correct byte-order. The memory area may not even be writeable. The array flags and data-type of this array should be respected when passing this attribute to arbitrary C-code to avoid trouble that can include Python crashing. User Beware! The value of this attribute is exactly the same as self._array_interface_['data'][0].
- shape (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the C-integer corresponding to dtype('p') on this platform. This base-type could be c_int, c_long, or c_longlong depending on the platform. The c_intp type is defined accordingly in numpy.ctypeslib. The ctypes array contains the shape of the underlying array.
- strides (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the same as for the shape attribute. This ctypes array contains the strides information from the underlying array. This strides information is important for showing how many bytes must be jumped to get to the next element in the array.
- data_as(obj): Return the data pointer cast to a particular c-types object. For example, calling self._as_parameter_ is equivalent to self.data_as(ctypes.c_void_p). Perhaps you want to use the data as a pointer to a ctypes array of floating-point data: self.data_as(ctypes.POINTER(ctypes.c_double)).
- shape_as(obj): Return the shape tuple as an array of some other c-types type. For example: self.shape_as(ctypes.c_short).
- strides_as(obj): Return the strides tuple as an array of some other c-types type. For example: self.strides_as(ctypes.c_longlong).

Be careful using the ctypes attribute - especially on temporary arrays or arrays constructed on the fly. For example, calling ( $a+b$ ) . ctypes.data_as (ctypes.c_void_p) returns a pointer to memory that is invalid because the array created as $(a+b)$ is deallocated before the next Python statement. You can avoid this problem using either $c=a+b$ or $c t=(a+b) . c t y p e s$. In the latter case, ct will hold a reference to the array until ct is deleted or re-assigned.

If the ctypes module is not available, then the ctypes attribute of array objects still returns something useful, but ctypes objects are not returned and errors may be raised instead. In particular, the object will still have the as parameter attribute which will return an integer equal to the data attribute.

```
>>> import ctypes
>>> x
array([[0, 1],
    [2, 3]])
>>> x.ctypes.data
30439712
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long))
<ctypes.LP_c_long object at 0x01F01300>
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long)).contents
c_long(0)
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_longlong)).contents
c_longlong(4294967296L)
>>> x.ctypes.shape
<numpy.core._internal.c_long_Array_2 object at 0x01FFD580>
>>> x.ctypes.shape_as(ctypes.c_long)
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides_as(ctypes.c_longlong)
<numpy.core._internal.c_longlong_Array_2 object at 0x01F01300>
```

data

OperatorArray.data
Python buffer object pointing to the start of the array's data.

## dtype

OperatorArray.dtype
Data-type of the array's elements.
None
d : numpy dtype object
numpy.dtype

```
>>> x
array([[0, 1],
    [2, 3]])
>>> x.dtype
dtype('int32')
>>> type(x.dtype)
<type 'numpy.dtype'>
```


## flags

OperatorArray.flags
Information about the memory layout of the array.
C_CONTIGUOUS (C) The data is in a single, C-style contiguous segment.
F_CONTIGUOUS (F) The data is in a single, Fortran-style contiguous segment.
OWNDATA (O) The array owns the memory it uses or borrows it from another object.

WRITEABLE (W) The data area can be written to. Setting this to False locks the data, making it readonly. A view (slice, etc.) inherits WRITEABLE from its base array at creation time, but a view of a writeable array may be subsequently locked while the base array remains writeable. (The opposite is not true, in that a view of a locked array may not be made writeable. However, currently, locking a base object does not lock any views that already reference it, so under that circumstance it is possible to alter the contents of a locked array via a previously created writeable view onto it.) Attempting to change a non-writeable array raises a RuntimeError exception.

ALIGNED (A) The data and all elements are aligned appropriately for the hardware.
UPDATEIFCOPY (U) This array is a copy of some other array. When this array is deallocated, the base array will be updated with the contents of this array.

FNC F_CONTIGUOUS and not C_CONTIGUOUS.
FORC F_CONTIGUOUS or C_CONTIGUOUS (one-segment test).

## BEHAVED (B) ALIGNED and WRITEABLE.

CARRAY (CA) BEHAVED and C_CONTIGUOUS.
FARRAY (FA) BEHAVED and F_CONTIGUOUS and not C_CONTIGUOUS.
The flags object can be accessed dictionary-like (as in a.flags ['WRITEABLE']), or by using lowercased attribute names (as in a.flags.writeable). Short flag names are only supported in dictionary access.

Only the UPDATEIFCOPY, WRITEABLE, and ALIGNED flags can be changed by the user, via direct assignment to the attribute or dictionary entry, or by calling ndarray.setflags.

The array flags cannot be set arbitrarily:

- UPDATEIFCOPY can only be set False.
- ALIGNED can only be set True if the data is truly aligned.
- WRITEABLE can only be set True if the array owns its own memory or the ultimate owner of the memory exposes a writeable buffer interface or is a string.

Arrays can be both C-style and Fortran-style contiguous simultaneously. This is clear for 1-dimensional arrays, but can also be true for higher dimensional arrays.

Even for contiguous arrays a stride for a given dimension arr.strides [dim] may be arbitrary if arr.shape[dim] == 1 or the array has no elements. It does not generally hold that self. strides [-1] == self.itemsize for C-style contiguous arrays or self.strides[0] == self.itemsize for Fortran-style contiguous arrays is true.
flat

OperatorArray.flat
A 1-D iterator over the array.
This is a numpy.flatiter instance, which acts similarly to, but is not a subclass of, Python's built-in iterator object.
flatten : Return a copy of the array collapsed into one dimension.
flatiter

```
>>> x = np.arange(1, 7).reshape(2, 3)
>>> x
array([[1, 2, 3],
    [4, 5, 6]])
>>> x.flat[3]
4
>>> x.T
array([[1, 4],
    [2, 5],
    [3, 6]])
>>> x.T.flat[3]
5
>>> type(x.flat)
<type 'numpy.flatiter'>
```

An assignment example:

```
>>> x.flat = 3; x
array([[3, 3, 3],
    [3, 3, 3]])
>>> x.flat[[1,4]] = 1; x
array([[3, 1, 3],
    [3, 1, 3]])
```

imag

OperatorArray.imag
The imaginary part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.imag
array([ 0. , 0.70710678])
>>> x.imag.dtype
dtype('float64')
```


## itemsize

OperatorArray.itemsize
Length of one array element in bytes.

```
>>> x = np.array([1,2,3], dtype=np.float64)
>>> x.itemsize
8
>>> x = np.array([1,2,3], dtype=np.complex128)
>>> x.itemsize
16
```


## nbytes

OperatorArray.nbytes
Total bytes consumed by the elements of the array.
Does not include memory consumed by non-element attributes of the array object.

```
>>> x = np.zeros((3,5,2), dtype=np.complex128)
>>> x.nbytes
480
>>> np.prod(x.shape) * x.itemsize
480
```

ndim

OperatorArray.ndim
Number of array dimensions.

```
>>> x = np.array([1, 2, 3])
>>> x.ndim
1
>>> y = np.zeros((2, 3, 4))
>>> y.ndim
3
```

real

OperatorArray.real
The real part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.real
array([ 1. , 0.70710678])
>>> x.real.dtype
dtype('float64')
```

numpy.real : equivalent function
shape

OperatorArray.shape
Tuple of array dimensions.
May be used to "reshape" the array, as long as this would not require a change in the total number of elements

```
>>> x = np.array([1, 2, 3, 4])
>>> x.shape
(4,)
>>> y = np.zeros((2, 3, 4))
>>> y.shape
(2, 3, 4)
>>> y.shape = (3, 8)
>>> y
array([[[ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.]])
>>> y.shape = (3, 6)
Traceback (most recent call last):
```

```
    File "<stdin>", line 1, in <module>
ValueError: total size of new array must be unchanged
```


## size

OperatorArray.size
Number of elements in the array.
Equivalent to np.prod (a.shape), i.e., the product of the array's dimensions.

```
>>> x = np.zeros((3, 5, 2), dtype=np.complex128)
>>> x.size
30
>>> np.prod(x.shape)
30
```


## strides

## OperatorArray.strides

Tuple of bytes to step in each dimension when traversing an array.
The byte offset of element (i[0], i[1], ..., $i[n]$ ) in an array $a$ is:

```
offset = sum(np.array(i) * a.strides)
```

A more detailed explanation of strides can be found in the "ndarray.rst" file in the NumPy reference guide.
Imagine an array of 32-bit integers (each 4 bytes):

```
x = np.array([[0, 1, 2, 3, 4],
    [5, 6, 7, 8, 9]], dtype=np.int 32)
```

This array is stored in memory as 40 bytes, one after the other (known as a contiguous block of memory). The strides of an array tell us how many bytes we have to skip in memory to move to the next position along a certain axis. For example, we have to skip 4 bytes ( 1 value) to move to the next column, but 20 bytes ( 5 values) to get to the same position in the next row. As such, the strides for the array $x$ will be $(20,4)$.
numpy.lib.stride_tricks.as_strided

```
>>> y = np.reshape(np.arange (2*3*4), (2,3,4))
>>> y
array([[[ 0, 1, 2, 3],
    [4, 5, 6, 7],
    [ 8, 9, 10, 11]],
    [[12, 13, 14, 15],
    [16, 17, 18, 19],
    [20, 21, 22, 23]]])
>>> y.strides
(48, 16, 4)
>>> y[1,1,1]
17
>>> offset=sum(y.strides * np.array((1,1,1)))
```

```
>>> offset/y.itemsize
```

17

```
>>> x = np.reshape(np.arange(5*6*7*8), (5, 6, 7, 8)).transpose (2, 3, 1,0)
>>> x.strides
(32, 4, 224, 1344)
>>> i = np.array ([3,5,2,2])
>>> offset = sum(i * x.strides)
>>> x[3,5,2,2]
813
>>> offset / x.itemsize
813
```


## stat

Python equivalents of statistical Excel functions.

## Functions

| xaverage |
| :--- |
| xmax |
| xmin |

xaverage

```
xaverage (*args)
```

xmax
xmax (*args)
xmin
xmin(*args)
text

Python equivalents of text Excel functions.

## Functions

| xfind |  |
| :--- | :--- |
| xleft | Continued on next page |

Table 64 - continued from previous page


## Classes

TrimArray

TrimArray
class TrimArray

## Methods

| all | Returns True if all elements evaluate to True. |
| :--- | :--- |
| any | Returns True if any of the elements of $a$ evaluate to <br> True. |
| argmax | Return indices of the maximum values along the <br> given axis. |
| argmin | Return indices of the minimum values along the <br> given axis of $a$. |
| argpartition | Returns the indices that would partition this array. |
| argsort | Returns the indices that would sort this array. |
| astype | Copy of the array, cast to a specified type. |
| byteswap | Swap the bytes of the array elements |
|  | Continued on next page |

Table 66 - continued from previous page

| choose | Use an index array to construct a new array from a set of choices. |
| :---: | :---: |
| clip | Return an array whose values are limited to [min, max]. |
| collapse |  |
| compress | Return selected slices of this array along given axis. |
| conj | Complex-conjugate all elements. |
| conjugate | Return the complex conjugate, element-wise. |
| copy | Return a copy of the array. |
| cumprod | Return the cumulative product of the elements along the given axis. |
| cumsum | Return the cumulative sum of the elements along the given axis. |
| diagonal | Return specified diagonals. |
| dot | Dot product of two arrays. |
| dump | Dump a pickle of the array to the specified file. |
| dumps | Returns the pickle of the array as a string. |
| fill | Fill the array with a scalar value. |
| flatten | Return a copy of the array collapsed into one dimension. |
| getfield | Returns a field of the given array as a certain type. |
| item | Copy an element of an array to a standard Python scalar and return it. |
| itemset | Insert scalar into an array (scalar is cast to array's dtype, if possible) |
| $\max$ | Return the maximum along a given axis. |
| mean | Returns the average of the array elements along given axis. |
| min | Return the minimum along a given axis. |
| newbyteorder | Return the array with the same data viewed with a different byte order. |
| nonzero | Return the indices of the elements that are non-zero. |
| partition | Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. |
| prod | Return the product of the array elements over the given axis |
| ptp | Peak to peak (maximum - minimum) value along a given axis. |
| put | Set a.flat $[\mathrm{n}]=$ values $[\mathrm{n}]$ for all $n$ in indices. |
| ravel | Return a flattened array. |
| repeat | Repeat elements of an array. |
| reshape | Returns an array containing the same data with a new shape. |
| resize | Change shape and size of array in-place. |
| round | Return $a$ with each element rounded to the given number of decimals. |
| searchsorted | Find indices where elements of v should be inserted in a to maintain order. |

Continued on next page

Table 66 - continued from previous page

| setfield | Put a value into a specified place in a field defined by <br> a data-type. |
| :--- | :--- |
| setflags | Set array flags WRITEABLE, ALIGNED, and UP- <br> DATEIFCOPY, respectively. |
| sort | Sort an array, in-place. |
| squeeze | Remove single-dimensional entries from the shape <br> of $a$. |
| std | Returns the standard deviation of the array elements <br> along given axis. |
| sum | Return the sum of the array elements over the given <br> axis. |
| swapaxes | Return a view of the array with axis $I$ and axis2 in- <br> terchanged. |
| take | Return an array formed from the elements of $a$ at the <br> given indices. |
| tobytes | Construct Python bytes containing the raw data bytes <br> in the array. |
| tofile | Write array to a file as text or binary (default). <br> Return the array as a (possibly nested) list. <br> tolistConstruct Python bytes containing the raw data bytes <br> in the array. |
| tostring | Return the sum along diagonals of the array. <br> traceReturns a view of the array with axes transposed. <br> given axis. |
| trariance of the array elements, along |  |
| var | New view of array with the same data. |
| view |  |

all

TrimArray.all (axis=None, out=None, keepdims=False)
Returns True if all elements evaluate to True.
Refer to numpy.all for full documentation.
numpy.all : equivalent function
any

TrimArray.any (axis=None, out=None, ,eepdims=False)
Returns True if any of the elements of $a$ evaluate to True.
Refer to numpy.any for full documentation.
numpy.any : equivalent function
argmax

TrimArray.argmax (axis=None, out=None)
Return indices of the maximum values along the given axis.
Refer to numpy.argmax for full documentation.
numpy.argmax : equivalent function

## argmin

TrimArray.argmin (axis=None, out=None)
Return indices of the minimum values along the given axis of $a$.
Refer to numpy.argmin for detailed documentation.
numpy.argmin : equivalent function

## argpartition

TrimArray. argpartition (kth, axis=-1, kind='introselect', order=None)
Returns the indices that would partition this array.
Refer to numpy.argpartition for full documentation.
New in version 1.8.0.
numpy.argpartition : equivalent function
argsort

TrimArray.argsort (axis=-1, kind='quicksort', order=None)
Returns the indices that would sort this array.
Refer to numpy.argsort for full documentation.
numpy.argsort : equivalent function
astype

TrimArray.astype (dtype, order=' $K^{\prime}$ ', casting='unsafe', subok=True, copy=True)
Copy of the array, cast to a specified type.
dtype [str or dtype] Typecode or data-type to which the array is cast.
order [ $\{$ ' C ', ' F ', ' A ', ' K '\}, optional] Controls the memory layout order of the result. ' C ' means C order, ' $F$ ' means Fortran order, ' $A$ ' means ' $F$ ' order if all the arrays are Fortran contiguous, ' $C$ ' order otherwise, and ' K ' means as close to the order the array elements appear in memory as possible. Default is ' K '.
casting [\{'no', 'equiv', 'safe', 'same_kind', 'unsafe'\}, optional] Controls what kind of data casting may occur. Defaults to 'unsafe' for backwards compatibility.

- 'no' means the data types should not be cast at all.
- 'equiv' means only byte-order changes are allowed.
- 'safe' means only casts which can preserve values are allowed.
- 'same_kind' means only safe casts or casts within a kind, like float64 to float32, are allowed.
- 'unsafe' means any data conversions may be done.
subok [bool, optional] If True, then sub-classes will be passed-through (default), otherwise the returned array will be forced to be a base-class array.
copy [bool, optional] By default, astype always returns a newly allocated array. If this is set to false, and the dtype, order, and subok requirements are satisfied, the input array is returned instead of a copy.
arr_t [ndarray] Unless copy is False and the other conditions for returning the input array are satisfied (see description for copy input parameter), arr_t is a new array of the same shape as the input array, with dtype, order given by dtype, order.

Starting in NumPy 1.9, astype method now returns an error if the string dtype to cast to is not long enough in 'safe' casting mode to hold the max value of integer/float array that is being casted. Previously the casting was allowed even if the result was truncated.

ComplexWarning When casting from complex to float or int. To avoid this, one should use a.real. astype ( t ).

```
>>> x = np.array([1, 2, 2.5])
>>> x
array([ 1. , 2. , 2.5])
```

```
>>> x.astype(int)
array([1, 2, 2])
```


## byteswap

TrimArray.byteswap (inplace)
Swap the bytes of the array elements
Toggle between low-endian and big-endian data representation by returning a byteswapped array, optionally swapped in-place.
inplace [bool, optional] If True, swap bytes in-place, default is False.
out [ndarray] The byteswapped array. If inplace is True, this is a view to self.

```
>>> A = np.array([1, 256, 8755], dtype=np.int16)
>>> map(hex, A)
['0x1', '0x100', '0x2233']
>>> A.byteswap(True)
array([ 256, 1, 13090], dtype=int16)
>>> map (hex, A)
['0x100', '0x1', '0x3322']
```

Arrays of strings are not swapped

```
>>> A = np.array(['ceg', 'fac'])
>>> A.byteswap()
array(['ceg', 'fac'],
    dtype='|S3')
```


## choose

TrimArray. choose (choices, out=None, mode='raise')
Use an index array to construct a new array from a set of choices.
Refer to numpy.choose for full documentation.
numpy.choose : equivalent function

## clip

TrimArray.clip ( min=None, max=None, out=None)
Return an array whose values are limited to [min, max]. One of max or min must be given.
Refer to numpy.clip for full documentation.
numpy.clip : equivalent function
collapse

TrimArray.collapse (shape)
compress

TrimArray. compress (condition, axis=None, out=None)
Return selected slices of this array along given axis.
Refer to numpy.compress for full documentation.
numpy.compress : equivalent function
conj

TrimArray.conj()
Complex-conjugate all elements.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function
conjugate

TrimArray. conjugate()
Return the complex conjugate, element-wise.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function
copy

TrimArray.copy (order='C')
Return a copy of the array.
order [\{ ' C ', ' F ', ' A ', ' K '\}, optional] Controls the memory layout of the copy. ' C ' means C -order, ' F ' means F -order, ' A ' means ' F ' if $a$ is Fortran contiguous, ' C ' otherwise. ' K ' means match the layout of $a$ as closely as possible. (Note that this function and :func:numpy.copy are very similar, but have different default values for their order= arguments.)
numpy.copy numpy.copyto
$\ggg x=n p \cdot \operatorname{array}([[1,2,3],[4,5,6]]$, order='F')

```
>>> y = x.copy()
```

```
>>> x.fill(0)
```

```
>>> x
array([[0, 0, 0],
    [0, 0, 0]])
```

```
>>> y
array([[1, 2, 3],
    [4, 5, 6]])
```

>>> Y.flags['C_CONTIGUOUS']
True

## cumprod

TrimArray. cumprod (axis=None, dtype=None, out=None)
Return the cumulative product of the elements along the given axis.
Refer to numpy.cumprod for full documentation.
numpy.cumprod : equivalent function

## cumsum

TrimArray. cumsum (axis=None, dtype=None, out=None)
Return the cumulative sum of the elements along the given axis.
Refer to numpy.cumsum for full documentation.
numpy.cumsum : equivalent function

## diagonal

TrimArray.diagonal (offset=0, axis $1=0$, axis $2=1$ )
Return specified diagonals. In NumPy 1.9 the returned array is a read-only view instead of a copy as in previous NumPy versions. In a future version the read-only restriction will be removed.
Refer to numpy.diagonal () for full documentation.
numpy.diagonal : equivalent function
dot

TrimArray.dot ( $b$, out=None)
Dot product of two arrays.
Refer to numpy.dot for full documentation.
numpy.dot : equivalent function

```
>>> a = np.eye(2)
>>> b = np.ones((2, 2)) * 2
>>> a.dot(b)
array([[ 2., 2.],
    [ 2., 2.]])
```

This array method can be conveniently chained:

```
>>> a.dot(b).dot(b)
array([[[ 8., 8.],
    [ 8., 8.]])
```

dump

TrimArray.dump (file)
Dump a pickle of the array to the specified file. The array can be read back with pickle.load or numpy.load.
file [str] A string naming the dump file.

## dumps

TrimArray.dumps()
Returns the pickle of the array as a string. pickle.loads or numpy.loads will convert the string back to an array.

None
fill

TrimArray.fill (value)
Fill the array with a scalar value.
value [scalar] All elements of $a$ will be assigned this value.

```
>>> a = np.array([1, 2])
>>> a.fill(0)
>>> a
array([0, 0])
>>> a = np.empty (2)
>>> a.fill(1)
>>> a
array([ 1., 1.])
```

flatten

TrimArray.flatten (order='C')
Return a copy of the array collapsed into one dimension.
order [ $\left\{{ }^{\prime} \mathrm{C}\right.$ ', ' F ', ' $A$ ', ' K '\}, optional] ' C ' means to flatten in row-major ( C -style) order. ' F ' means to flatten in column-major (Fortran- style) order. 'A' means to flatten in column-major order if $a$ is Fortran contiguous in memory, row-major order otherwise. ' K ' means to flatten $a$ in the order the elements occur in memory. The default is ' C '.
$\mathbf{y}$ [ndarray] A copy of the input array, flattened to one dimension.
ravel : Return a flattened array. flat: A 1-D flat iterator over the array.

```
>>> a = np.array([[1,2], [3,4]])
>>> a.flatten()
array([1, 2, 3, 4])
>>> a.flatten('F')
array([1, 3, 2, 4])
```


## getfield

TrimArray.getfield (dtype, offset=0)
Returns a field of the given array as a certain type.
A field is a view of the array data with a given data-type. The values in the view are determined by the given type and the offset into the current array in bytes. The offset needs to be such that the view dtype fits in the array dtype; for example an array of dtype complex 128 has 16-byte elements. If taking a view with a 32-bit integer ( 4 bytes), the offset needs to be between 0 and 12 bytes.
dtype [str or dtype] The data type of the view. The dtype size of the view can not be larger than that of the array itself.
offset [int] Number of bytes to skip before beginning the element view.

```
>>> x = np.diag([1.+1.j]*2)
>>> x[1, 1] = 2 + 4.j
>>> x
array([[ 1.+1.j, 0.+0.j],
    [ 0.+0.j, 2.+4.j]])
>>> x.getfield(np.float64)
array([[ 1., 0.],
    [ 0., 2.]])
```

By choosing an offset of 8 bytes we can select the complex part of the array for our view:

```
>>> x.getfield(np.float64, offset=8)
array([[ 1., 0.],
    [0., 4.]])
```

item

TrimArray.item(*args)
Copy an element of an array to a standard Python scalar and return it.
*args : Arguments (variable number and type)

- none: in this case, the method only works for arrays with one element (a.size $==1$ ), which element is copied into a standard Python scalar object and returned.
- int_type: this argument is interpreted as a flat index into the array, specifying which element to copy and return.
- tuple of int_types: functions as does a single int_type argument, except that the argument is interpreted as an nd-index into the array.
$\mathbf{z}$ [Standard Python scalar object] A copy of the specified element of the array as a suitable Python scalar
When the data type of $a$ is longdouble or clongdouble, item() returns a scalar array object because there is no available Python scalar that would not lose information. Void arrays return a buffer object for item(), unless fields are defined, in which case a tuple is returned.
item is very similar to a[args], except, instead of an array scalar, a standard Python scalar is returned. This can be useful for speeding up access to elements of the array and doing arithmetic on elements of the array using Python's optimized math.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.item(3)
2
>>> x.item(7)
5
>>> x.item((0, 1))
1
>>> x.item((2, 2))
3
```


## itemset

## TrimArray.itemset (*args)

Insert scalar into an array (scalar is cast to array's dtype, if possible)
There must be at least 1 argument, and define the last argument as item. Then, a.itemset (*args) is equivalent to but faster than a [args] = item. The item should be a scalar value and args must select a single item in the array $a$.
$\ * \operatorname{args}$ [Arguments] If one argument: a scalar, only used in case $a$ is of size 1. If two arguments: the last argument is the value to be set and must be a scalar, the first argument specifies a single array element location. It is either an int or a tuple.

Compared to indexing syntax, itemset provides some speed increase for placing a scalar into a particular location in an ndarray, if you must do this. However, generally this is discouraged: among other problems, it complicates the appearance of the code. Also, when using itemset (and item) inside a loop, be sure to assign the methods to a local variable to avoid the attribute look-up at each loop iteration.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.itemset (4, 0)
>>> x.itemset((2, 2), 9)
>>> x
array([[3, 1, 7],
    [2, 0, 3],
    [8, 5, 9]])
```

max

TrimArray.max (axis=None, out=None)
Return the maximum along a given axis.
Refer to numpy.amax for full documentation.
numpy.amax : equivalent function
mean

TrimArray.mean (axis=None, dtype=None, out=None, , eepdims $=$ False )
Returns the average of the array elements along given axis.
Refer to numpy.mean for full documentation.
numpy.mean : equivalent function
min

TrimArray.min(axis=None, out=None, keepdims=False)
Return the minimum along a given axis.
Refer to numpy.amin for full documentation.
numpy.amin : equivalent function
newbyteorder

TrimArray.newbyteorder (new_order='S')
Return the array with the same data viewed with a different byte order.
Equivalent to:
arr.view (arr.dtype.newbytorder(new_order))
Changes are also made in all fields and sub-arrays of the array data type.
new_order [string, optional] Byte order to force; a value from the byte order specifications below. new_order codes can be any of:

- 'S' - swap dtype from current to opposite endian
- $\left\{\right.$ ' $^{\prime}$ ', 'L' $\}$ - little endian
- $\{‘>$ ', 'B'\} - big endian
- $\left\{{ }^{\prime}=\right.$ ', ' N ' $\}$ - native order
- \{'l', 'I'\} - ignore (no change to byte order)

The default value ('S') results in swapping the current byte order. The code does a case-insensitive check on the first letter of new_order for the alternatives above. For example, any of 'B' or 'b' or 'biggish' are valid to specify big-endian.
new_arr [array] New array object with the dtype reflecting given change to the byte order.

## nonzero

TrimArray.nonzero()
Return the indices of the elements that are non-zero.
Refer to numpy.nonzero for full documentation.
numpy.nonzero : equivalent function

## partition

TrimArray.partition (kth, axis=-1, kind='introselect', order=None)
Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. All elements smaller than the kth element are moved before this element and all equal or greater are moved behind it. The ordering of the elements in the two partitions is undefined.

New in version 1.8.0.
kth [int or sequence of ints] Element index to partition by. The kth element value will be in its final sorted position and all smaller elements will be moved before it and all equal or greater elements behind it. The order all elements in the partitions is undefined. If provided with a sequence of kth it will partition all elements indexed by kth of them into their sorted position at once.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'introselect'\}, optional] Selection algorithm. Default is 'introselect'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.partition : Return a parititioned copy of an array. argpartition : Indirect partition. sort : Full sort.
See np.partition for notes on the different algorithms.

```
>>> a = np.array([3, 4, 2, 1])
>>> a.partition(a, 3)
>>> a
array([2, 1, 3, 4])
```

```
>>> a.partition((1, 3))
array([1, 2, 3, 4])
```

prod

TrimArray.prod (axis=None, dtype=None, out=None, keepdims=False)
Return the product of the array elements over the given axis
Refer to numpy.prod for full documentation.
numpy.prod : equivalent function
ptp

TrimArray.ptp (axis=None, out=None)
Peak to peak (maximum - minimum) value along a given axis.
Refer to numpy.ptp for full documentation.
numpy.ptp : equivalent function
put

TrimArray.put (indices, values, mode='raise')
Seta.flat $[\mathrm{n}]=$ values $[\mathrm{n}]$ for all $n$ in indices.
Refer to numpy.put for full documentation.
numpy.put : equivalent function
ravel

TrimArray. ravel ([order])
Return a flattened array.
Refer to numpy.ravel for full documentation.
numpy.ravel : equivalent function
ndarray.flat : a flat iterator on the array.
repeat

TrimArray.repeat (repeats, axis $=$ None)
Repeat elements of an array.
Refer to numpy.repeat for full documentation.
numpy.repeat : equivalent function
reshape

TrimArray. reshape (shape, order $=$ ' $C$ ')
Returns an array containing the same data with a new shape.
Refer to numpy.reshape for full documentation.
numpy.reshape : equivalent function
resize

TrimArray.resize (new_shape, refcheck=True)
Change shape and size of array in-place.
new_shape [tuple of ints, or $n$ ints] Shape of resized array.
refcheck [bool, optional] If False, reference count will not be checked. Default is True.

None
ValueError If $a$ does not own its own data or references or views to it exist, and the data memory must be changed.

SystemError If the order keyword argument is specified. This behaviour is a bug in NumPy.
resize : Return a new array with the specified shape.
This reallocates space for the data area if necessary.
Only contiguous arrays (data elements consecutive in memory) can be resized.
The purpose of the reference count check is to make sure you do not use this array as a buffer for another Python object and then reallocate the memory. However, reference counts can increase in other ways so if you are sure that you have not shared the memory for this array with another Python object, then you may safely set refcheck to False.
Shrinking an array: array is flattened (in the order that the data are stored in memory), resized, and reshaped:

```
>>> a = np.array([[0, 1], [2, 3]], order='C')
>>> a.resize((2, 1))
>>> a
array([[0],
    [1]])
```

```
>>> a = np.array([[0, 1], [2, 3]], order='F')
>>> a.resize((2, 1))
>>> a
array([[0],
    [2]])
```

Enlarging an array: as above, but missing entries are filled with zeros:

```
>>> b = np.array([[0, 1], [2, 3]])
>>> b.resize(2, 3) # new_shape parameter doesn't have to be a tuple
>>> b
array([[0, 1, 2],
    [3, 0, 0]])
```

Referencing an array prevents resizing...

```
>>> c=a
>>> a.resize((1, 1))
Traceback (most recent call last):
ValueError: cannot resize an array that has been referenced ...
```

Unless refcheck is False:

```
>>> a.resize((1, 1), refcheck=False)
>>> a
array([[0]])
>>> c
array([[0]])
```


## round

TrimArray. round (decimals=0, out=None)
Return $a$ with each element rounded to the given number of decimals.
Refer to numpy.around for full documentation.
numpy.around : equivalent function

## searchsorted

TrimArray. searchsorted ( $v$, side='left', sorter=None)
Find indices where elements of $v$ should be inserted in a to maintain order.
For full documentation, see numpy.searchsorted
numpy.searchsorted : equivalent function

## setfield

TrimArray.setfield(val, dtype, offset=0)
Put a value into a specified place in a field defined by a data-type.
Place val into $a$ 's field defined by dtype and beginning offset bytes into the field.
val [object] Value to be placed in field.
dtype [dtype object] Data-type of the field in which to place val.
offset [int, optional] The number of bytes into the field at which to place val.
None
getfield

```
>>> x = np.eye(3)
>>> x.getfield(np.float64)
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
>>> x.setfield(3, np.int32)
>>> x.getfield(np.int32)
array([[3, 3, 3],
    [3, 3, 3],
    [3, 3, 3]])
>>> x
array([[ 1.00000000e+000, 1.48219694e-323, 1.48219694e-323],
    [ 1.48219694e-323, 1.00000000e+000, 1.48219694e-323],
    [ 1.48219694e-323, 1.48219694e-323, 1.00000000e+000]])
>>> x.setfield(np.eye(3), np.int32)
>>> x
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
```


## setflags

TrimArray.setflags (write=None, align=None, uic=None)
Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively.
These Boolean-valued flags affect how numpy interprets the memory area used by $a$ (see Notes below). The ALIGNED flag can only be set to True if the data is actually aligned according to the type. The UPDATEIFCOPY flag can never be set to True. The flag WRITEABLE can only be set to True if the array owns its own memory, or the ultimate owner of the memory exposes a writeable buffer interface, or is a string. (The exception for string is made so that unpickling can be done without copying memory.)
write [bool, optional] Describes whether or not $a$ can be written to.
align [bool, optional] Describes whether or not $a$ is aligned properly for its type.
uic [bool, optional] Describes whether or not $a$ is a copy of another "base" array.
Array flags provide information about how the memory area used for the array is to be interpreted. There are 6 Boolean flags in use, only three of which can be changed by the user: UPDATEIFCOPY, WRITEABLE, and ALIGNED.

WRITEABLE (W) the data area can be written to;
ALIGNED (A) the data and strides are aligned appropriately for the hardware (as determined by the compiler);
UPDATEIFCOPY ( U ) this array is a copy of some other array (referenced by .base). When this array is deallocated, the base array will be updated with the contents of this array.

All flags can be accessed using their first (upper case) letter as well as the full name.

```
>>> y
array([[3, 1, 7],
    [2, 0, 0],
    [8, 5, 9]])
>>> y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : True
    ALIGNED : True
    UPDATEIFCOPY : False
>>> y.setflags(write=0, align=0)
>>> y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : False
    ALIGNED : False
    UPDATEIFCOPY : False
>>> Y.setflags(uic=1)
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: cannot set UPDATEIFCOPY flag to True
```

sort

TrimArray.sort (axis=-1, kind='quicksort', order=None)
Sort an array, in-place.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'quicksort', 'mergesort', 'heapsort'\}, optional] Sorting algorithm. Default is 'quicksort'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.sort : Return a sorted copy of an array. argsort : Indirect sort. lexsort : Indirect stable sort on multiple keys. searchsorted : Find elements in sorted array. partition: Partial sort.

See sort for notes on the different sorting algorithms.

```
>>> a = np.array([[1,4], [3,1]])
>>> a.sort(axis=1)
>>> a
array([[1, 4],
    [1, 3]])
>>> a.sort(axis=0)
>>> a
array([[1, 3],
    [1, 4]])
```

Use the order keyword to specify a field to use when sorting a structured array:

```
>>> a = np.array([('a', 2), ('c', 1)], dtype=[('x', 'S1'), ('y', int)])
>>> a.sort(order='y')
>>> a
array([('c', 1), ('a', 2)],
    dtype=[('x', '|S1'), ('y', '<i4')])
```


## squeeze

TrimArray.squeeze (axis=None)
Remove single-dimensional entries from the shape of $a$.
Refer to numpy.squeeze for full documentation.
numpy.squeeze : equivalent function
std

TrimArray.std (axis=None, dtype=None, out=None, ddof=0, keepdims=False)
Returns the standard deviation of the array elements along given axis.
Refer to numpy.std for full documentation.
numpy.std : equivalent function
sum

TrimArray.sum (axis=None, dtype=None, out=None, keepdims $=$ False )
Return the sum of the array elements over the given axis.
Refer to numpy.sum for full documentation.
numpy.sum : equivalent function

## swapaxes

TrimArray.swapaxes (axis1, axis2)
Return a view of the array with axis1 and axis 2 interchanged.
Refer to numpy.swapaxes for full documentation.
numpy.swapaxes : equivalent function

## take

TrimArray.take (indices, axis=None, out=None, mode='raise')
Return an array formed from the elements of $a$ at the given indices.
Refer to numpy.take for full documentation.
numpy.take : equivalent function

## tobytes

TrimArray.tobytes (order='C')
Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.

New in version 1.9.0.
order [ $\{$ ' $C$ ', ' $F$ ', None $\}$, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01 \x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01 \x00\x00\x00\x03\x00\x00\x00'
```

tofile

TrimArray.tofile (fid, sep="", format="\%s")
Write array to a file as text or binary (default).
Data is always written in ' C ' order, independent of the order of $a$. The data produced by this method can be recovered using the function fromfile().
fid [file or str] An open file object, or a string containing a filename.
sep [str] Separator between array items for text output. If "" (empty), a binary file is written, equivalent to file.write(a.tobytes()).
format [str] Format string for text file output. Each entry in the array is formatted to text by first converting it to the closest Python type, and then using "format" \% item.

This is a convenience function for quick storage of array data. Information on endianness and precision is lost, so this method is not a good choice for files intended to archive data or transport data between machines with different endianness. Some of these problems can be overcome by outputting the data as text files, at the expense of speed and file size.

## tolist

TrimArray.tolist()
Return the array as a (possibly nested) list.
Return a copy of the array data as a (nested) Python list. Data items are converted to the nearest compatible Python type.
none
y [list] The possibly nested list of array elements.
The array may be recreated, $a=n p . a r r a y(a . t o l i s t())$.

```
>>> a = np.array([1, 2])
>>> a.tolist()
[1, 2]
>>> a = np.array([[1, 2], [3, 4]])
>>> list(a)
[array([1, 2]), array([3, 4])]
>>> a.tolist()
[[1, 2], [3, 4]]
```


## tostring

TrimArray.tostring (order='C')
Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.

This function is a compatibility alias for tobytes. Despite its name it returns bytes not strings.
order [ $\{$ ' $C$ ', ' $F$ ', None $\}$, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01 \x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01\x00\x00\x00\x03\x00\x00\x00'
```


## trace

TrimArray.trace (offset=0, axis $1=0$, axis $2=1$, dtype $=$ None, out $=$ None )
Return the sum along diagonals of the array.
Refer to numpy.trace for full documentation.
numpy.trace : equivalent function

## transpose

TrimArray.transpose (*axes)
Returns a view of the array with axes transposed.
For a 1-D array, this has no effect. (To change between column and row vectors, first cast the 1-D array into a matrix object.) For a 2-D array, this is the usual matrix transpose. For an n-D array, if axes are given, their order indicates how the axes are permuted (see Examples). If axes are not provided and a.shape $=(i[0], i[1], \ldots i[n-2]$, $i[n-1])$, then a.transpose(). shape $=$ (i[n-1], i[n-2], ... i[1], i[0]).
axes : None, tuple of ints, or $n$ ints

- None or no argument: reverses the order of the axes.
- tuple of ints: $i$ in the $j$-th place in the tuple means $a$ 's $i$-th axis becomes a.transpose()'s $j$-th axis.
- $n$ ints: same as an $n$-tuple of the same ints (this form is intended simply as a "convenience" alternative to the tuple form)
out [ndarray] View of $a$, with axes suitably permuted.
ndarray.T : Array property returning the array transposed.

```
>>> a = np.array([[1, 2], [3, 4]])
>>> a
array([[1, 2],
    [3, 4]])
>>> a.transpose()
array([[1, 3],
    [2, 4]])
>>> a.transpose((1, 0))
array([[1, 3],
    [2, 4]])
>>> a.transpose(1, 0)
array([[1, 3],
    [2, 4]])
```

var

TrimArray.var (axis=None, dtype=None, out=None, ddof=0, keepdims=False)
Returns the variance of the array elements, along given axis.
Refer to numpy.var for full documentation.
numpy.var : equivalent function

## view

TrimArray.view $($ dtype $=$ None, type $=$ None $)$
New view of array with the same data.
dtype [data-type or ndarray sub-class, optional] Data-type descriptor of the returned view, e.g., float 32 or int16. The default, None, results in the view having the same data-type as $a$. This argument can also be specified as an ndarray sub-class, which then specifies the type of the returned object (this is equivalent to setting the type parameter).
type [Python type, optional] Type of the returned view, e.g., ndarray or matrix. Again, the default None results in type preservation.
a.view () is used two different ways:
a.view (some_dtype) or a.view (dtype=some_dtype) constructs a view of the array's memory with a different data-type. This can cause a reinterpretation of the bytes of memory.
a.view(ndarray_subclass) or a.view(type=ndarray_subclass) just returns an instance of ndarray_subclass that looks at the same array (same shape, dtype, etc.) This does not cause a reinterpretation of the memory.

For a.view (some_dtype), if some_dtype has a different number of bytes per entry than the previous dtype (for example, converting a regular array to a structured array), then the behavior of the view cannot be predicted just from the superficial appearance of a (shown by print (a)). It also depends on exactly how a is stored in memory. Therefore if a is C-ordered versus fortran-ordered, versus defined as a slice or transpose, etc., the view may give different results.

```
>>> x = np.array([(1, 2)], dtype=[('a', np.int8), ('b', np.int8)])
```

Viewing array data using a different type and dtype:

```
>>> y = x.view(dtype=np.int16, type=np.matrix)
>>> y
matrix([[513]], dtype=int16)
>>> print(type(y))
<class 'numpy.matrixlib.defmatrix.matrix'>
```

Creating a view on a structured array so it can be used in calculations

```
>>> x = np.array([(1, 2), (3,4)], dtype=[('a', np.int8), ('b', np.int8)])
>>> xv = x.view(dtype=np.int8).reshape(-1,2)
>>> xV
array([[1, 2],
    [3, 4]], dtype=int8)
>>> xv.mean(0)
array([ 2., 3.])
```

Making changes to the view changes the underlying array

```
>>> xv[0,1] = 20
>>> print(x)
[(1, 20) (3, 4)]
```

Using a view to convert an array to a recarray:

```
>>> z = x.view(np.recarray)
>>> z.a
array([1], dtype=int8)
```

Views share data:

```
>>> x[0] = (9, 10)
>>> z[0]
(9, 10)
```

Views that change the dtype size (bytes per entry) should normally be avoided on arrays defined by slices, transposes, fortran-ordering, etc.:

```
>>> x = np.array([[1,2,3],[4,5,6]], dtype=np.int16)
>>> y = x[:, 0:2]
>>> y
array([[1, 2],
    [4, 5]], dtype=int16)
>>> y.view(dtype=[('width', np.int16), ('length', np.int16)])
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: new type not compatible with array.
>>> z = y.copy()
>>> z.view(dtype=[('width', np.int16), ('length', np.int16)])
array([[(1, 2)],
    [(4, 5)]], dtype=[('width', '<i2'), ('length', '<i2')])
```

init ()

Initialize self. See help(type(self)) for accurate signature.

## Attributes

| T | Same as self.transpose(), except that self is returned <br> if self.ndim < 2. |
| :--- | :--- |
| base | Base object if memory is from some other object. |
| ctypes | An object to simplify the interaction of the array with <br> the ctypes module. |
| data | Python buffer object pointing to the start of the ar- <br> ray's data. |
| dtype | Data-type of the array's elements. |
| flags | Information about the memory layout of the array. |
| flat | A 1-D iterator over the array. |
| imag | The imaginary part of the array. |
| itemsize | Length of one array element in bytes. |
| nbytes | Total bytes consumed by the elements of the array. |
| ndim | Number of array dimensions. |
| real | The real part of the array. |
| shape | Tuple of array dimensions. |
| size | Number of elements in the array. |
| strides | Tuple of bytes to step in each dimension when <br> traversing an array. |

## T

TrimArray. T
Same as self.transpose(), except that self is returned if self.ndim $<2$.

```
>>> x = np.array([[1.,2.],[3.,4.]])
>>> x
array([[ 1., 2.],
    [ 3., 4.]])
>>> x.T
array([[ 1., 3.],
    [ 2., 4.]])
>>> x = np.array([1.,2.,3.,4.])
>>> x
array([ 1., 2., 3., 4.])
>>> x.T
array([ 1., 2., 3., 4.])
```

base

## TrimArray.base

Base object if memory is from some other object.
The base of an array that owns its memory is None:

```
>>> x = np.array([1,2,3,4])
>>> x.base is None
True
```

Slicing creates a view, whose memory is shared with x :

```
>>> y = x[2:]
>>> y.base is x
True
```

ctypes

TrimArray.ctypes
An object to simplify the interaction of the array with the ctypes module.
This attribute creates an object that makes it easier to use arrays when calling shared libraries with the ctypes module. The returned object has, among others, data, shape, and strides attributes (see Notes below) which themselves return ctypes objects that can be used as arguments to a shared library.

None
c [Python object] Possessing attributes data, shape, strides, etc.
numpy.ctypeslib
Below are the public attributes of this object which were documented in "Guide to NumPy" (we have omitted undocumented public attributes, as well as documented private attributes):

- data: A pointer to the memory area of the array as a Python integer. This memory area may contain data that is not aligned, or not in correct byte-order. The memory area may not even be writeable. The array flags and data-type of this array should be respected when passing this attribute to arbitrary C-code to avoid trouble that can include Python crashing. User Beware! The value of this attribute is exactly the same as self._array_interface_['data'][0].
- shape (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the C-integer corresponding to dtype(' p ') on this platform. This base-type could be c_int, c_long, or c_longlong
depending on the platform. The c_intp type is defined accordingly in numpy.ctypeslib. The ctypes array contains the shape of the underlying array.
- strides (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the same as for the shape attribute. This ctypes array contains the strides information from the underlying array. This strides information is important for showing how many bytes must be jumped to get to the next element in the array.
- data_as(obj): Return the data pointer cast to a particular c-types object. For example, calling self._as_parameter_ is equivalent to self.data_as(ctypes.c_void_p). Perhaps you want to use the data as a pointer to a ctypes array of floating-point data: self.data_as(ctypes.POINTER(ctypes.c_double)).
- shape_as(obj): Return the shape tuple as an array of some other c-types type. For example: self.shape_as(ctypes.c_short).
- strides_as(obj): Return the strides tuple as an array of some other c-types type. For example: self.strides_as(ctypes.c_longlong).

Be careful using the ctypes attribute - especially on temporary arrays or arrays constructed on the fly. For example, calling ( $a+b$ ) . ctypes.data_as (ctypes.c_void_p) returns a pointer to memory that is invalid because the array created as $(a+b)$ is deallocated before the next Python statement. You can avoid this problem using either $c=a+b$ or $c t=(a+b) . c t y p e s$. In the latter case, $c t$ will hold a reference to the array until ct is deleted or re-assigned.

If the ctypes module is not available, then the ctypes attribute of array objects still returns something useful, but ctypes objects are not returned and errors may be raised instead. In particular, the object will still have the as parameter attribute which will return an integer equal to the data attribute.

```
>>> import ctypes
>>> x
array([[0, 1],
    [2, 3]])
>>> x.ctypes.data
30439712
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long))
<ctypes.LP_c_long object at 0x01F01300>
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long)).contents
c_long(0)
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_longlong)).contents
c_longlong(4294967296L)
>>> x.ctypes.shape
<numpy.core._internal.c_long_Array_2 object at 0x01FFD580>
>>> x.ctypes.shape_as(ctypes.c_long)
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides_as(ctypes.c_longlong)
<numpy.core._internal.c_longlong_Array_2 object at 0x01F01300>
```

data

TrimArray.data
Python buffer object pointing to the start of the array's data.
dtype

TrimArray.dtype
Data-type of the array's elements.
None
d: numpy dtype object
numpy.dtype

```
>>> x
array([[0, 1],
    [2, 3]])
>>> x.dtype
dtype('int32')
>>> type(x.dtype)
<type 'numpy.dtype'>
```


## flags

TrimArray.flags
Information about the memory layout of the array.
C_CONTIGUOUS (C) The data is in a single, C-style contiguous segment.
F_CONTIGUOUS (F) The data is in a single, Fortran-style contiguous segment.
OWNDATA (O) The array owns the memory it uses or borrows it from another object.
WRITEABLE (W) The data area can be written to. Setting this to False locks the data, making it readonly. A view (slice, etc.) inherits WRITEABLE from its base array at creation time, but a view of a writeable array may be subsequently locked while the base array remains writeable. (The opposite is not true, in that a view of a locked array may not be made writeable. However, currently, locking a base object does not lock any views that already reference it, so under that circumstance it is possible to alter the contents of a locked array via a previously created writeable view onto it.) Attempting to change a non-writeable array raises a RuntimeError exception.

ALIGNED (A) The data and all elements are aligned appropriately for the hardware.
UPDATEIFCOPY (U) This array is a copy of some other array. When this array is deallocated, the base array will be updated with the contents of this array.
FNC F_CONTIGUOUS and not C_CONTIGUOUS.
FORC F_CONTIGUOUS or C_CONTIGUOUS (one-segment test).
BEHAVED (B) ALIGNED and WRITEABLE.
CARRAY (CA) BEHAVED and C_CONTIGUOUS.
FARRAY (FA) BEHAVED and F_CONTIGUOUS and not C_CONTIGUOUS.
The flags object can be accessed dictionary-like (as in a.flags ['WRITEABLE']), or by using lowercased attribute names (as in a.flags.writeable). Short flag names are only supported in dictionary access.

Only the UPDATEIFCOPY, WRITEABLE, and ALIGNED flags can be changed by the user, via direct assignment to the attribute or dictionary entry, or by calling ndarray.setflags.
The array flags cannot be set arbitrarily:

- UPDATEIFCOPY can only be set False.
- ALIGNED can only be set True if the data is truly aligned.
- WRITEABLE can only be set True if the array owns its own memory or the ultimate owner of the memory exposes a writeable buffer interface or is a string.

Arrays can be both C-style and Fortran-style contiguous simultaneously. This is clear for 1-dimensional arrays, but can also be true for higher dimensional arrays.

Even for contiguous arrays a stride for a given dimension arr.strides [dim] may be arbitrary if arr.shape[dim] == 1 or the array has no elements. It does not generally hold that self. strides[-1] == self.itemsize for C-style contiguous arrays or self.strides[0] == self.itemsize for Fortran-style contiguous arrays is true.
flat

TrimArray.flat
A 1-D iterator over the array.
This is a numpy.flatiter instance, which acts similarly to, but is not a subclass of, Python's built-in iterator object.
flatten : Return a copy of the array collapsed into one dimension.
flatiter

```
>>> x = np.arange(1, 7).reshape(2, 3)
>>> x
array([[1, 2, 3],
    [4, 5, 6]])
>>> x.flat[3]
4
>>> x.T
array([[1, 4],
    [2, 5],
    [3, 6]])
>>> x.T.flat[3]
5
>>> type(x.flat)
<type 'numpy.flatiter'>
```

An assignment example:

```
>>> x.flat = 3; x
array([[3, 3, 3],
    [3, 3, 3]])
>>> x.flat[[1,4]] = 1; x
array([[3, 1, 3],
    [3, 1, 3]])
```

imag

TrimArray.imag
The imaginary part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.imag
array([ 0. , 0.70710678])
>>> x.imag.dtype
dtype('float64')
```


## itemsize

TrimArray.itemsize
Length of one array element in bytes.

```
>>> x = np.array([1,2,3], dtype=np.float64)
>>> x.itemsize
8
>>> x = np.array([1,2,3], dtype=np.complex128)
>>> x.itemsize
16
```

nbytes

TrimArray.nbytes
Total bytes consumed by the elements of the array.
Does not include memory consumed by non-element attributes of the array object.

```
>>> x = np.zeros((3,5,2), dtype=np.complex128)
>>> x.nbytes
480
>>> np.prod(x.shape) * x.itemsize
480
```

ndim

TrimArray.ndim
Number of array dimensions.

```
>>> x = np.array([1, 2, 3])
>>> x.ndim
1
>>> y = np.zeros((2, 3, 4))
>>> y.ndim
3
```

real

TrimArray.real
The real part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.real
array([ 1. , 0.70710678])
>>> x.real.dtype
dtype('float64')
```

numpy.real : equivalent function
shape

TrimArray.shape
Tuple of array dimensions.
May be used to "reshape" the array, as long as this would not require a change in the total number of elements

```
>>> x = np.array([1, 2, 3, 4])
>>> x.shape
(4,)
>>> y = np.zeros((2, 3, 4))
>>> y.shape
(2, 3, 4)
>>> y.shape = (3, 8)
>>> y
array([[ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.]])
>>> y.shape = (3, 6)
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: total size of new array must be unchanged
```

size

TrimArray.size
Number of elements in the array.
Equivalent to np.prod (a.shape), i.e., the product of the array's dimensions.

```
>>> x = np.zeros((3, 5, 2), dtype=np.complex128)
>>> x.size
30
>>> np.prod(x.shape)
30
```

strides

TrimArray.strides
Tuple of bytes to step in each dimension when traversing an array.
The byte offset of element (i[0], i[1], ..., $i[n]$ ) in an array $a$ is:

```
offset = sum(np.array(i) * a.strides)
```

A more detailed explanation of strides can be found in the "ndarray.rst" file in the NumPy reference guide.
Imagine an array of 32-bit integers (each 4 bytes):

```
x = np.array([[0, 1, 2, 3, 4],
    [5, 6, 7, 8, 9]], dtype=np.int32)
```

This array is stored in memory as 40 bytes, one after the other (known as a contiguous block of memory). The strides of an array tell us how many bytes we have to skip in memory to move to the next position along a certain axis. For example, we have to skip 4 bytes ( 1 value) to move to the next column, but 20 bytes ( 5 values) to get to the same position in the next row. As such, the strides for the array $x$ will be $(20,4)$.
numpy.lib.stride_tricks.as_strided

```
>>> y = np.reshape(np.arange(2*3*4), (2,3,4))
>>> y
array([[[ 0, 1, 2, 3],
    [4, 5, 6, 7],
    [ 8, 9, 10, 11]],
    [[12, 13, 14, 15],
    [16, 17, 18, 19],
    [20, 21, 22, 23]]])
>>> y.strides
(48, 16, 4)
>>> y[1,1,1]
17
>>> offset=sum(y.strides * np.array((1,1,1)))
>>> offset/y.itemsize
17
```

```
>>> x = np.reshape(np.arange(5*6*7*8), (5, 6, 7, 8)).transpose (2, 3, 1,0)
>>> x.strides
(32, 4, 224, 1344)
>>> i = np.array ([3,5,2,2])
>>> offset = sum(i * x.strides)
>>> x[3,5,2,2]
813
>>> offset / x.itemsize
813
```


## Functions

| flatten |  |
| :--- | :--- |
| get_error |  |
| is_number |  |
| not_implemented |  |
| parse_ranges |  |
| raise_errors |  |
| replace_empty | Helps call a numpy universal function (ufunc). |
| wrap_func |  |
| wrap_ranges_func |  |

flatten
flatten $(l$, check $=<$ function is_number $>$ )
get_error
get_error (*vals)
is_number
is_number (number)
not_implemented
not_implemented (*args, **kwargs)
parse_ranges
parse_ranges (*args, **kw)
raise_errors
raise_errors(*args)
replace_empty
replace_empty $(x$, empty $=0)$
wrap_func
wrap_func (func, ranges=False)
wrap_ranges_func
wrap_ranges_func (func, n_out=1)
wrap_ufunc
wrap_ufunc (func, input_parser=<function <lambda>>, check_error=<function get_error>, args_parser $=<$ function <lambda>>, otype $=<$ function $<$ lambda>>, ranges $=$ False, **kw)
Helps call a numpy universal function (ufunc).

## Classes

Array
Array

## class Array

## Methods

| all | Returns True if all elements evaluate to True. |
| :---: | :---: |
| any | Returns True if any of the elements of $a$ evaluate to True. |
| argmax | Return indices of the maximum values along the given axis. |
| argmin | Return indices of the minimum values along the given axis of $a$. |
| argpartition | Returns the indices that would partition this array. |
| argsort | Returns the indices that would sort this array. |
| astype | Copy of the array, cast to a specified type. |
| byteswap | Swap the bytes of the array elements |
| choose | Use an index array to construct a new array from a set of choices. |
| clip | Return an array whose values are limited to [min, max]. |
| collapse |  |
| compress | Return selected slices of this array along given axis. |
| conj | Complex-conjugate all elements. |
| conjugate | Return the complex conjugate, element-wise. |
| copy | Return a copy of the array. |
| cumprod | Return the cumulative product of the elements along the given axis. |
| cumsum | Return the cumulative sum of the elements along the given axis. |
| diagonal | Return specified diagonals. |
| dot | Dot product of two arrays. |
| dump | Dump a pickle of the array to the specified file. |
| dumps | Returns the pickle of the array as a string. |
| fill | Fill the array with a scalar value. |
| flatten | Return a copy of the array collapsed into one dimension. |
| getfield | Returns a field of the given array as a certain type. |
| item | Copy an element of an array to a standard Python scalar and return it. |
| itemset | Insert scalar into an array (scalar is cast to array's dtype, if possible) |
| $\max$ | Return the maximum along a given axis. |
| mean | Returns the average of the array elements along given axis. |
| min | Return the minimum along a given axis. |
|  | Continued on next page |

Table 70 - continued from previous page

| newbyteorder | Return the array with the same data viewed with a <br> different byte order. |
| :--- | :--- |
| nonzero | Return the indices of the elements that are non-zero. |
| partition | Rearranges the elements in the array in such a way <br> that value of the element in kth position is in the po- <br> sition it would be in a sorted array. |
| prod | Return the product of the array elements over the <br> given axis |
| ptp | Peak to peak (maximum - minimum) value along a <br> given axis. |
| put | Set a.flat $[$ n] $=$ values [n] for all $n$ in in- <br> dices. |
| ravel | Return a flattened array. |
| repeat | Repeat elements of an array. <br> reshape <br> resize <br> round <br> shape. |
| searchsorted array containing the same data with a new |  |
| trace | Change shape and size of array in-place. <br> transpose the given <br> var |
| neturn $a$ with each element rounded to the of decimals. |  |

all

Array.all (axis=None, out=None, keepdims=False)
Returns True if all elements evaluate to True.

Refer to numpy.all for full documentation.
numpy.all : equivalent function

## any

Array. any (axis=None, out=None, keepdims=False)
Returns True if any of the elements of $a$ evaluate to True.
Refer to numpy.any for full documentation.
numpy.any : equivalent function

## argmax

Array. argmax (axis=None, out=None $)$
Return indices of the maximum values along the given axis.
Refer to numpy.argmax for full documentation.
numpy.argmax : equivalent function
argmin

Array.argmin (axis=None, out=None)
Return indices of the minimum values along the given axis of $a$.
Refer to numpy.argmin for detailed documentation.
numpy.argmin : equivalent function

## argpartition

Array.argpartition (kth, axis=-1, kind='introselect', order=None)
Returns the indices that would partition this array.
Refer to numpy.argpartition for full documentation.
New in version 1.8.0.
numpy.argpartition : equivalent function
argsort

Array.argsort (axis=-1, kind='quicksort', order=None)
Returns the indices that would sort this array.
Refer to numpy.argsort for full documentation.
numpy.argsort : equivalent function

## astype

Array. astype (dtype, order =' $K^{\prime}$, casting='unsafe', subok=True, copy=True)
Copy of the array, cast to a specified type.
dtype [str or dtype] Typecode or data-type to which the array is cast.
order [ $\{$ ' C ', ' F ', ' A ', ' K ' \}, optional] Controls the memory layout order of the result. ' C ' means C order, ' F ' means Fortran order, ' $A$ ' means ' F ' order if all the arrays are Fortran contiguous, ' C ' order otherwise, and ' K ' means as close to the order the array elements appear in memory as possible. Default is ' K '.
casting [\{'no', 'equiv', 'safe', 'same_kind', 'unsafe'\}, optional] Controls what kind of data casting may occur. Defaults to 'unsafe' for backwards compatibility.

- 'no' means the data types should not be cast at all.
- 'equiv' means only byte-order changes are allowed.
- 'safe' means only casts which can preserve values are allowed.
- 'same_kind' means only safe casts or casts within a kind, like float64 to float32, are allowed.
- 'unsafe' means any data conversions may be done.
subok [bool, optional] If True, then sub-classes will be passed-through (default), otherwise the returned array will be forced to be a base-class array.
copy [bool, optional] By default, astype always returns a newly allocated array. If this is set to false, and the dtype, order, and subok requirements are satisfied, the input array is returned instead of a copy.
arr_t [ndarray] Unless copy is False and the other conditions for returning the input array are satisfied (see description for copy input parameter), arr_t is a new array of the same shape as the input array, with dtype, order given by dtype, order.

Starting in NumPy 1.9, astype method now returns an error if the string dtype to cast to is not long enough in 'safe' casting mode to hold the max value of integer/float array that is being casted. Previously the casting was allowed even if the result was truncated.

ComplexWarning When casting from complex to float or int. To avoid this, one should use a. real. astype(t).

```
>>> x = np.array([1, 2, 2.5])
>>> x
array([ 1. , 2. , 2.5])
```

```
>>> x.astype(int)
array([1, 2, 2])
```


## byteswap

## Array.byteswap (inplace)

Swap the bytes of the array elements
Toggle between low-endian and big-endian data representation by returning a byteswapped array, optionally swapped in-place.
inplace [bool, optional] If True, swap bytes in-place, default is False.
out [ndarray] The byteswapped array. If inplace is True, this is a view to self.

```
>>> A = np.array([1, 256, 8755], dtype=np.int16)
>>> map(hex, A)
['0x1', '0x100', '0x2233']
>>> A.byteswap(True)
array([ 256, 1, 13090], dtype=int16)
>>> map(hex, A)
['0x100', '0x1', '0x3322']
```

Arrays of strings are not swapped

```
>>> A = np.array(['ceg', 'fac'])
>>> A.byteswap()
array(['ceg', 'fac'],
    dtype='|S3')
```


## choose

Array. choose (choices, out=None, mode='raise')
Use an index array to construct a new array from a set of choices.
Refer to numpy.choose for full documentation.
numpy.choose : equivalent function
clip

Array.clip (min=None, max=None, out=None)
Return an array whose values are limited to [min, max]. One of max or min must be given.
Refer to numpy.clip for full documentation.
numpy.clip : equivalent function
collapse

Array.collapse (shape)
compress

Array.compress (condition, axis=None, out=None)
Return selected slices of this array along given axis.
Refer to numpy.compress for full documentation.
numpy.compress : equivalent function
conj

Array.conj()
Complex-conjugate all elements.

Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function

## conjugate

Array. conjugate ()
Return the complex conjugate, element-wise.
Refer to numpy.conjugate for full documentation.
numpy.conjugate : equivalent function
copy

Array. copy (order ='C')
Return a copy of the array.
order [ $\{$ ' C ', ' F ', ' A ', ' K '\}, optional] Controls the memory layout of the copy. ' C ' means C -order, ' F ' means F-order, 'A' means ' F ' if $a$ is Fortran contiguous, ' C ' otherwise. ' K ' means match the layout of $a$ as closely as possible. (Note that this function and :func:numpy.copy are very similar, but have different default values for their order $=$ arguments.)
numpy.copy numpy.copyto

```
>>> x = np.array([[1,2,3],[4,5,6]], order='F')
```

```
>>> y = x.copy()
```

>>> x.fill(0)

```
>>> x
array([[0, 0, 0],
    [0, 0, 0]])
```

```
>>> y
array([[1, 2, 3],
    [4, 5, 6]])
```

>>> y.flags['C_CONTIGUOUS']
True
cumprod

Array. cumprod (axis=None, dtype=None, out=None)
Return the cumulative product of the elements along the given axis.
Refer to numpy.cumprod for full documentation.
numpy.cumprod : equivalent function

## cumsum

Array. cumsum (axis=None, dtype=None, out=None)
Return the cumulative sum of the elements along the given axis.
Refer to numpy.cumsum for full documentation.
numpy.cumsum : equivalent function

## diagonal

Array. diagonal (offset $=0$, axis $1=0$, axis $2=1$ )
Return specified diagonals. In NumPy 1.9 the returned array is a read-only view instead of a copy as in previous NumPy versions. In a future version the read-only restriction will be removed.

Refer to numpy. diagonal () for full documentation.
numpy.diagonal : equivalent function
dot

Array.dot ( $b$, out=None)
Dot product of two arrays.
Refer to numpy.dot for full documentation.
numpy.dot : equivalent function

```
>>> a = np.eye(2)
>>> b = np.ones((2, 2)) * 2
>>> a.dot(b)
array([[ 2., 2.],
    [ 2., 2.]])
```

This array method can be conveniently chained:

```
>>> a.dot(b).dot(b)
array([[ 8., 8.],
    [ 8., 8.]])
```

dump

Array. dump (file)
Dump a pickle of the array to the specified file. The array can be read back with pickle.load or numpy.load.
file [str] A string naming the dump file.
dumps

Array.dumps ()
Returns the pickle of the array as a string. pickle.loads or numpy.loads will convert the string back to an array.

None

## fill

Array.fill (value)
Fill the array with a scalar value.
value [scalar] All elements of $a$ will be assigned this value.

```
>>> a = np.array([1, 2])
>>> a.fill(0)
>>> a
array([0, 0])
>>> a = np.empty(2)
>>> a.fill(1)
>>> a
array([ 1., 1.])
```


## flatten

Array.flatten (order='C')
Return a copy of the array collapsed into one dimension.
order [\{ ' C ', ' F ', ' A ', ' K '\}, optional] ' C ' means to flatten in row-major (C-style) order. ' F ' means to flatten in column-major (Fortran- style) order. 'A' means to flatten in column-major order if $a$ is Fortran contiguous in memory, row-major order otherwise. ' K ' means to flatten $a$ in the order the elements occur in memory. The default is ' C '.
$\mathbf{y}$ [ndarray] A copy of the input array, flattened to one dimension.
ravel : Return a flattened array. flat: A 1-D flat iterator over the array.

```
>>> a = np.array([[1,2], [3,4]])
>>> a.flatten()
array([1, 2, 3, 4])
>>> a.flatten('F')
array([1, 3, 2, 4])
```

getfield

Array.getfield (dtype, offset=0)
Returns a field of the given array as a certain type.
A field is a view of the array data with a given data-type. The values in the view are determined by the given type and the offset into the current array in bytes. The offset needs to be such that the view dtype fits in the array dtype; for example an array of dtype complex 128 has 16-byte elements. If taking a view with a 32-bit integer ( 4 bytes), the offset needs to be between 0 and 12 bytes.
dtype [str or dtype] The data type of the view. The dtype size of the view can not be larger than that of the array itself.
offset [int] Number of bytes to skip before beginning the element view.

```
>>> x = np.diag([1.+1.j]*2)
>>> x[1, 1] = 2 + 4.j
>>> x
```

```
array([[ 1.+1.j, 0.+0.j],
    [ 0.+0.j, 2.+4.j]])
>>> x.getfield(np.float64)
array([[ 1., 0.],
    [ 0., 2.]])
```

By choosing an offset of 8 bytes we can select the complex part of the array for our view:

```
>>> x.getfield(np.float64, offset=8)
array([[ 1., 0.],
    [ 0., 4.]])
```

item

## Array.item (*args)

Copy an element of an array to a standard Python scalar and return it.
*args : Arguments (variable number and type)

- none: in this case, the method only works for arrays with one element (a.size $==1$ ), which element is copied into a standard Python scalar object and returned.
- int_type: this argument is interpreted as a flat index into the array, specifying which element to copy and return.
- tuple of int_types: functions as does a single int_type argument, except that the argument is interpreted as an nd-index into the array.
$\mathbf{z}$ [Standard Python scalar object] A copy of the specified element of the array as a suitable Python scalar
When the data type of $a$ is longdouble or clongdouble, item() returns a scalar array object because there is no available Python scalar that would not lose information. Void arrays return a buffer object for item(), unless fields are defined, in which case a tuple is returned.
item is very similar to a[args], except, instead of an array scalar, a standard Python scalar is returned. This can be useful for speeding up access to elements of the array and doing arithmetic on elements of the array using Python's optimized math.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.item(3)
2
>>> x.item(7)
5
>>> x.item((0, 1))
1
>>> x.item((2, 2))
3
```


## itemset

Array.itemset (*args)
Insert scalar into an array (scalar is cast to array's dtype, if possible)
There must be at least 1 argument, and define the last argument as item. Then, a.itemset (*args) is equivalent to but faster than a [args] = item. The item should be a scalar value and args must select a single item in the array $a$.
\*args [Arguments] If one argument: a scalar, only used in case $a$ is of size 1 . If two arguments: the last argument is the value to be set and must be a scalar, the first argument specifies a single array element location. It is either an int or a tuple.

Compared to indexing syntax, itemset provides some speed increase for placing a scalar into a particular location in an ndarray, if you must do this. However, generally this is discouraged: among other problems, it complicates the appearance of the code. Also, when using itemset (and item) inside a loop, be sure to assign the methods to a local variable to avoid the attribute look-up at each loop iteration.

```
>>> x = np.random.randint(9, size=(3, 3))
>>> x
array([[3, 1, 7],
    [2, 8, 3],
    [8, 5, 3]])
>>> x.itemset (4, 0)
>>> x.itemset((2, 2), 9)
>>> x
array([[3, 1, 7],
    [2, 0, 3],
    [8, 5, 9]])
```

max
Array.max (axis=None, out=None)

Return the maximum along a given axis.
Refer to numpy.amax for full documentation.
numpy.amax : equivalent function
mean

Array.mean (axis=None, dtype=None, out=None, keepdims=False)
Returns the average of the array elements along given axis.
Refer to numpy.mean for full documentation.
numpy.mean : equivalent function
min

Array.min (axis=None, out=None, keepdims=False)
Return the minimum along a given axis.
Refer to numpy.amin for full documentation.
numpy.amin : equivalent function

## newbyteorder

Array. newbyteorder (new_order='S')
Return the array with the same data viewed with a different byte order.
Equivalent to:
arr.view (arr.dtype.newbytorder(new_order))

Changes are also made in all fields and sub-arrays of the array data type.
new_order [string, optional] Byte order to force; a value from the byte order specifications below. new_order codes can be any of:

- 'S' - swap dtype from current to opposite endian
- \{ '<', 'L'\} - little endian
- $\{‘>$ ', 'B' $\}$ - big endian
- $\{$ ' $=$ ', ' $N$ ' $\}$ - native order
- \{'l', 'I' \} - ignore (no change to byte order)

The default value (' $S$ ') results in swapping the current byte order. The code does a case-insensitive check on the first letter of new_order for the alternatives above. For example, any of ' B ' or ' $b$ ' or 'biggish' are valid to specify big-endian.
new_arr [array] New array object with the dtype reflecting given change to the byte order.

## nonzero

## Array.nonzero()

Return the indices of the elements that are non-zero.
Refer to numpy.nonzero for full documentation.
numpy.nonzero : equivalent function

## partition

Array.partition (kth, axis=-1, kind='introselect', order=None)
Rearranges the elements in the array in such a way that value of the element in kth position is in the position it would be in a sorted array. All elements smaller than the kth element are moved before this element and all equal or greater are moved behind it. The ordering of the elements in the two partitions is undefined.

New in version 1.8.0.
kth [int or sequence of ints] Element index to partition by. The kth element value will be in its final sorted position and all smaller elements will be moved before it and all equal or greater elements behind it. The order all elements in the partitions is undefined. If provided with a sequence of kth it will partition all elements indexed by kth of them into their sorted position at once.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{ 'introselect'\}, optional] Selection algorithm. Default is 'introselect'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.partition : Return a parititioned copy of an array. argpartition : Indirect partition. sort : Full sort.
See np.partition for notes on the different algorithms.

```
>>> a = np.array([3, 4, 2, 1])
>>> a.partition(a, 3)
>>> a
array([2, 1, 3, 4])
```

```
>>> a.partition((1, 3))
```

$\operatorname{array}([1,2,3,4])$
prod

Array.prod (axis=None, dtype=None, out=None, keepdims=False)
Return the product of the array elements over the given axis
Refer to numpy.prod for full documentation.
numpy.prod : equivalent function
ptp

Array.ptp (axis=None, out=None)
Peak to peak (maximum - minimum) value along a given axis.
Refer to numpy.ptp for full documentation.
numpy.ptp : equivalent function
put

Array.put (indices, values, mode='raise')
Seta.flat [n] = values [n] for all $n$ in indices.
Refer to numpy.put for full documentation.
numpy.put : equivalent function
ravel

Array.ravel ([order])
Return a flattened array.
Refer to numpy.ravel for full documentation.
numpy.ravel : equivalent function
ndarray.flat : a flat iterator on the array.
repeat
Array.repeat (repeats, axis=None)
Repeat elements of an array.
Refer to numpy.repeat for full documentation.
numpy.repeat : equivalent function
reshape

Array. reshape (shape, order='C')
Returns an array containing the same data with a new shape.
Refer to numpy.reshape for full documentation.
numpy.reshape : equivalent function
resize

Array.resize (new_shape, refcheck=True)
Change shape and size of array in-place.
new_shape [tuple of ints, or $n$ ints] Shape of resized array.
refcheck [bool, optional] If False, reference count will not be checked. Default is True.
None
ValueError If $a$ does not own its own data or references or views to it exist, and the data memory must be changed.

SystemError If the order keyword argument is specified. This behaviour is a bug in NumPy.
resize : Return a new array with the specified shape.
This reallocates space for the data area if necessary.
Only contiguous arrays (data elements consecutive in memory) can be resized.
The purpose of the reference count check is to make sure you do not use this array as a buffer for another Python object and then reallocate the memory. However, reference counts can increase in other ways so if you are sure that you have not shared the memory for this array with another Python object, then you may safely set refcheck to False.

Shrinking an array: array is flattened (in the order that the data are stored in memory), resized, and reshaped:

```
>>> a = np.array([[0, 1], [2, 3]], order='C')
>>> a.resize((2, 1))
>>> a
array([[0],
    [1]])
```

```
>>> a = np.array([[0, 1], [2, 3]], order='F')
>>> a.resize((2, 1))
>>> a
array([[0],
    [2]])
```

Enlarging an array: as above, but missing entries are filled with zeros:

```
>>> b = np.array([[0, 1], [2, 3]])
>>> b.resize(2, 3) # new_shape parameter doesn't have to be a tuple
>>> b
array([[0, 1, 2],
    [3, 0, 0]])
```

Referencing an array prevents resizing...

```
>>> c=a
>>> a.resize((1, 1))
Traceback (most recent call last):
. .
ValueError: cannot resize an array that has been referenced ...
```

Unless refcheck is False:

```
>>> a.resize((1, 1), refcheck=False)
>>> a
array([[0]])
>>> c
array([[0]])
```

round

Array.round (decimals=0, out=None)
Return $a$ with each element rounded to the given number of decimals.
Refer to numpy.around for full documentation.
numpy.around : equivalent function

## searchsorted

Array. searchsorted ( $v$, side='left', sorter=None)
Find indices where elements of v should be inserted in a to maintain order.
For full documentation, see numpy.searchsorted
numpy.searchsorted : equivalent function

## setfield

Array.setfield (val, dtype, offset=0)
Put a value into a specified place in a field defined by a data-type.
Place val into $a$ 's field defined by dtype and beginning offset bytes into the field.
val [object] Value to be placed in field.
dtype [dtype object] Data-type of the field in which to place val.
offset [int, optional] The number of bytes into the field at which to place val.

None
getfield

```
>>> x = np.eye(3)
>>> x.getfield(np.float64)
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
>>> x.setfield(3, np.int32)
>>> x.getfield(np.int32)
array([[3, 3, 3],
    [3, 3, 3],
    [3, 3, 3]])
>>> x
array([[ 1.00000000e+000, 1.48219694e-323, 1.48219694e-323],
    [ 1.48219694e-323, 1.00000000e+000, 1.48219694e-323],
    [ 1.48219694e-323, 1.48219694e-323, 1.00000000e+000]])
>>> x.setfield(np.eye(3), np.int32)
>>> x
array([[ 1., 0., 0.],
    [ 0., 1., 0.],
    [ 0., 0., 1.]])
```

setflags

Array.setflags (write=None, align=None, uic=None)
Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively.
These Boolean-valued flags affect how numpy interprets the memory area used by $a$ (see Notes below). The ALIGNED flag can only be set to True if the data is actually aligned according to the type. The UPDATEIFCOPY flag can never be set to True. The flag WRITEABLE can only be set to True if the array owns its own memory, or the ultimate owner of the memory exposes a writeable buffer interface, or is a string. (The exception for string is made so that unpickling can be done without copying memory.)
write [bool, optional] Describes whether or not $a$ can be written to.
align [bool, optional] Describes whether or not $a$ is aligned properly for its type.
uic [bool, optional] Describes whether or not $a$ is a copy of another "base" array.
Array flags provide information about how the memory area used for the array is to be interpreted. There are 6 Boolean flags in use, only three of which can be changed by the user: UPDATEIFCOPY, WRITEABLE, and ALIGNED.

WRITEABLE (W) the data area can be written to;
ALIGNED (A) the data and strides are aligned appropriately for the hardware (as determined by the compiler);
UPDATEIFCOPY ( U ) this array is a copy of some other array (referenced by .base). When this array is deallocated, the base array will be updated with the contents of this array.

All flags can be accessed using their first (upper case) letter as well as the full name.

```
>>> y
array([[3, 1, 7],
    [2, 0, 0],
    [8, 5, 9]])
```

```
>>> y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : True
    ALIGNED : True
    UPDATEIFCOPY : False
>>> y.setflags(write=0, align=0)
>>> y.flags
    C_CONTIGUOUS : True
    F_CONTIGUOUS : False
    OWNDATA : True
    WRITEABLE : False
    ALIGNED : False
    UPDATEIFCOPY : False
>>> Y.setflags(uic=1)
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: cannot set UPDATEIFCOPY flag to True
```


## sort

Array.sort (axis=-1, kind='quicksort', order=None)
Sort an array, in-place.
axis [int, optional] Axis along which to sort. Default is -1 , which means sort along the last axis.
kind [\{'quicksort', 'mergesort', 'heapsort'\}, optional] Sorting algorithm. Default is 'quicksort'.
order [str or list of str, optional] When $a$ is an array with fields defined, this argument specifies which fields to compare first, second, etc. A single field can be specified as a string, and not all fields need be specified, but unspecified fields will still be used, in the order in which they come up in the dtype, to break ties.
numpy.sort : Return a sorted copy of an array. argsort : Indirect sort. lexsort : Indirect stable sort on multiple keys. searchsorted : Find elements in sorted array. partition: Partial sort.

See sort for notes on the different sorting algorithms.

```
>>> a = np.array([[1,4], [3,1]])
>>> a.sort(axis=1)
>>> a
array([[1, 4],
    [1, 3]])
>>> a.sort(axis=0)
>>> a
array([[1, 3],
        [1, 4]])
```

Use the order keyword to specify a field to use when sorting a structured array:

```
>> a = np.array([('a', 2), ('c', 1)], dtype=[('x', 'S1'), ('y', int)])
>>> a.sort(order='y')
>>> a
array([('c', 1), ('a', 2)],
    dtype=[('x', '|S1'), ('y', '<i4')])
```


## squeeze

Array.squeeze (axis=None)
Remove single-dimensional entries from the shape of $a$.
Refer to numpy.squeeze for full documentation.
numpy.squeeze : equivalent function
std

Array.std (axis=None, dtype=None, out=None, ddof=0, keepdims=False)
Returns the standard deviation of the array elements along given axis.
Refer to numpy.std for full documentation.
numpy.std : equivalent function
sum

Array.sum (axis=None, dtype=None, out=None, keepdims=False)
Return the sum of the array elements over the given axis.
Refer to numpy.sum for full documentation.
numpy.sum : equivalent function
swapaxes

Array.swapaxes (axis1, axis2)
Return a view of the array with axis1 and axis2 interchanged.
Refer to numpy.swapaxes for full documentation.
numpy.swapaxes : equivalent function

## take

Array.take (indices, axis=None, out=None, mode='raise')
Return an array formed from the elements of $a$ at the given indices.
Refer to numpy.take for full documentation.
numpy.take : equivalent function

## tobytes

Array.tobytes (order='C')
Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.

New in version 1.9.0.
order [ $\{$ ' $C$ ', ' $F$ ', None $\}$, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01\x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x01\x00\x00\x00\x03\x00\x00\x00'
```


## tofile

Array.tofile (fid, sep="", format="\%s")
Write array to a file as text or binary (default).
Data is always written in ' $C$ ' order, independent of the order of $a$. The data produced by this method can be recovered using the function fromfile().
fid [file or str] An open file object, or a string containing a filename.
sep [str] Separator between array items for text output. If "" (empty), a binary file is written, equivalent to file.write (a.tobytes()).
format [str] Format string for text file output. Each entry in the array is formatted to text by first converting it to the closest Python type, and then using "format" \% item.

This is a convenience function for quick storage of array data. Information on endianness and precision is lost, so this method is not a good choice for files intended to archive data or transport data between machines with different endianness. Some of these problems can be overcome by outputting the data as text files, at the expense of speed and file size.

## tolist

Array.tolist()
Return the array as a (possibly nested) list.
Return a copy of the array data as a (nested) Python list. Data items are converted to the nearest compatible Python type.
none
$\mathbf{y}$ [list] The possibly nested list of array elements.
The array may be recreated, $a=n p . \operatorname{array}(a . t o l i s t())$.

```
>>> a = np.array([1, 2])
>>> a.tolist()
[1, 2]
>>> a = np.array([[1, 2], [3, 4]])
>>> list(a)
[array([1, 2]), array([3, 4])]
>>> a.tolist()
[[1, 2], [3, 4]]
```


## tostring

Array.tostring (order='C')
Construct Python bytes containing the raw data bytes in the array.
Constructs Python bytes showing a copy of the raw contents of data memory. The bytes object can be produced in either ' C ' or 'Fortran', or 'Any' order (the default is ' C '-order). 'Any' order means C-order unless the F_CONTIGUOUS flag in the array is set, in which case it means 'Fortran' order.

This function is a compatibility alias for tobytes. Despite its name it returns bytes not strings.
order [ $\left\{{ }^{\prime} C\right.$ ', ' $F$ ', None $\}$, optional] Order of the data for multidimensional arrays: C, Fortran, or the same as for the original array.
$\mathbf{s}$ [bytes] Python bytes exhibiting a copy of $a$ 's raw data.

```
>>> x = np.array([[0, 1], [2, 3]])
>>> x.tobytes()
b'\x00\x00\x00\x00\x01\x00\x00\x00\x02\x00\x00\x00\x03\x00\x00\x00'
>>> x.tobytes('C') == x.tobytes()
True
>>> x.tobytes('F')
b'\x00\x00\x00\x00\x02\x00\x00\x00\x0 \ \x00\x00\x00\x03\x00\x00\x00'
```

trace

Array.trace $($ offset $=0$, axis $1=0$, axis $2=1$, dtype $=$ None, out $=$ None $)$
Return the sum along diagonals of the array.
Refer to numpy.trace for full documentation.
numpy.trace : equivalent function

## transpose

## Array.transpose (*axes)

Returns a view of the array with axes transposed.
For a 1-D array, this has no effect. (To change between column and row vectors, first cast the 1-D array into a matrix object.) For a 2-D array, this is the usual matrix transpose. For an n-D array, if axes are given, their order indicates how the axes are permuted (see Examples). If axes are not provided anda.shape $=(i[0], i[1], \ldots i[n-2], i[n-1])$, then a.transpose(). shape $=$ (i[n-1], i[n-2], ... i[1], i[0]).
axes : None, tuple of ints, or $n$ ints

- None or no argument: reverses the order of the axes.
- tuple of ints: $i$ in the $j$-th place in the tuple means $a$ 's $i$-th axis becomes a.transpose()'s $j$-th axis.
- $n$ ints: same as an n-tuple of the same ints (this form is intended simply as a "convenience" alternative to the tuple form)
out [ndarray] View of $a$, with axes suitably permuted.
ndarray. T : Array property returning the array transposed.

```
>>> a = np.array([[1, 2], [3, 4]])
>>> a
array([[1, 2],
    [3, 4]])
>>> a.transpose()
array([[1, 3],
    [2, 4]])
>>> a.transpose((1, 0))
array([[1, 3],
    [2, 4]])
>>> a.transpose(1, 0)
array([[1, 3],
    [2, 4]])
```

var

Array.var $($ axis $=$ None, dtype=None, out=None, ddof=0, keepdims $=$ False $)$
Returns the variance of the array elements, along given axis.
Refer to numpy.var for full documentation.
numpy.var : equivalent function
view

Array.view (dtype=None, type=None)
New view of array with the same data.
dtype [data-type or ndarray sub-class, optional] Data-type descriptor of the returned view, e.g., float 32 or int16. The default, None, results in the view having the same data-type as $a$. This argument can also be specified as an ndarray sub-class, which then specifies the type of the returned object (this is equivalent to setting the type parameter).
type [Python type, optional] Type of the returned view, e.g., ndarray or matrix. Again, the default None results in type preservation.
a.view () is used two different ways:
a.view (some_dtype) or a.view (dtype=some_dtype) constructs a view of the array's memory with a different data-type. This can cause a reinterpretation of the bytes of memory.
a.view (ndarray_subclass) or a.view (type=ndarray_subclass) just returns an instance of ndarray_subclass that looks at the same array (same shape, dtype, etc.) This does not cause a reinterpretation of the memory.

For a.view (some_dtype), if some_dtype has a different number of bytes per entry than the previous dtype (for example, converting a regular array to a structured array), then the behavior of the view cannot be predicted just from the superficial appearance of a (shown by print (a)). It also depends on exactly how a is stored in memory. Therefore if a is C-ordered versus fortran-ordered, versus defined as a slice or transpose, etc., the view may give different results.

```
>>> x = np.array([(1, 2)], dtype=[('a', np.int8), ('b', np.int8)])
```

Viewing array data using a different type and dtype:

```
>>> y = x.view(dtype=np.int16, type=np.matrix)
>>> y
matrix([[513]], dtype=int16)
>>> print(type(y))
<class 'numpy.matrixlib.defmatrix.matrix'>
```

Creating a view on a structured array so it can be used in calculations

```
>>> x = np.array([(1, 2), (3,4)], dtype=[('a', np.int8), ('b', np.int8)])
>>> xv = x.view(dtype=np.int8).reshape(-1,2)
>>> xv
array([[1, 2],
    [3, 4]], dtype=int8)
>>> xv.mean(0)
array([ 2., 3.])
```

Making changes to the view changes the underlying array

```
>>> xv[0,1] = 20
>>> print(x)
[(1, 20) (3, 4)]
```

Using a view to convert an array to a recarray:

```
>>> z = x.view(np.recarray)
>>> z.a
array([1], dtype=int8)
```

Views share data:

```
>>> x[0] = (9, 10)
>>> z[0]
(9, 10)
```

Views that change the dtype size (bytes per entry) should normally be avoided on arrays defined by slices, transposes, fortran-ordering, etc.:

```
>>> x = np.array([[1,2,3],[4,5,6]], dtype=np.int16)
>>> y = x[:, 0:2]
>>> y
array([[1, 2],
    [4, 5]], dtype=int16)
>>> y.view(dtype=[('width', np.int16), ('length', np.int16)])
Traceback (most recent call last):
    File "<stdin>", line 1, in <module>
ValueError: new type not compatible with array.
>>> z = y.copy()
>>> z.view(dtype=[('width', np.int16), ('length', np.int16)])
array([[(1, 2)],
    [(4, 5)]], dtype=[('width', '<i2'), ('length', '<i2')])
```

_init__()

Initialize self. See help(type(self)) for accurate signature.

## Attributes

| T | Same as self.transpose(), except that self is returned <br> if self.ndim < 2. |
| :--- | :--- |
| base | Base object if memory is from some other object. |
| ctypes | An object to simplify the interaction of the array with <br> the ctypes module. |
| data | Python buffer object pointing to the start of the ar- <br> ray's data. |
| dtype | Data-type of the array's elements. |
| flags | Information about the memory layout of the array. |
| flat | A 1-D iterator over the array. |
| imag | The imaginary part of the array. |
| itemsize | Length of one array element in bytes. |
| nbytes | Total bytes consumed by the elements of the array. |
| ndim | Number of array dimensions. |
| real | The real part of the array. |
| shape | Tuple of array dimensions. |
| size | Number of elements in the array. |
| strides | Tuple of bytes to step in each dimension when <br> traversing an array. |

T

Array. $\mathbf{T}$
Same as self.transpose(), except that self is returned if self.ndim $<2$.

```
>>> x = np.array([[1.,2.],[3.,4.]])
>>> x
array([[ 1., 2.],
    [ 3., 4.]])
>>> x.T
array([[ 1., 3.],
    [ 2., 4.]])
>>> x = np.array([1.,2.,3.,4.])
>>> x
array([ 1., 2., 3., 4.])
>>> x.T
array([ 1., 2., 3., 4.])
```

base

Array.base
Base object if memory is from some other object.
The base of an array that owns its memory is None:

```
>>> x = np.array([1,2,3,4])
>>> x.base is None
True
```

Slicing creates a view, whose memory is shared with x :

```
>>> y = x[2:]
>>> y.base is x
True
```


## ctypes

## Array.ctypes

An object to simplify the interaction of the array with the ctypes module.
This attribute creates an object that makes it easier to use arrays when calling shared libraries with the ctypes module. The returned object has, among others, data, shape, and strides attributes (see Notes below) which themselves return ctypes objects that can be used as arguments to a shared library.

## None

c [Python object] Possessing attributes data, shape, strides, etc.
numpy.ctypeslib
Below are the public attributes of this object which were documented in "Guide to NumPy" (we have omitted undocumented public attributes, as well as documented private attributes):

- data: A pointer to the memory area of the array as a Python integer. This memory area may contain data that is not aligned, or not in correct byte-order. The memory area may not even be writeable. The array flags and data-type of this array should be respected when passing this attribute to arbitrary C-code to avoid trouble that can include Python crashing. User Beware! The value of this attribute is exactly the same as self._array_interface_['data'][0].
- shape (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the C-integer corresponding to dtype(' p ') on this platform. This base-type could be c_int, c_long, or c_longlong depending on the platform. The c_intp type is defined accordingly in numpy.ctypeslib. The ctypes array contains the shape of the underlying array.
- strides (c_intp*self.ndim): A ctypes array of length self.ndim where the basetype is the same as for the shape attribute. This ctypes array contains the strides information from the underlying array. This strides information is important for showing how many bytes must be jumped to get to the next element in the array.
- data_as(obj): Return the data pointer cast to a particular c-types object. For example, calling self._as_parameter_ is equivalent to self.data_as(ctypes.c_void_p). Perhaps you want to use the data as a pointer to a ctypes array of floating-point data: self.data_as(ctypes.POINTER(ctypes.c_double)).
- shape_as(obj): Return the shape tuple as an array of some other c-types type. For example: self.shape_as(ctypes.c_short).
- strides_as(obj): Return the strides tuple as an array of some other c-types type. For example: self.strides_as(ctypes.c_longlong).

Be careful using the ctypes attribute - especially on temporary arrays or arrays constructed on the fly. For example, calling $(a+b)$. ctypes.data_as (ctypes.c_void_p) returns a pointer to memory that is invalid because the array created as $(a+b)$ is deallocated before the next Python statement. You can avoid this problem using either $c=a+b$ or $c t=(a+b) . c t y p e s$. In the latter case, ct will hold a reference to the array until ct is deleted or re-assigned.

If the ctypes module is not available, then the ctypes attribute of array objects still returns something useful, but ctypes objects are not returned and errors may be raised instead. In particular, the object will still have the as parameter attribute which will return an integer equal to the data attribute.

```
>>> import ctypes
>>> x
array([[0, 1],
    [2, 3]])
>>> x.ctypes.data
30439712
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long))
<ctypes.LP_c_long object at 0x01F01300>
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_long)).contents
c_long(0)
>>> x.ctypes.data_as(ctypes.POINTER(ctypes.c_longlong)).contents
c_longlong(4294967296L)
>>> x.ctypes.shape
<numpy.core._internal.c_long_Array_2 object at 0x01FFD580>
>>> x.ctypes.shape_as(ctypes.c_long)
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides
<numpy.core._internal.c_long_Array_2 object at 0x01FCE620>
>>> x.ctypes.strides_as(ctypes.c_longlong)
<numpy.core._internal.c_longlong_Array_2 object at 0x01F01300>
```

data

Array.data
Python buffer object pointing to the start of the array's data.
dtype

Array.dtype
Data-type of the array's elements.
None
d : numpy dtype object
numpy.dtype

```
>>> x
array([[0, 1],
    [2, 3]])
>>> x.dtype
dtype('int32')
>>> type(x.dtype)
<type 'numpy.dtype'>
```


## flags

## Array.flags

Information about the memory layout of the array.
C_CONTIGUOUS (C) The data is in a single, C-style contiguous segment.
F_CONTIGUOUS (F) The data is in a single, Fortran-style contiguous segment.
OWNDATA (O) The array owns the memory it uses or borrows it from another object.

WRITEABLE (W) The data area can be written to. Setting this to False locks the data, making it readonly. A view (slice, etc.) inherits WRITEABLE from its base array at creation time, but a view of a writeable array may be subsequently locked while the base array remains writeable. (The opposite is not true, in that a view of a locked array may not be made writeable. However, currently, locking a base object does not lock any views that already reference it, so under that circumstance it is possible to alter the contents of a locked array via a previously created writeable view onto it.) Attempting to change a non-writeable array raises a RuntimeError exception.

ALIGNED (A) The data and all elements are aligned appropriately for the hardware.
UPDATEIFCOPY ( $\mathbf{U}$ ) This array is a copy of some other array. When this array is deallocated, the base array will be updated with the contents of this array.

FNC F_CONTIGUOUS and not C_CONTIGUOUS.
FORC F_CONTIGUOUS or C_CONTIGUOUS (one-segment test).

## BEHAVED (B) ALIGNED and WRITEABLE.

CARRAY (CA) BEHAVED and C_CONTIGUOUS.
FARRAY (FA) BEHAVED and F_CONTIGUOUS and not C_CONTIGUOUS.
The flags object can be accessed dictionary-like (as in a.flags ['WRITEABLE']), or by using lowercased attribute names (as in a.flags.writeable). Short flag names are only supported in dictionary access.

Only the UPDATEIFCOPY, WRITEABLE, and ALIGNED flags can be changed by the user, via direct assignment to the attribute or dictionary entry, or by calling ndarray.setflags.

The array flags cannot be set arbitrarily:

- UPDATEIFCOPY can only be set False.
- ALIGNED can only be set True if the data is truly aligned.
- WRITEABLE can only be set True if the array owns its own memory or the ultimate owner of the memory exposes a writeable buffer interface or is a string.

Arrays can be both C-style and Fortran-style contiguous simultaneously. This is clear for 1-dimensional arrays, but can also be true for higher dimensional arrays.

Even for contiguous arrays a stride for a given dimension arr.strides [dim] may be arbitrary if arr.shape[dim] == 1 or the array has no elements. It does not generally hold that self. strides [-1] == self.itemsize for C-style contiguous arrays or self.strides[0] == self.itemsize for Fortran-style contiguous arrays is true.
flat

## Array.flat

A 1-D iterator over the array.
This is a numpy.flatiter instance, which acts similarly to, but is not a subclass of, Python's built-in iterator object.
flatten : Return a copy of the array collapsed into one dimension.
flatiter

```
>>> x = np.arange(1, 7).reshape(2, 3)
>>> x
array([[1, 2, 3],
    [4, 5, 6]])
>>> x.flat[3]
4
>>> x.T
array([[1, 4],
    [2, 5],
    [3, 6]])
>>> x.T.flat[3]
5
>>> type(x.flat)
<type 'numpy.flatiter'>
```

An assignment example:

```
>>> x.flat = 3; x
array([[3, 3, 3],
    [3, 3, 3]])
>>> x.flat[[1,4]] = 1; x
array([[3, 1, 3],
    [3, 1, 3]])
```

imag

## Array.imag

The imaginary part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.imag
array([ 0. , 0.70710678])
>>> x.imag.dtype
dtype('float64')
```


## itemsize

## Array.itemsize

Length of one array element in bytes.

```
>>> x = np.array([1,2,3], dtype=np.float64)
>>> x.itemsize
8
>>> x = np.array([1,2,3], dtype=np.complex128)
>>> x.itemsize
16
```

nbytes

Array.nbytes
Total bytes consumed by the elements of the array.
Does not include memory consumed by non-element attributes of the array object.

```
>>> x = np.zeros((3,5,2), dtype=np.complex128)
>>> x.nbytes
480
>>> np.prod(x.shape) * x.itemsize
480
```

ndim

Array.ndim
Number of array dimensions.

```
>>> x = np.array([1, 2, 3])
>>> x.ndim
1
>>> y = np.zeros((2, 3, 4))
>>> y.ndim
3
```

real

Array.real
The real part of the array.

```
>>> x = np.sqrt([1+0j, 0+1j])
>>> x.real
array([ 1. , 0.70710678])
>>> x.real.dtype
dtype('float64')
```

numpy.real : equivalent function
shape

Array.shape
Tuple of array dimensions.
May be used to "reshape" the array, as long as this would not require a change in the total number of elements

```
>>> x = np.array([1, 2, 3, 4])
>>> x.shape
(4,)
>>> y = np.zeros((2, 3, 4))
>>> y.shape
(2, 3, 4)
>>> y.shape = (3, 8)
>>> y
array([[[ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.],
    [ 0., 0., 0., 0., 0., 0., 0., 0.]])
>>> y.shape = (3, 6)
Traceback (most recent call last):
```

```
File "<stdin>", line 1, in <module>
ValueError: total size of new array must be unchanged
```


## size

Array.size
Number of elements in the array.
Equivalent to np.prod (a.shape), i.e., the product of the array's dimensions.

```
>>> x = np.zeros((3, 5, 2), dtype=np.complex128)
>>> x.size
30
>>> np.prod(x.shape)
30
```


## strides

## Array.strides

Tuple of bytes to step in each dimension when traversing an array.
The byte offset of element (i[0], i[1], ..., $i[n]$ ) in an array $a$ is:

```
offset = sum(np.array(i) * a.strides)
```

A more detailed explanation of strides can be found in the "ndarray.rst" file in the NumPy reference guide.
Imagine an array of 32-bit integers (each 4 bytes):
$x=n p \cdot \operatorname{array}([[0,1,2,3,4]$,
$[5,6,7,8,9]]$, dtype=np.int 32)
This array is stored in memory as 40 bytes, one after the other (known as a contiguous block of memory). The strides of an array tell us how many bytes we have to skip in memory to move to the next position along a certain axis. For example, we have to skip 4 bytes ( 1 value) to move to the next column, but 20 bytes ( 5 values) to get to the same position in the next row. As such, the strides for the array $x$ will be $(20,4)$.
numpy.lib.stride_tricks.as_strided

```
>>> y = np.reshape(np.arange (2*3*4), (2, 3, 4))
>>> y
array([[[ 0, 1, 2, 3],
    [4, 5, 6, 7],
    [ 8, 9, 10, 11]],
    [[12, 13, 14, 15],
    [16, 17, 18, 19],
    [20, 21, 22, 23]]])
>>> y.strides
(48, 16, 4)
>>> y[1,1,1]
17
>>> offset=sum(y.strides * np.array((1,1,1)))
```

>>> offset/y.itemsize
17

```
>>> x = np.reshape(np.arange(5*6*7*8), (5, 6, 7, 8)).transpose (2, 3, 1,0)
>>> x.strides
(32, 4, 224, 1344)
>>> i = np.array([3,5,2,2])
>>> offset = sum(i * x.strides)
>>> x[3,5,2,2]
813
>>> offset / x.itemsize
813
```

reshape (shape, order='C')
Returns an array containing the same data with a new shape.
Refer to numpy.reshape for full documentation.
numpy.reshape : equivalent function

### 2.1.7.6 ranges

It provides Ranges class.

## Classes

## Ranges

## Ranges

class Ranges (ranges=(), values=None, is_set=False, all_values=True)

## Methods

| _init___range | Initialize self. |
| :--- | :--- |
| format_range |  |
| get_rang |  |
| push |  |
| pushes |  |
| set_value |  |
| simplify |  |

__init__

Ranges.__init__(ranges=(), values=None, is_set=False, all_values=True) Initialize self. See help(type(self)) for accurate signature.

```
format_range
static Ranges.format_range(*args, **kwargs)
get_range
static Ranges.get_range (format_range,ref, context=None)
push
Ranges.push (ref,value=empty, context=None)
pushes
Ranges.pushes(refs, values=(), context=None)
set_value
Ranges.set_value(rng,value=empty)
simplify
Ranges.simplify()
__init__ (ranges=(), values=None, is_set=False, all_values=True)
    Initialize self. See help(type(self)) for accurate signature.
```


## Attributes

input_fields
value
input_fields
Ranges.input_fields = ('excel', 'sheet', 'n1', 'n2', 'r1', 'r2')
value
Ranges.value

### 2.1.7.7 cell

It provides Cell class.

## Functions

| format_output |
| :--- |
| wrap_cell_func |
| format_output |
| format_output (rng, value) |
| wrap_cell_func |
| wrap_cell_func (func, parse_args=<function <lambda>>, parse_kwargs=<function <lambda>>) |
| Classes |


| Cell |
| :--- |
| CellWrapper |
| RangesAssembler |

## Cell

class Cell (reference, value, context $=$ None)

## Methods

| _init__ | Initialize self. |
| :--- | :--- |
| add |  |
| compile |  |
| update_inputs |  |

__init__

Cell.__init__(reference, value, context=None)
Initialize self. See help(type(self)) for accurate signature.
add

Cell.add (dsp, context=None)
compile

Cell.compile(references=None)

## update_inputs

Cell.update_inputs (references=None)
__init__(reference, value, context=None)
Initialize self. See help(type(self)) for accurate signature.

## Attributes

$\qquad$
output

Cell. output

## CellWrapper

class CellWrapper (func, parse_args, parse_kwargs)

## Methods

| _init___cycles | Initialize self. |
| :--- | :--- |

$\qquad$

CellWrapper.__init__(func, parse_args, parse_kwargs)
Initialize self. See help(type(self)) for accurate signature.
check_cycles

CellWrapper.check_cycles (cycle)
__init__(func, parse_args, parse_kwargs)
Initialize self. See help(type(self)) for accurate signature.

RangesAssembler
class RangesAssembler (ref, context=None)

## Methods

| pinit_ | Initialize self. |
| :--- | :--- |
| push |  |

RangesAssembler.__init__(ref, context=None)
Initialize self. See help(type(self)) for accurate signature.
push

RangesAssembler.push (cell)
__init__(ref, context=None)
Initialize self. See help(type(self)) for accurate signature.

## Attributes

## output

output

RangesAssembler.output

### 2.1.7.8 excel

It provides Excel model class.

## Classes

## ExcelModel

## ExcelModel

## class ExcelModel

## Methods

| init__ | Initialize self. |
| :--- | :--- |
| add_book |  |
| add_cell |  |
| add_sheet |  |
| compile |  |
| complete |  |
| load |  |
| push |  |
| pushes |  |

Table 83 - continued from previous page
solve_circular
write
$\qquad$

ExcelModel.__init__()
Initialize self. See help(type(self)) for accurate signature.
add_book

ExcelModel.add_book(book, context=None, data_only=False)
add_cell

ExcelModel.add_cell(cell, context, references=None, formula_references=None, formula_ranges $=$ None, external_links=None)
add_sheet

ExcelModel.add_sheet (worksheet, context)
compile

ExcelModel. compile (inputs, outputs)
complete

ExcelModel.complete()
finish

ExcelModel.finish (complete=True, circular=False)
load

ExcelModel.load (filename)
loads

ExcelModel.loads (*file_names)
push

ExcelModel.push (worksheet, context)

```
pushes
ExcelModel.pushes(*worksheets, context=None)
solve_circular
ExcelModel.solve_circular()
write
ExcelModel.write (books=None, solution=None)
__init__()
    Initialize self. See help(type(self)) for accurate signature.
compile_class
        alias of schedula.utils.dsp.DispatchPipe
```


### 2.1.8 Changelog

### 2.1.8.1 v0.1.4 (2018-10-19)

Fix

- (tokens) \#20: Improve Number regex.


### 2.1.8.2 v0.1.3 (2018-10-09)

Feat

- (excel) \#16: Solve circular references.
- (setup): Add donate url.

Fix

- (functions) \#18: Enable check_error in IF function just for the first argument.
- (functions) \#18: Disable input_parser in IF function to return any type of values.
- (rtd): Define fpath from prj_dir for rtd.
- (rtd): Add missing requirements openpyxl for rtd.
- (setup): Patch to use sphinxcontrib.restbuilder in setup long_description.


## Other

- Update documentation.
- Replace excel with Excel.
- Create PULL_REQUEST_TEMPLATE.md.
- Update issue templates.
- Update copyright.
- (doc): Update author mail.


### 2.1.8.3 v0.1.2 (2018-09-12)

Feat

- (functions) \#14: Add ROW and COLUMN.
- (cell): Pass cell reference when compiling cell + new function struct with dict to add inputs like CELL.

Fix

- (ranges): Replace system max size with excel max row and col.
- (tokens): Correct number regex.


### 2.1.8.4 v0.1.1 (2018-09-11)

## Feat

- (contrib): Add contribution instructions.
- (setup): Add additional project_urls.
- (setup): Update Development Status to 4 - Beta.

Fix

- (init) \#15: Replace FUNCTIONS and OPERATORS objs with get_functions, SUBMODULES.
- (doc): Correct link docs_status.


### 2.1.8.5 v0.1.0 (2018-07-20)

Feat

- (readme) \#6, \#7: Add examples.
- (doc): Add changelog.
- (test): Add info of executed test of test_excel_model.
- (functions) \#11: Add HEX2OCT, HEX2BIN, HEX2DEC, OCT2HEX, OCT2BIN, OCT2DEC, BIN2HEX, BIN2OCT, BIN2DEC, DEC2HEX, DEC2OCT, and DEC2BIN functions.
- (setup) \#13: Add extras_require to setup file.

Fix

- (excel): Use DispatchPipe to compile a sub model of excel workbook.
- (range) \#11: Correct range regex to avoid parsing of function like ranges (e.g., HEX2DEC).
2.1.8.6 v0.0.10 (2018-06-05)

Feat

- (look): Simplify _get_type_id function.

Fix

- (functions): Correct ImportError for FUNCTIONS.
- (operations): Correct behaviour of the basic operations.
2.1.8.7 v0.0.9 (2018-05-28)

Feat

- (excel): Improve performances pre-calculating the range format.
- (core): Improve performances using DispatchPipe instead SubDispatchPipe when compiling formulas.
- (function): Improve performances setting errstate outside vectorization.
- (core): Improve performances of range2parts function (overall $50 \%$ faster).

Fix

- (ranges): Minimize conversion str to int and vice versa.
- (functions) \#10: Avoid returning shapeless array.


### 2.1.8.8 v0.0.8 (2018-05-23)

Feat

- (functions): Add MATCH, LOOKUP, HLOOKUP, VLOOKUP functions.
- (excel): Add method to compile ExcelModel.
- (travis): Run coveralls in python 3.6.
- (functions): Add FIND,'LEFT‘,'LEN‘,'LOWER','MID‘,'REPLACE‘,'RIGHT', 'TRIM‘, and‘UPPER‘ functions.
- (functions): Add $I R R$ function.
- (formulas): Custom reshape to Array class.
- (functions): Add ISO.CEILING, SQRTPI, TRUNC functions.
- (functions): Add ROUND, ROUNDDOWN, ROUNDUP, SEC, SECH, SIGN functions.
- (functions): Add DECIMAL, EVEN, MROUND, ODD, RAND, RANDBETWEEN functions.
- (functions): Add FACT and FACTDOUBLE functions.
- (functions): Add ARABIC and ROMAN functions.
- (functions): Parametrize function wrap_ufunc.
- (functions): Split function raise_errors adding get_error function.
- (ranges): Add custom default and error value for defining ranges Arrays.
- (functions): Add LOG10 function + fix $L O G$.
- (functions): Add CSC and CSCH functions.
- (functions): Add COT and COTH functions.
- (functions): Add FLOOR, FLOOR.MATH, and FLOOR.PRECISE functions.
- (test): Improve log message of test cell.

Fix

- (rtd): Update installation file for read the docs.
- (functions): Remove unused functions.
- (formulas): Avoid too broad exception.
- (functions.math): Drop scipy dependency for calculate factorial2.
- (functions.logic): Correct error behaviour of if and iferror functions + add BroadcastError.
- (functions.info): Correct behaviour of iserr function.
- (functions): Correct error behaviour of average function.
- (functions): Correct iserror and iserr returning a custom Array.
- (functions): Now xceiling function returns np.nan instead Error.errors['\#NUM!'].
- (functions): Correct is_number function, now returns False when number is a bool.
- (test): Ensure same order of workbook comparisons.
- (functions): Correct behaviour of min max and int function.
- (ranges): Ensure to have a value with correct shape.
- (parser): Change order of parsing to avoid TRUE and FALSE parsed as ranges or errors as strings.
- (function):Remove unused kwargs n_out.
- (parser): Parse error string as formulas.
- (readme): Remove downloads_count because it is no longer available.


## Other

- Refact: Update Copyright + minor pep.
- Excel returns 1-indexed string positions???
- Added common string functions.
- Merge pull request \#9 from ecatkins/irr.
- Implemented IRR function using numpy.


### 2.1.8.9 v0.0.7 (2017-07-20)

Feat

- (appveyor): Add python 3.6.
- (functions) \#4: Add sumproduct function.

Fix

- (install): Force update setuptools>=36.0.1.
- (functions): Correct iserror iserr functions.
- (ranges): Replace '\#N/A' with '’ as empty value when assemble values.
- (functions) \#4: Remove check in ufunc when inputs have different size.
- (functions) \#4: Correct power, arctan 2 , and mod error results.
- (functions) \#4: Simplify ufunc code.
- (test) \#4: Check that all results are in the output.
- (functions) \#4: Correct atan2 argument order.
- (range) \#5: Avoid parsing function name as range when it is followed by (.
- (operator) \#3: Replace strip with replace.
- (operator) \#3: Correct valid operators like ${ }^{\wedge}$ - or ${ }^{*}+$.


## Other

- Made the ufunc wrapper work with multi input functions, e.g., power, mod, and atan2.
- Created a workbook comparison method in TestExcelModel.
- Added MIN and MAX to the test.xlsx.
- Cleaned up the ufunc wrapper and added min and max to the functions list.
- Relaxed equality in TestExcelModel and made some small fixes to functions.py.
- Added a wrapper for numpy ufuncs, mapped some Excel functions to ufuncs and provided tests.


### 2.1.8.10 v0.0.6 (2017-05-31)

Fix

- (plot): Update schedula to 0.1.12.
- (range): Sheet name without commas has this [^Wd][w.] format.


### 2.1.8.11 v0.0.5 (2017-05-04)

Fix

- (doc): Update schedula to 0.1.11.


### 2.1.8.12 v0.0.4 (2017-02-10)

Fix

- (regex): Remove deprecation warnings.


### 2.1.8.13 v0.0.3 (2017-02-09)

Fix

- (appveyor): Setup of lxml.
- (excel): Remove deprecation warning openpyxl.
- (requirements): Update schedula requirement 0.1.9.


### 2.1.8.14 v0.0.2 (2017-02-08)

Fix

- (setup): setup fails due to long description.
- (excel): Remove deprecation warning remove_sheet $\rightarrow$ remove.


## chapter 3

Indices and tables

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