# **Guide to NumPy**

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# Contents

Ι	Nι	ımPy	from Python	11
1	Orig	gins of	f NumPy	12
2	Obj	ect Es	ssentials	17
	2.1	Data-	Type Descriptors	18
	2.2	Basic	indexing (slicing)	22
	2.3	Memo	ory Layout of ndarray	24
		2.3.1	Contiguous Memory Layout	25
		2.3.2	Discontiguous memory layout	26
	2.4	Unive	rsal Functions for arrays	29
	2.5	Summ	nary of new features	30
	2.6	Summ	nary of differences with Numeric	32
		2.6.1	The list of necessary changes:	33
		2.6.2	Updating code that uses Numeric using convertcode	35
		2.6.3	Recommended changes	35
3	The	Array	y Object	37
	3.1	ndarr	ray Object Attributes	37
		3.1.1	Memory Layout attributes	37
		3.1.2	Data Type attributes	41
		3.1.3	Other attributes	42
		3.1.4	Array Interface attributes	43
	3.2	ndarr	ay Methods	44
		3.2.1	Array conversion	44
		3.2.2	Array shape manipulation	48
		3.2.3	Array item selection and manipulation	51
		3.2.4	Array calculation	55
	3.3	Array	Special Methods	59
		3.3.1	Methods for standard library functions	59

3.3.2       Basic customization       61         3.3.3       Container customization       63         3.3.4       Arithmetic customization       64         3.3.4.1       Binary       64         3.3.4.2       In-place       66         3.3.4.3       Unary operations       67         3.3.4.4       Array indexing       67         3.3.5       Basic Slicing       68         3.3.6       Advanced selection       70         3.3.6.1       Integer       70         3.3.6.2       Boolean       72         3.3.7       Flat Iterator indexing       73         4       Basic Routines       74         4.1       Creating arrays       74         4.2       Operations on two or more arrays       78         4.3       Printing arrays       80         4.4       Functions redundant with methods       81         4.5       Dealing with data types       82
3.3.4 Arithmetic customization       64         3.3.4.1 Binary       64         3.3.4.2 In-place       66         3.3.4.3 Unary operations       67         3.3.4.4 Array indexing       67         3.3.5 Basic Slicing       68         3.3.6 Advanced selection       70         3.3.6.1 Integer       70         3.3.6.2 Boolean       72         3.3.7 Flat Iterator indexing       73         4 Basic Routines       74         4.1 Creating arrays       74         4.2 Operations on two or more arrays       78         4.3 Printing arrays       80         4.4 Functions redundant with methods       81
3.3.4.2 In-place       66         3.3.4.3 Unary operations       67         3.3.4.4 Array indexing       67         3.3.5 Basic Slicing       68         3.3.6 Advanced selection       70         3.3.6.1 Integer       70         3.3.6.2 Boolean       72         3.3.7 Flat Iterator indexing       73         4 Basic Routines       74         4.1 Creating arrays       74         4.2 Operations on two or more arrays       78         4.3 Printing arrays       80         4.4 Functions redundant with methods       81
3.3.4.3 Unary operations 67 3.3.4.4 Array indexing 67 3.3.5 Basic Slicing 68 3.3.6 Advanced selection 70 3.3.6.1 Integer 70 3.3.6.2 Boolean 72 3.3.7 Flat Iterator indexing 73 4 Basic Routines 74 4.1 Creating arrays 74 4.2 Operations on two or more arrays 78 4.3 Printing arrays 80 4.4 Functions redundant with methods 81
3.3.4.3 Unary operations       67         3.3.4.4 Array indexing       67         3.3.5 Basic Slicing       68         3.3.6 Advanced selection       70         3.3.6.1 Integer       70         3.3.6.2 Boolean       72         3.3.7 Flat Iterator indexing       73         4 Basic Routines       74         4.1 Creating arrays       74         4.2 Operations on two or more arrays       78         4.3 Printing arrays       80         4.4 Functions redundant with methods       81
3.3.5       Basic Slicing       68         3.3.6       Advanced selection       70         3.3.6.1       Integer       70         3.3.6.2       Boolean       72         3.3.7       Flat Iterator indexing       73         4       Basic Routines       74         4.1       Creating arrays       74         4.2       Operations on two or more arrays       78         4.3       Printing arrays       80         4.4       Functions redundant with methods       81
3.3.5       Basic Slicing       68         3.3.6       Advanced selection       70         3.3.6.1       Integer       70         3.3.6.2       Boolean       72         3.3.7       Flat Iterator indexing       73         4       Basic Routines       74         4.1       Creating arrays       74         4.2       Operations on two or more arrays       78         4.3       Printing arrays       80         4.4       Functions redundant with methods       81
3.3.6.1 Integer       70         3.3.6.2 Boolean       72         3.3.7 Flat Iterator indexing       73         4 Basic Routines       74         4.1 Creating arrays       74         4.2 Operations on two or more arrays       78         4.3 Printing arrays       80         4.4 Functions redundant with methods       81
3.3.6.1 Integer       70         3.3.6.2 Boolean       72         3.3.7 Flat Iterator indexing       73         4 Basic Routines       74         4.1 Creating arrays       74         4.2 Operations on two or more arrays       78         4.3 Printing arrays       80         4.4 Functions redundant with methods       81
3.3.7 Flat Iterator indexing       73         4 Basic Routines       74         4.1 Creating arrays       74         4.2 Operations on two or more arrays       78         4.3 Printing arrays       80         4.4 Functions redundant with methods       81
4 Basic Routines       74         4.1 Creating arrays       74         4.2 Operations on two or more arrays       78         4.3 Printing arrays       80         4.4 Functions redundant with methods       81
4 Basic Routines       74         4.1 Creating arrays       74         4.2 Operations on two or more arrays       78         4.3 Printing arrays       80         4.4 Functions redundant with methods       81
4.1 Creating arrays744.2 Operations on two or more arrays784.3 Printing arrays804.4 Functions redundant with methods81
4.2 Operations on two or more arrays
4.3 Printing arrays
4.4 Functions redundant with methods 81
4.5 Dealing with data types
4.9 Deaning with data types
5 Additional Convenience Routines 84
5.1 Shape functions
5.2 Basic functions
5.3 Polynomial functions
5.4 Set Operations
5.5 Array construction using index tricks
5.6 Other indexing devices
5.7 Two-dimensional functions
5.8 More data type functions
5.9 Functions that behave like ufuncs
5.10 Miscellaneous Functions
6 Scalar objects 108
6.1 Attributes of array scalars
6.2 Methods of array scalars
6.3 Defining New Types

7	Dat	a-type	e Objects	113
	7.1	Attrib	outes	. 113
	7.2	Const	ruction	. 115
	7.3	Metho	ods	. 119
8	Sta	ndard	Classes	121
	8.1	Specia	al attributes and methods recognized by NumPy	. 122
	8.2	Matri	x Objects	. 122
	8.3	Memo	ory-mapped-file arrays	. 125
	8.4	Chara	acter arrays (numpy.char)	. 126
	8.5	Recor	d Arrays (numpy.rec)	. 127
	8.6	Maske	ed Arrays (numpy.ma)	. 130
	8.7	UserA	array	. 130
	8.8	Array	Iterators	. 131
		8.8.1	Default iteration	. 131
		8.8.2	Flat iteration	. 131
		8.8.3	N-dimensional enumeration	. 132
		8.8.4	Iterator for broadcasting	. 132
9	Uni	versal	Functions	134
	9.1		iption	. 134
	0.1	9.1.1	Broadcasting	
		9.1.2	Output type determination	
		9.1.3	Use of internal buffers	
		9.1.4	Error handling	
	9.2	0	attributes	
	9.3		ng Rules	
	9.4		Object methods	
	0.1	9.4.1	Reduce	
		9.4.2	Accumulate	
		9.4.3	Reduceat	
		9.4.4	Outer	
	9.5	_	able ufuncs	
	5.0	9.5.1	Math operations	
		9.5.2	Trigonometric functions	
		9.5.2 $9.5.3$	Bit-twiddling functions	
		9.5.4	Comparison functions	
		9.5.5	Floating functions	. 191

<b>10</b>	Basi	ic Mod	lules	153
	10.1	Linear	${\bf Algebra} \; ({\tt linalg}) \; \ldots \; $	153
	10.2	Discret	te Fourier Transforms (dft)	155
	10.3	Rando	m Numbers (random)	159
		10.3.1	Discrete Distributions	160
		10.3.2	Continuous Distributions	162
		10.3.3	Miscellaneous utilities	169
11	Test	ing an	d Packaging	170
	11.1	Testing	g	170
		,	y Distutils	
			misc_util	
		11.2.2	Other modules	179
	11.3	Conver	rsion of .src files	180
		11.3.1	Fortran files	181
			11.3.1.1 Named repeat rule	181
			11.3.1.2 Short repeat rule	
			11.3.1.3 Pre-defined names	181
		11.3.2	Other files	182
II	$\mathbf{C}$	-API		183
			on Types and C-Structures	183 184
	New	y Pyth		184
	New	v <b>Pyth</b> New P	on Types and C-Structures	<b>184</b> 185
	New	New P	on Types and C-Structures  Tython Types Defined	184 185 185
	New	New P 12.1.1 12.1.2	on Types and C-Structures  ython Types Defined	184 185 185 187
	New	New P 12.1.1 12.1.2 12.1.3	on Types and C-Structures  ython Types Defined	184 185 185 187 193
	New	New P 12.1.1 12.1.2 12.1.3 12.1.4	on Types and C-Structures  Tython Types Defined	184 185 185 187 193 196
	New	New P 12.1.1 12.1.2 12.1.3 12.1.4 12.1.5	on Types and C-Structures  Tython Types Defined	184 185 185 187 193 196 197
	New	New P 12.1.1 12.1.2 12.1.3 12.1.4 12.1.5 12.1.6	on Types and C-Structures  Tython Types Defined  PyArray_Type  PyArrayDescr_Type  PyUFunc_Type  PyArrayIter_Type  PyArrayMultiIter_Type	184 185 185 187 193 196 197
	New 12.1	New P 12.1.1 12.1.2 12.1.3 12.1.4 12.1.5 12.1.6 12.1.7	on Types and C-Structures  ython Types Defined  PyArray_Type  PyArrayDescr_Type  PyUFunc_Type  PyArrayIter_Type  PyArrayMultiIter_Type  PyArrayFlags_Type	184 185 185 187 193 196 197 198
	New 12.1	New P 12.1.1 12.1.2 12.1.3 12.1.4 12.1.5 12.1.6 12.1.7 Other	on Types and C-Structures  Tython Types Defined  PyArray_Type  PyArrayDescr_Type  PyUFunc_Type  PyArrayIter_Type  PyArrayMultiIter_Type  PyArrayFlags_Type  ScalarArrayTypes	184 185 187 193 196 197 198 198
	New 12.1	New P 12.1.1 12.1.2 12.1.3 12.1.4 12.1.5 12.1.6 12.1.7 Other 12.2.1	on Types and C-Structures  ython Types Defined  PyArray_Type  PyArrayDescr_Type  PyUFunc_Type  PyArrayIter_Type  PyArrayMultiIter_Type  PyArrayFlags_Type  ScalarArrayTypes  C-Structures	184 185 187 193 196 197 198 198
	New 12.1	New P 12.1.1 12.1.2 12.1.3 12.1.4 12.1.5 12.1.6 12.1.7 Other 12.2.1 12.2.2	on Types and C-Structures  Tython Types Defined  PyArray_Type  PyArrayDescr_Type  PyUFunc_Type  PyArrayIter_Type  PyArrayMultiIter_Type  PyArrayFlags_Type  ScalarArrayTypes  C-Structures  PyArray_Dims	184 185 187 193 196 197 198 199 199
	New 12.1	New P 12.1.1 12.1.2 12.1.3 12.1.4 12.1.5 12.1.6 12.1.7 Other 12.2.1 12.2.2 12.2.3	on Types and C-Structures  Tython Types Defined  PyArray_Type  PyArrayDescr_Type  PyUFunc_Type  PyArrayIter_Type  PyArrayMultiIter_Type  PyArrayFlags_Type  ScalarArrayTypes  C-Structures  PyArray_Dims  PyArray_Chunk	184 185 187 193 196 197 198 199 199 200
	New 12.1	New P 12.1.1 12.1.2 12.1.3 12.1.4 12.1.5 12.1.6 12.1.7 Other 12.2.1 12.2.2 12.2.3	on Types and C-Structures  Tython Types Defined  PyArray_Type  PyArrayDescr_Type  PyUFunc_Type  PyArrayIter_Type  PyArrayMultiIter_Type  PyArrayFlags_Type  ScalarArrayTypes  C-Structures  PyArray_Dims  PyArray_Chunk  PyArrayInterface	184 185 187 193 196 197 198 199 199 200 201
	New 12.1	New P 12.1.1 12.1.2 12.1.3 12.1.4 12.1.5 12.1.6 12.1.7 Other 12.2.1 12.2.2 12.2.3	on Types and C-Structures  Tython Types Defined  PyArray_Type  PyArrayDescr_Type  PyUFunc_Type  PyArrayIter_Type  PyArrayMultiIter_Type  PyArrayFlags_Type  ScalarArrayTypes  C-Structures  PyArray_Dims  PyArray_Interface  Internally used structures	184 185 187 193 196 197 198 199 199 200 201 201

13	Con	$\mathbf{plete}$	API		203
	13.1	Config	uration d	efines	. 203
		13.1.1	Guarant	eed to be defined $\dots$	. 203
		13.1.2	Possible	defines	. 204
	13.2	Array	Data Typ	es	. 204
		13.2.1	Enumera	ated Types	. 205
		13.2.2	Defines		. 205
			13.2.2.1	Max and min values for integers	. 205
			13.2.2.2	Number of bits in data types	. 206
			13.2.2.3	Bit-width references to enumerated typenums	. 206
			13.2.2.4	Integer that can hold a pointer	. 206
		13.2.3	C-type n	ames	. 207
			13.2.3.1	Boolean	. 207
			13.2.3.2	(Un)Signed Integer	. 207
			13.2.3.3	(Complex) Floating point	. 207
			13.2.3.4	Bit-width names	. 208
		13.2.4	Printf Fo	ormatting	. 208
	13.3	Array	API		. 208
		13.3.1	Array st	ructure and data access	. 208
			13.3.1.1	Data access	. 210
		13.3.2	Creating	arrays	. 210
			13.3.2.1	From scratch	. 210
			13.3.2.2	From other objects	. 213
		13.3.3	Dealing	with types	. 218
			13.3.3.1	General check of Python Type	. 218
			13.3.3.2	Data-type checking	. 219
			13.3.3.3	Converting data types	. 221
			13.3.3.4	New data types	. 224
			13.3.3.5	Special functions for PyArray_OBJECT	. 225
		13.3.4	Array fla	gs	. 225
			13.3.4.1	Basic Array Flags	. 225
			13.3.4.2	Combinations of array flags	. 226
			13.3.4.3	Flag-like constants	. 226
			13.3.4.4	Flag checking	. 226
		13.3.5	Array m	ethod alternative API	. 228
			13.3.5.1	Conversion	. 228
			13.3.5.2	Shape Manipulation	. 230
			13.3.5.3	Item selection and manipulation $\dots \dots$ .	. 231

			13.3.5.4 Calculation	233
		13.3.6	Functions	235
			13.3.6.1 Array Functions	235
			13.3.6.2 Other functions	237
		13.3.7	Array Iterators	237
		13.3.8	Broadcasting (multi-iterators)	238
		13.3.9	Array Scalars	239
		13.3.10	Data-type descriptors	241
		13.3.11	Conversion Utilities	243
			13.3.11.1 For use with PyArg_ParseTuple $\ \ldots \ \ldots \ \ldots$	243
			13.3.11.2 Other conversions	244
		13.3.12	? Miscellaneous	245
			13.3.12.1 Importing the API	245
			13.3.12.2 Internal Flexibility	246
			13.3.12.3 Memory management	247
			13.3.12.4 Threading support	248
			13.3.12.5 Priority	249
			13.3.12.6 Default buffers	249
			13.3.12.7 Other constants	250
			13.3.12.8 Miscellaneous Macros	250
			13.3.12.9 Enumerated Types	251
	13.4	UFunc	API	251
		13.4.1	Constants	251
		13.4.2	Macros	252
		13.4.3	Functions	252
		13.4.4	Generic functions	254
	13.5	Import	ing the API	256
14	How	to ex	tend NumPy	258
	14.1	Writing	g an extension module	258
			ed subroutine	
		-		260
				260
			· · ·	262
				262
	14.4			264
				264
			Creating a brand-new ndarray	
			V	

		14.4.3	Getting at ndarray memory and accessing elements of the	
			ndarray	8
	14.5	Examp	ple	6
<b>15</b>	Bey	$\mathbf{ond}  \mathbf{th}$	ne Basics 27	′1
	15.1	Iterati	ng over elements in the array	1
		15.1.1	Basic Iteration	1
		15.1.2	Iterating over all but one axis	2
		15.1.3	Iterating over multiple arrays	′3
		15.1.4	Broadcasting over multiple arrays	<b>'</b> 4
	15.2	Creati	ng a new universal function	<b>'</b> 4
	15.3	User-d	lefined data-types	7
		15.3.1	Adding the new data-type	78
		15.3.2	Registering a casting function	8
		15.3.3	Registering coercion rules	'9
		15.3.4	Registering a ufunc loop	30
	15.4	Subtyp	ping the ndarray in C	30
		15.4.1	Creating sub-types	31
		15.4.2	Specific features of ndarray sub-typing 28	32
			15.4.2.1 The _array_finalize attribute 28	32
			15.4.2.2 The _array_priority_ attribute 28	3
			15.4.2.3 The _array_wrap_ attribute	3
16	Thi	rd-part	ty tools 28	4
	16.1	Calling	g other compiled libraries from Python	34
		16.1.1	Hand-generated wrappers	35
		16.1.2	Using f2py	35
	16.2	Other	tools installed separately	35
		16.2.1	Using weave	35
		16.2.2	Using PyRex	35
		16.2.3	Using ctypes	35
		16.2.4	Other tools	35
			16.2.4.1 SWIG	35
			16.2.4.2 Boost	35
			16.2.4.3 SIP	35
			16.2.4.4 PyFort	35
			16.2.4.5 Instant	35

7 Code Explanations		
17.1 Code generation		286
17.2 Array Scalars		286
17.3 N-d Array Iteration		286
17.4 Advanced Indexing		286
17.5 Universal Functions		286

# List of Tables

2.1	Built-in array-scalar types corresponding to data-types for an ndar-	
	ray. The bold-face types correspond to standard Python types. The	
	object_type is special because arrays with dtype='O' do not return	
	an array scalar on item access but instead return the actual object	
	referenced in the array	21
3.1	Attributes of the ndarray	38
3.2	Array conversion methods	48
3.3	Array item selection and shape manipulation methods. If axis is an	
	argument, then the calculation is performed along that axis. An axis	
	value of None means the array is flattened before calculation proceeds.	56
3.4	Array object calculation methods. If axis is an argument, then the	
	calculation is performed along that axis. An axis value of None means	
	the array is flattened before calculation proceeds	60
6.1	Array scalar types that inherit from basic Python types. The into	
	array data type might also inherit from the IntType if it has the same	
	number of bits as the int_ array data type on your platform	108
9.1	Universal function (ufunc) attributes	138
10.1	Functions in numpy dual (both in NumPy and SciPy)	153

# Part I NumPy from Python

# Chapter 1

# Origins of NumPy

NumPy builds on (and is a successor to) the successful Numeric array object. Its goal is to create a useful environment for scientific computing. In order to better understand the people surrounding NumPy and (its library-package) SciPy, I will explain a little about how SciPy and NumPy originated. In 1998, as a graduate student studying biomedical imaging at the Mayo Clinic in Rochester, MN, I came across Python and its numerical extension (Numeric) while I was looking for ways to analyze large data sets for Magnetic Resonance Imaging and Ultrasound using a high-level language. I quickly fell in love with Python programming which is a remarkable statement to make about a programming language. If I had not seen others with the same view, I might have seriously doubted my sanity. I became rather involved in the Numeric Python community, adding the C-API chapter to the Numeric documentation (for which Paul Dubois graciously made me a co-author).

As I progressed with my thesis work, programming in Python was so enjoyable that I felt inhibited when I worked with other programming frameworks. As a result, when a task I needed to perform was not available in the core language, or in the Numeric extension, I looked around and found C or Fortran code that performed the needed task, wrapped it into Python (either by hand or using SWIG), and used the new functionality in my programs.

Along the way, I learned a great deal about the underlying structure of Numeric and grew to admire it's simple but elegant structures that grew out of the mechanism by which Python allows itself to be extended.



## NOTE

Numeric was originally written in 1995 largely by Jim Hugunin while he was a graduate student at MIT. He received help from many people including Jim Fulton, David Ascher, Paul DuBois, and Konrad Hinsen. These individuals and many others added comments, criticisms, and code which helped the Numeric extension reach stability. Jim Hugunin did not stay long as an active member of the community — moving on to write Jython and, later, Iron Python.

By operating in this need-it-make-it fashion I ended up with a substantial library of extension modules that helped Python + Numeric become easier to use in a scientific setting. These early modules included raw input-output functions, a special function library, an integration library, an ordinary differential equation solver, some least-squares optimizers, and sparse matrix solvers. While I was doing this laborious work, Pearu Peterson noticed that a lot of the routines I was wrapping were written in Fortran and there was no simplified wrapping mechanism for Fortran subroutines (like SWIG for C). He began the task of writing f2py which made it possible to easily wrap Fortran programs into Python. I helped him a little bit, mostly with testing and contributing early function-call-back code, but he put forth the brunt of the work. His result was simply amazing to me. I've always been impressed with f2py, especially because I knew how much effort writing and maintaining extension modules could be. Anybody serious about scientific computing with Python will appreciate that f2py is distributed along with NumPy.

When I finished my Ph.D. in 2001, Eric Jones (who had recently completed his Ph.D. at Duke) contacted me because he had a collection of Python modules he had developed as part of his thesis work as well. He wanted to combine his modules with mine into one super package. Together with Pearu Peterson we joined our efforts, and SciPy was born in 2001. Since then, many people have contributed module code to SciPy including Fernando Perez, Prabhu Ramachandran, Charles Harris, David Cooke, Gary Strangman, and Jean-Sebastien Roy. Others such as Travis Vaught, David Morrill, Jeff Whitaker, and Louis Luangkesorn have contributed testing and build support.

At the start of 2005, SciPy was at release 0.3 and relatively stable for an early version number. Part of the reason it was difficult to stabilize SciPy was that the array object upon which SciPy builds was undergoing a bit of an upheveal. At about the same time as SciPy was being built, some Numeric users were hitting up against the limited capabilities of Numeric. In particular, the ability to deal with memory

mapped files (and associated alignment and swapping issues), record arrays, and altered error checking modes were important but limited or non-existent in Numeric. As a result, numarray was created by Perry Greenfield, Todd Miller, and Rick White at the Space Science Telescope Institute as a replacement for Numeric. Numarray used a very different implementation scheme as a mix of Python classes and C code (which led to extreme slow downs in certain common uses). While improving some capabilities, it was slow to pick up on the more advanced features of Numeric's universal functions (ufuncs) — never re-creating the C-API that SciPy depended on. This made it difficult for SciPy to "convert" to numarray.

Many newcomers to scientific computing with Python were told that numarray was the future and started developing for it. Very useful tools were developed that could not be used with Numeric (because of numarray's change in C-API), and therefore could not be used easily in SciPy. This state of affairs was very discouraging for me personally as it left the community fragmented. Some developed for numarray, others developed as part of SciPy. A few people even rejected adopting Python for scientific computing entirely because of the split. In addition, I estimate that quite a few Python users simply stayed away from both SciPy and numarray, leaving the community smaller than it could have been given the number of people that use Python for science and engineering purposes.

It should be recognized that the split was not intentional, but simply an outgrowth of the different and exacting demands of scientific computing users. My describing these events should not be construed as assigning blame to anyone. I very much admire and appreciate everyone I've met who is involved with scientific computing and Python. Using a stretched biological metaphor, it is only through the process of dividing and merging that better results are born. I think this is definitely the case with NumPy.

In early 2005, I decided to begin an effort to help bring the diverging community together under a common framework if it were possible. I first looked at numarray to see what could be done to add the missing features to make NumPy work with it as a core array object. After a couple of days of studying numarray, I was not enthusiastic about this approach. My familiarity with the Numeric code base no doubt biased my opinion, but it seemed to me that the features of Numarray could be added back to Numeric with a few fundamental changes to the core object. This would make the transition of SciPy to a more enhanced array object much easier in my mind.

Therefore, I began to construct this hybrid array object complete with an enhanced set of universal (broadcasting) functions that could deal with it. Along the way, quite a few new features and significant enhancements were added to the array

object and its surrounding infrastructure. This book describes the result of that year-long effort which culminated with the release of NumPy in early 2006. I first named the new package, SciPy Core, and used the scipy namespace. However, after a few months of testing under that name, it became clear that a separate namespace was needed for the new package. As a result, a rapid search for a new name resulted in actually coming back to the NumPy name which was the unofficial name of Numerical Python but never the actual namespace. Because the new package builds on the code-base of and is a successor to Numeric, I think the NumPy name is fitting and hopefully not too confusing to new users.

This book only briefly outlines some of the infrastructure that surrounds the basic objects in NumPy to provide the additional functionality contained in the older Numeric package (i.e. LinearAlgebra, RandomArray, FFT). This infrastructure in NumPy includes basic linear algebra routines, Fourier transform capabilities, and random number generators. In addition, the f2py module is described in its own documentation, and so is only briefly mentioned in the second part of the book. There are also extensions to the standard Python distutils and testing frameworks included with NumPy that are useful in constructing your own packages built on top of NumPy. The central purpose of this book, however, is to describe and document the basic NumPy system that is available under the numpy namespace.



#### NOTE

The numpy namespace includes all names under the numpy.core and numpy.lib namespaces as well. Thus, import numpy will also import the names from numpy.core and numpy.lib (along with fft, ifft, rand, and randn from from the other standard libraries). This is the recommended way to use numpy.

The following table gives a brief outline of the sub-packages contained in numpy package.

Sub-Package	Purpose	Comments
core	basic objects	all names exported to numpy
lib	additional utilities all names exported to	
linalg	basic linear algebra	old LinearAlgebra from Numeric
dft	discrete Fourier transforms	old FFT from Numeric
random	random number generators	old RandomArray from Numeric
distutils	enhanced build and distribution	improvements built on standard distutils
testing	unit-testing	utility functions useful for testing
f2py	automatic wrapping of Fortran code	a useful utility needed by SciPy

# Chapter 2

# Object Essentials

NumPy provides two fundamental objects: an N-dimensional array object (ndarray) and a universal function object (ufunc). There are other objects that build on top of these which you may find useful in your work, and these will be discussed later. The current chapter will provide background information on just the ndarray and the ufunc that will be important for understanding the attributes and methods to be discussed later.

An N-dimensional array is a homogeneous collection of "items" indexed using N integers. There are two essential pieces of information that define an N-dimensional array: 1) the shape of the array, and 2) the kind of item the array is composed of. The shape of the array is a tuple of N integers, one for each dimension, that provides information on how far the index can vary along that dimension. It is also necessary to specify the kind of item the array is composed of. Because every ndarray is a homogeneous collection of exactly the same data-type, every item takes up the same size block of memory, and each block of memory in the array is interpreted in exactly the same way<sup>1</sup>.



#### TIP

All arrays in base NumPy are indexed starting at 0 and ending at M-1 following the Python convention.

For example, consider the following piece of code:

<sup>&</sup>lt;sup>1</sup>By using OBJECT arrays, one can effectively have heterogeneous arrays, but the system still sees each element of the array as exactly the same thing (a reference to a Python object).

```
>>> a = array([[1,2,3],[4,5,6]])
>>> a.shape
(2, 3)
>>> a.dtype
dtype('<i4')
```



## NOTE

for all code in this book it is assumed that you have first entered from numpy import \*. In addition, any previously defined arrays are still defined for subsequent examples.

This code defines an array of size  $2 \times 3$  composed of 4-byte (little-endian) integer elements (on my 32-bit AMD platform). We can index into this two-dimensional array using two integers: the first integer running from 0 to 1 inclusive and the second from 0 to 2 inclusive. For example, index (1,1) selects the element with value 5:

```
>>> a[1,1]
5
```

All code shown in the shaded-boxes in this book has been (automatically) executed on a particular version of NumPy. The output of the code shown below shows which version of NumPy was used to create all of the output in your copy of this book.

```
>>> import numpy; print numpy._version_
0.9.9.2627
```

# 2.1 Data-Type Descriptors

In NumPy, an indexeray is an N-dimensional array of items where each item takes up a fixed number of bytes. Typically, this fixed number of bytes represents an integer or a floating-point number. However, this fixed number of bytes could also represent an arbitrary record made up of any collection of other data types. NumPy achieves this flexibility through the use of a data-type (dtype) object. Every array has an associated dtype object which describes the layout of the array data. Every dtype object, in turn, has an associated Python type-object (.type) that determines exactly what type of Python object is returned when an element of the array is

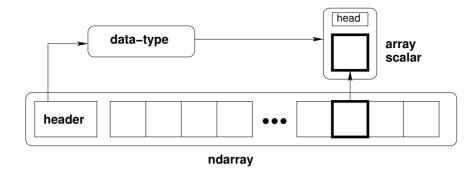


Figure 2.1: Conceptual diagram showing the relationship between the three fundamental objects used to describe the data in an array: 1) the ndarray itself, 2) the data-type object that describes the layout of a single fixed-size element of the array, 3) the array-scalar Python object that is returned when a single element of the array is accessed.

accessed. The dtype objects are flexible enough to contain references to arrays of other dtype objects and, therefore, can be used to define nested records. This advanced functionality will be described in better detail later as it is mainly useful for the recarray (record array) subclass that will also be defined later. However, all ndarrays can enjoy the flexibility provided by the dtype objects. Figure 2.1 provides a conceptual diagram showing the relationship between the ndarray, its associated data-type object, and an array-scalar that is returned when a single-element of the array is accessed. Note that the data-type points to the type-object of the array scalar. An array scalar is returned using the type-object and a particular element of the ndarray.

Every dtype object is based on one of 21 built-in dtype objects. These built-in objects allow numeric operations on a wide-variety of integer, floating-point, and complex data types. Associated with each data-type is a Python type object whose instances are array scalars. This type-object can be obtained using the **type** attribute of the dtype object. Python typically defines only one data-type of a particular data class (one integer type, one floating-point type, etc.). This can be convenient for some applications that don't need to be concerned with all the ways data can be represented in a computer. For scientific applications, however, this is not always true. As a result, in NumPy, their are 21 different fundamental Python data-type-descriptor objects built-in. These descriptors are mostly based on the types available in the C language that CPython is written in. However, there are a few types that are extremely flexible, such as string, unicode, and void.

The fundamental data-types are shown in Table 2.1. Along with their (mostly) C-derived names, the integer, float, and complex data-types are also available using

a bit-width convention so that an array of the right size can always be ensured (e.g. int8, float64, complex128). The C-like names are also accessible using a character code which is also shown in the table. Names for the data types that would clash with standard Python object names are followed by a trailing underscore, '-'. These data types are so named because they use the same underlying precision as the corresponding Python data types. Most scientific users should be able to use the array-enhanced scalar objects in place of the Python objects. The array-enhanced scalars inherit from the Python objects they can replace and should act like them under all circumstances (except for how errors are calculated in math computations).



#### TIP

The array types **bool\_**, **int\_**, **complex\_**, **float\_**, **object\_**, **unicode\_**, and **str\_** are enhanced-scalars. They are very similar to the standard Python types (without the trailing underscore) and inherit from them (except for bool\_ and object\_). They can be used in place of the standard Python types whenever desired. Whenever a data type is required, as an argument, the standard Python types are recognized as well.

Three of the data types are flexible in that they can have items that are of an arbitrary size: the str\_type, the unicode\_type, and the void type. While, you can specify an arbitrary size for the type, every item in the array is still of that specified size. The void type, for example, allows for arbitrary records to be defined as elements of the array, and can be used to define exotic types built on top of the basic ndarray.



## NOTE

The two types intp and uintp are not separate types. They are names bound to a specific integer type just large enough to hold a memory address (a pointer) on the platform.

Table 2.1: Built-in array-scalar types corresponding to data-types for an ndarray. The bold-face types correspond to standard Python types. The object\_type is special because arrays with dtype='O' do not return an array scalar on item access but instead return the actual object referenced in the array.

Type	Bit-Width	Character
bool_	boolXX	'?'
byte	intXX	'b'
short		'n'
intc		'i'
$int_{-}$		'1'
longlong		'q'
intp		'n,
ubyte	uintXX	'B'
ushort		'H'
uintc		'I'
uint		'L'
ulonglong		'Q'
uintp		'P'
single	floatXX	'f'
float_		'd'
longfloat		'g'
csingle	complexXX	'F'
complex_		'D'
clongfloat		'G'
object_		'0'
str_		'S#'
unicode_		'U#'
void		'V#'



## WARNING

Numeric Compatibility: If you used old typecode characters in your Numeric code (which was never recommended), you will need to change some of them to the new characters. In particular, the needed changes are 'c->'S1', 'b'->'B', '1'->'b', 's'->'h', 'w'->'H', and 'u'->'I'. These changes make the typecharacter convention more consistent with other Python modules such as the struct module.

The fundamental data-types are arranged into a hierarchy of Python typeobjects shown in Figure 2.2. Each of the leaves on this hierarchy correspond to actual data-types that arrays can have (in other words, there is a built in dtype object associated with each of these new types). They also correspond to new Python objects that can be created. These new objects are "scalar" types corresponding to each fundamental data-type. Their purpose is to smooth out the rough edges that result when mixing scalar and array operations. These scalar objects will be discussed in more detail in Chapter 6. The other types in the hierarchy define particular categories of types. These categories can be useful for testing whether or not dtype.type is of a particular class (using issubclass).

# 2.2 Basic indexing (slicing)

Indexing is a powerful tool in Python and NumPy takes full advantage of this power. In fact, some of capabilities of Python's indexing were first established by the needs of Numeric users<sup>2</sup>. Indexing is also called slicing in Python, and slicing for an **ndarray** works very similarly as it does for other Python sequences. There are three big differences: 1) slicing can be done over multiple dimensions, 2) exactly one ellipsis object can be used to indicate several dimensions at once, 3) slicing cannot be used to expand the size of an array (unlike lists).

A few examples should make slicing more clear. Suppose A is a  $10 \times 20$  array, then A[3] is the same as A[3,:] and represents the 4th length-20 "row" of the array. On the other hand, A[:,3] represents the 4th length-10 "column" of the array. Every third element of the 4th column can be selected as A[::3,3]. Ellipses can be used to replace zero or more ":" terms. In other words, an Ellipsis object expands to zero or more full slice objects (":") so that the total number of dimensions in the slicing

<sup>&</sup>lt;sup>2</sup>For example, the ability to index with a comma separated list of objects and have it correspond to indexing with a tuple is a feature added to Python at the request of the NumPy community. The Ellipsis object was also added to Python explicitly for the NumPy community. Extended slicing (wherein a step can be provided) was also a feature added to Python because of Numeric.

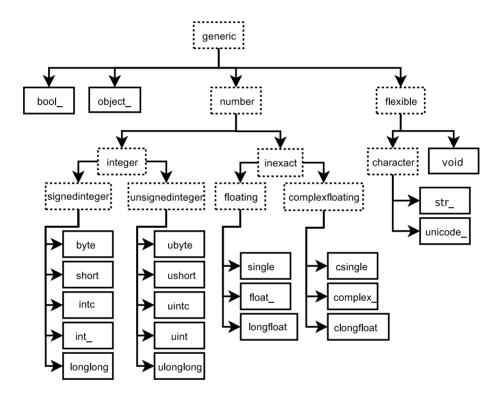


Figure 2.2: Hierarchy of type objects representing the array data types. Not shown are the two integer types intp and uintp which just point to the integer type that holds a pointer for the platform. All the number types can be obtained using bit-width names as well.

tuple matches the number of dimensions in the array. Thus, if A is  $10 \times 20 \times 30 \times 40$ , then A[3:,...,4] is equivalent to A[3:,:,:,4] while A[...,3] is equivalent to A[:,:,:,3]. The following code illustrates some of these concepts:

```
>>> print a[...,3]
[[ 3 8 13 18]
[23 28 33 38]
[43 48 53 58]]
>>> print a[1,...,3]
[23 28 33 38]
>>> print a[:,:,2]
[[ 2 7 12 17]
[22 27 32 37]
[42 47 52 57]]
>>> print a[0,::2,::2]
[[ 0 2 4]
[10 12 14]]
```

# 2.3 Memory Layout of ndarray

On a fundamental level, an N-dimensional array object is just a one-dimensional sequence of memory with fancy indexing code that maps an N-dimensional index into

a one-dimensional index. The one-dimensional index is necessary on some level because that is how memory is addressed in a computer. The fancy indexing, however, can be very helpful for translating our ideas into computer code. This is because many concepts we wish to model on a computer have a natural representation as an N-dimensional array. While this is especially true in science and engineering, it is also applicable to many other arenas which can be appreciated by considering the popularity of the spreadsheet as well as "image processing" applications.



## WARNING

Some high-level languages give pre-eminence to a particular use of 2-dimensional arrays as Matrices. In NumPy, however, the core object is the more general N-dimensional array. NumPy defines a matrix object as a sub-class of the N-dimensional array.

In order to more fully understand the array object along with its attributes and methods it is important to learn more about how an N-dimensional array is represented in the computer's memory. A complete understanding of this layout is only essential for optimizing algorithms operating on general purpose arrays. But, even for the casual user, an understanding of memory layout will help to explain the use of certain array attributes that may otherwise be mysterious.

## 2.3.1 Contiguous Memory Layout

There is a fundamental ambiguity in how the mapping to a one-dimensional index can take place which is illustrated for a 2-dimensional array in Figure 2.3. In that figure, each block represents a chunk of memory that is needed for representing the underlying array element. For example, each block could represent the 8 bytes needed to represent a double-precision floating point number. In the figure, two arrays are shown, a 4x3 array and a 3x4 array. Each of these arrays takes 12 blocks of memory shown as a single, contiguous segment. How this memory is used to form the abstract 2-dimensional array can vary, however, and the ndarray object supports both styles. Which style is in use can be interrogated by the use of the flags attribute which returns a dictionary of the state of array flags.

In the C-style of N-dimensional indexing shown on the left of Figure 2.3 the last N-dimensional index "varies the fastest." In other words, to move through computer memory sequentially, the last index is incremented first, followed by the second-to-last index and so forth. Some of the algorithms in NumPy that deal with N-dimensional arrays work best with this kind of data.

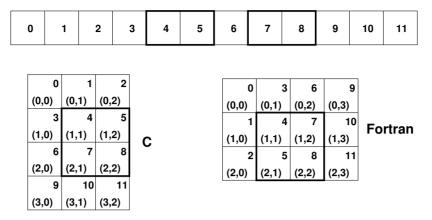


Figure 2.3: Options for memory layout of a 2-dimensional array.

In the Fortran-style of N-dimensional indexing shown on the right of Figure 2.3, the first N-dimensional index "varies the fastest." Thus, to move through computer memory sequentially, the first index is incremented first until it reaches the limit in that dimension, then the second index is incremented and the first index reset to zero. While NumPy can be compiled without the use of a Fortran compiler, several modules of the full installation of NumPy (available separately) rely on underlying algorithms written in Fortran. Algorithms that work on N-dimensional arrays that are written in Fortran typically expect Fortran-style arrays.

The two-styles of memory layout for arrays are connected through the transpose operation. Thus, if A is a (contiguous) C-style array, then the same block of memory can be used to represent  $A^T$  as a (contiguous) Fortran-style array. This kind of understanding can be useful when trying to optimize the wrapping of Fortran subroutines, or if a more detailed understanding of how to write algorithms for generally-indexed arrays is desired. But, fortunately, the casual user who does not care if an array is copied occasionally to get it into the right orientation needed for a particular algorithm can forget about how the array is stored in memory and just visualize it as an N-dimensional array (that is, after all, the whole point of creating the ndarray object in the first place).

# 2.3.2 Discontiguous memory layout

Both of the examples presented above are *single-segment* arrays where the entire array is visited by sequentially marching through memory one element at a time. When an algorithm in C or Fortran expects an N-dimensional array, this single segment (of a certain fundamental type) is usually what is expected along with

the shape N-tuple. With a single-segment of memory representing the array, the one-dimensional index into computer memory can always be computed from the N-dimensional index. This concept is explored further in the following paragraphs.

Let  $n_i$  be the value of the  $i^{\text{th}}$  index into an array whose shape is represented by the N integers  $d_i$  (i = 0...N-1). Then, the one-dimensional index into a C-style contiguous array is

$$n^C = \sum_{i=0}^{N-1} n_i \prod_{j=i+1}^{N-1} d_j$$

while the one-dimensional index into a Fortran-style contiguous array is

$$n^F = \sum_{i=0}^{N-1} n_i \prod_{j=0}^{i-1} d_j.$$

In these formulas we are assuming that

$$\prod_{j=k}^{m} d_j = d_k d_{k+1} \cdots d_{m-1} d_m$$

so that if m < k, the product is 1. While perfectly general, these formulas may be a bit confusing at first glimpse. Let's see how they expand out for determining the one-dimensional index corresponding to the element (1,3,2) of a  $4 \times 5 \times 6$  array. If the array is stored as Fortran contiguous, then

$$n^F = n_0 \cdot (1) + n_1 \cdot (4) + n_2 \cdot (4 \cdot 5)$$
  
= 1 + 3 \cdot 4 + 2 \cdot 20 = 53.

On the other hand, if the array is stored as C contiguous, then

$$n^C = n_0 \cdot (5 \cdot 6) + n_1 \cdot (6) + n_2 \cdot (1)$$
  
=  $1 \cdot 30 + 3 \cdot 6 + 2 \cdot 1 = 50$ .

The general pattern should be more clear from these examples.

The formulas for the one-dimensional index of the N-dimensional arrays reveal what results in an important generalization for memory layout. Notice that each formula can be written as

$$n^X = \sum_{i=0}^{N-1} n_i s_i^X$$

where  $s_i^X$  gives the **stride** for dimension  $i^3$ . Thus, for C and Fortran contiguous

<sup>&</sup>lt;sup>3</sup>Our definition of stride here is an element-based stride, while the strides attribute returns a

arrays respectively we have

$$s_i^C = \prod_{j=i+1}^{N-1} d_j = d_{i+1} d_{i+2} \cdots d_{N-1},$$
  
 $s_i^F = \prod_{j=0}^{i-1} d_j = d_0 d_1 \cdots d_{i-1}.$ 

The stride is how many elements in the underlying one-dimensional layout of the array one must jump in order to get to the next array element of a specific dimension in the N-dimensional layout. Thus, in a C-style  $4 \times 5 \times 6$  array one must jump over 30 elements to increment the first index by one, so 30 is the stride for the first dimension ( $s_0^C = 30$ ). If, for each array, we define a strides tuple with N integers, then we have pre-computed and stored an important piece of how to map the N-dimensional index to the one-dimensional one used by the computer.

In addition to providing a pre-computed table for index mapping, by allowing the strides tuple to consist of arbitrary integers we have provided a more general layout for the N-dimensional array. As long as we always use the strides information to move around in the N-dimensional array, we can use any convenient layout we wish for the underlying representation as long as it is regular enough to be defined by constant jumps in each dimension. The **ndarray** object of NumPy uses this strides information and therefore the underlying memory of an **ndarray** can be layed out dis-contiguously.



## NOTE

Several algorithms in NumPy work on arbitrarily strided arrays. However, some algorithms require single-segment arrays. When an irregularly strided array is passed in to such algorithms, a copy is automatically made.

An important situation where irregularly strided arrays occur is array indexing. Consider again Figure 2.3. In that figure a high-lighted sub-array is shown. Define C to be the  $4\times 3$  C contiguous array and F to be the  $3\times 4$  Fortran contiguous array. The highlighted areas can be written respectively as C[1:3,1:3] and F[1:3,1:3]. As evidenced by the corresponding highlighted region in the one-dimensional view of the memory, these sub-arrays are neither C contiguous nor Fortran contiguous. However, they can still be represented by an **ndarray** object using the same striding tuple as the original array used. Therefore, a regular indexing expression on an

byte-based stride. The byte-based stride is the element itemsize multiplied by the element-based stride.

ndarray can always produce an ndarray object without copying any data. This is sometimes referred to as the "view" feature of array indexing, and one can see that it is enabled by the use of striding information in the underlying ndarray object. The greatest benefit of this feature is that it allows indexing to be done very rapidly and without exploding memory usage (because no copies of the data are made).

# 2.4 Universal Functions for arrays

NumPy provides a wealth of mathematical functions that operate on then ndarray object. From algebraic functions such as addition and multiplication to trigonometric functions such as sin, and cos. Each universal function (ufunc) is an instance of a general class so that function behavior is the same. All ufuncs perform element-by-element operations over an array or a set of arrays (for multi-input functions). The ufuncs themselves and their methods are documented in Part 9.

One important aspect of ufunc behavior that should be introduced early, however, is the idea of **broadcasting**. Broadcasting is used in several places throughout NumPy and is therefore worth early exposure. To understand the idea of broadcasting, you first have to be conscious of the fact that all ufuncs are always element-by-element operations. In other words, suppose we have a ufunc with two inputs and one output (e.g. addition) and the inputs are both arrays of shape  $4 \times 6 \times 5$ . Then, the output is going to be  $4 \times 6 \times 5$ , and will be the result of applying the underlying function (e.g. +) to each pair of inputs to produce the output at the corresponding N-dimensional location.

Broadcasting allows ufuncs to deal in a meaningful way with inputs that do not have exactly the same shape. In particular, the first rule of broadcasting is that if all input arrays do not have the same number of dimensions, then a "1" will be repeatedly pre-pended to the shapes of the smaller arrays until all the arrays have the same number of dimensions. The second rule of broadcasting ensures that arrays with a size of 1 along a particular dimension act as if they had the size of the array with the largest shape along that dimension. The value of the array element is assumed to be the same along that dimension for the "broadcasted" array. After application of the broadcasting rules, the sizes of all arrays must match.

While a little tedious to explain, the broadcasting rules are easy to pick up by looking at a couple of examples. Suppose there is a **ufunc** with two inputs, A and B. Now supposed that A has shape  $4 \times 6 \times 5$  while B has shape  $4 \times 6 \times 1$ . The ufunc will proceed to compute the  $4 \times 6 \times 5$  output as if B had been  $4 \times 6 \times 5$  by assuming that B[...,k] = B[...,0] for k = 1,2,3,4.

Another example illustrates the idea of adding 1's to the beginning of the array

shape-tuple. Suppose A is the same as above, but B is a length 5 array. Because of the first rule, B will be interpreted as a  $1 \times 1 \times 5$  array, and then because of the second rule B will be interpreted as a  $4 \times 6 \times 5$  array by repeating the elements of B in the obvious way. If it is desired, instead, to add 1's to the end of the array shape, then dimensions can always be added using the newaxis name in NumPy.

One important aspect of broadcasting is the calculation of functions on regularly spaced grids. For example, suppose it is desired to show a portion of the multiplication table by computing the function a\*b on a grid with a running from 6 to 9 and b running from 12 to 16. The following code illustrates how this could be done using ufuncs and broadcasting.

```
>>> a = arange(6, 10); print a
[6 7 8 9]
>>> b = arange(12, 17); print b
[12 13 14 15 16]
>>> table = a[:,newaxis] * b
>>> print table
[[ 72 78 84 90 96]
[ 84 91 98 105 112]
[ 96 104 112 120 128]
[108 117 126 135 144]]
```

# 2.5 Summary of new features

More information about using arrays in Python can be found in the old Numeric documentation at http://numeric.scipy.org http://numeric.scipy.org. Quite a bit of that documentation is still accurate, especially in the discussion of array basics. There are significant differences, however, and this book seeks to explain them in detail. The following list tries to summarize the significant new features (over Numeric) available in the ndarray and ufunc objects of NumPy:

- 1. more data types (all standard C-data types plus complex floats, boolean, string, unicode, and void \*);
- 2. flexible data types where each array can have a different itemsize (but all elements of the same array still have the same itemsize);
- 3. there is a true Python type (contained in a hierarchy of types) for every datatype an array can have;

- 4. data-type objects define the data-type with support for data-type objects with fields and subarrays which allow record arrays with nested records;
- 5. many more array methods in addition to functional counterparts;
- 6. attributes more clearly distinguished from methods (attributes are intrinsic parts of an array so that setting them changes the array itself);
- 7. array scalars covering all data types which inherit from Python scalars when appropriate;
- 8. arrays can be misaligned, swapped, and in Fortran order in memory (facilitates memory-mapped arrays);
- 9. arrays can be more easily read from text files and created from buffers;
- 10. arrays can be quickly written to files in text and/or binary mode;
- 11. arrays inherit from big arrays which do not define the sequence, or buffer protocol and can therefore be very large on 64-bit platforms.
- fancy indexing can be done on arrays using integer sequences and boolean masks;
- 13. coercion rules are altered for mixed scalar / array operations so that scalars (anything that produces a 0-dimensional array internally) will not determine the output type in such cases.
- 14. when coercion is needed, temporary buffer-memory allocation is limited to a user-adjustable size;
- 15. errors are handled through the IEEE floating point status flags and there is flexibility on a per-thread level for handling these errors;
- 16. one can register an error callback function in Python to handle errors are set to 'call' for their error handling;
- 17. ufunc reduce, accumulate, and reduceat can take place using a different type then the array type if desired (without copying the entire array);
- 18. ufunc output arrays passed in can be a different type than expected from the calculation;
- 19. arbitrary classes can be passed through ufuncs (\_array\_wrap\_ and \_array\_priority\_);
- 20. ufuncs can be easily created from Python functions;

- 21. ufuncs have attributes to detail their behavior, including a dynamic doc string that automatically generates the calling signature;
- 22. several new ufuncs (frexp, modf, ldexp, isnan, isfinite, isinf, signbit);
- 23. new types can be registered with the system so that specialized ufunc loops can be written over new type objects;
- 24. C-API enhanced so that more of the functionality is available from compiled code;
- 25. C-API enhanced so array structure access can take place through macros;
- 26. new iterator objects created for easy handling in C of discontiguous arrays;
- 27. types have more functions associated with them (no magic function lists in the C-code). Any function needed is part of the type structure.

All of these enhancements will be documented more thoroughly in the remaining portions of this book.

# 2.6 Summary of differences with Numeric

An attempt was made to retain backwards compatibility with Numeric all the way to the C-level. This was mostly accomplished, with a few changes that needed to be made for consistency of the new system. If you are just starting out with NumPy, then this section may be skipped.



#### TIP

There is a module called convertcode.py that is distributed with NumPy. This script takes a Python filename <name>.py as an argument, saves a copy <name>.orig, and makes any needed changes to the script. This script only makes the necessary name replacement changes, and should handle many needs. The script is also availabel as a module numpy.convertcode.

Throughout this book, warnings are inserted when compatibility issues with old Numeric are raised. Here you can find a summary of all the differences that may need changing in your code to work with the new NumPy ndarray object. While you may not need to make any changes to get code to run with the ndarray object, you will likely want to make changes to take advantage of the new features of NumPy. Note that Numeric and NumPy can both be used together, however, so you can

use both simultaneously while you make the transition. In addition, if you have Numeric 24.0 or newer, they can even share the same memory. This makes it easy to use NumPy as well as third-party tools that have not made the switch yet.

## 2.6.1 The list of necessary changes:

#### 1. Importing

- (a) import Numeric -> import numpy as Numeric
- (b) import Numeric as XX -> import numpy as XX
- (c) from Numeric import <name1>,...<nameN> -> from numpy import <name1>,...,<nameN>
- (d) from Numeric import \* -> from numpy import \* (this may clobber more names and therefore require further fixes to your code but then you didn't do this regularly anyway did you). The recommended procedure if this replacement causes problems is to fix the use of from Numeric import \* to one of the previous three approaches and then continue.
- (e) Similar name changes need to be made for MLab (numpy.lib.mlab), LinearAlgebra (numpy.linalg), RandomArray (numpy.random), RNG (numpy.random), and FFT (numpy.dft).
- (f) multiarray and umath (if you used them directly) are now numpy.core.multiarray and numpy.core.umath.
- 2. The old names for functions that used to live under LinearAlgebra, Rando-mArray, and FFT are still there but they are not advertised in this book. The old interfaces for RNG are gone, for now. The functionality is available under numpy.random.
- 3. Method name changes and methods converted to attributes
  - (a) arr.typecode() -> arr.dtype.char
  - (b) arr.iscontiguous() -> arr.flags.contiguous
  - (c) arr.byteswapped() -> arr.byteswap()
  - (d)  $arr.toscalar() \rightarrow arr.item()$
  - (e) arr.itemsize() -> arr.itemsize
  - (f) arr.spacesaver() eliminated
  - (g) arr.savespace() eliminated

- 4. arr.flat now returns an indexable 1-D iterator. This behaves correctly when passed to a function, but if you expected methods or attributes on arr.flat besides .copy() then you will need to replace arr.flat with arr.ravel() (copies only when necessary) or arr.flatten() (always copies).
- 5. If you used the construct arr.shape=<tuple>, this will not work for array scalars. You cannot set the shape of an array-scalar (you can read it though). As a result, for more general code you should use arr=arr.reshape(<tuple>) which works for both array-scalars and arrays.
- 6. If your code should produce 0-d arrays (occasionally cropped up in Numeric). These no-longer have a length as they should be interpreted similarly to real scalars which don't have a length.
- 7. Some of the typecode characters have changed to be more consistent with other Python modules (array and struct). Numeric -> numpy
  - (a) 'b' -> 'B'
  - (b) '1' -> 'b'
  - (c) 's' -> 'h'
  - (d) 'w' -> 'H'
  - (e) 'u' -> 'I'
- 8. Keyword and argument changes
  - (a) All typecode= keywords have been changed to dtype=.
  - (b) The savespace keyword argument has been removed from all functions where it was present (array, sarray, asarray, ones, and zeros). The sarray function is equivalent to asarray.
- 9. Character arrays work a little differently now, but there is still such a thing as a 'c' array and it is nearly equivalent to an 'S1' array. The big difference is that on array creation it has the effect of ensuring that any string is interpreted as a list of characters as well as causing object assignment from short strings to pad with spaces as Numeric did. There is also a PyArray\_CHAR typecode equivalent to PyArray\_STRING with an itemsize of 1 (but easier to specify in C-code).
- 10. If you used type-equality testing on the objects returned from arrays, then you need to change this to isinstance testing. Thus type(a[0]) is float or type(a[0]) == float should be changed to isinstance(a[0], float).

11. Arrays cannot be tested for truth value unless they are empty (returns False) or have only one element. This means that if Z: where Z is an array will fail (unless Z is empty or has only one element). Also the 'and' and 'or' operations (which test for object truth value) will also fail on arrays of more than one element.

## 2.6.2 Updating code that uses Numeric using convertcode

It is not difficult to convert from Numeric to NumPy. For example all of SciPy was converted in about 2-3 days. The needed changes are largely search-and replace type changes, and there is a module (numpy.lib.convertcode) with functions that can make the process simpler. The two functions of interest are

#### convertfile (filename)

Convert the file with the given filename to use NumPy. A backup is first made and given the name filename.orig. Then, the file is converted and the updated code written over the top of the old file.

#### **convertall** (direc=os.path.curdir)

Converts all the ".py" files in the given directory to use NumPy. Backups of all the files are first made as explained for the convertille function.

# 2.6.3 Recommended changes

- 1. Convert typecharacters to bitwidth type names or c-type names.
- 2. Convert use of uppercase Int32, Float, etc., to lower case int32, float, etc.
- 3. Convert use of functions to method calls where appropriate (but notice the possibly different default arguments).
- 4. Look for ways to take advantage of advanced slicing.
- 5. Remove any kludges you inserted to eliminate problems with Numeric that are now gone.
- 6. Look for ways to take advantage of new features like expanded data-types.
- 7. See if you can inherit from the ndarray directly, rather than using UserArray (but if you were using UserArray in a multiple-inheritance hierarchy this is going to be more difficult).

8. Watch your usage of scalars extracted from arrays. Treating Numeric arrays like lists and then doing math on the elements 1 by 1 was always slower than using real lists. This is even more true in NumPy.

## Chapter 3

# The Array Object

### 3.1 ndarray Object Attributes

Array object attributes reflect information that is intrinsic to the array itself. Generally, accessing an array through its attributes allows you to get and sometimes set intrinsic properties of the array without creating a new array. The exposed attributes are the core parts of an array and only some of them can be reset meaningfully without creating a new array. Table 3.1 shows all the attributes with a brief description. Detailed information on each attribute is given below.



### WARNING

Numeric Compatibility: you should check your old use of the .flat attribute. This attribute now returns an iterator object which acts like a 1-d array in terms of indexing. while it does not share all the attributes or methods of an array, it will be interpreted as an array in functions that take objects and convert them to arrays. Furthermore, Any changes in an array converted from a 1-d iterator will be reflected back in the original array when the converted array is deleted.

### 3.1.1 Memory Layout attributes

### flags

Array flags provide information about how the memory area used for the array is to be interpreted. There are 6 boolean flags in use which govern whether or not:

Table 3.1: Attributes of the ndarray.

Attribute	Settable	Description
flags	No	Special array-connected dictionary-like object with attributes showing the state of flags in this array. Only the flags WRITEABLE, ALIGNED, and UPDATEIFCOPY can be modified by setting items in this special object.
shape	Yes	a tuple showing the array shape. An array can be reshaped by setting this attribute to a commensurate shape.
strides	Yes	a tuple showing how many bytes must be jumped in the data segment to get from one entry to the next
ndim	No	number of dimensions in array
data	Yes	a buffer object loosely wrapping the array data (only works for single-segment arrays).
size	No	the total number of elements
itemsize	No	the size (in bytes) of each element
nbytes	No	the total number of bytes used
base	No	the object this array is using for its data buffer, or None if it owns its own memory.
dtype	Yes	data-type object for this array
real	Yes	reference to real part of the array. Setting copies data to real part of current array.
imag	Yes	imaginary part, or read-only zero array if type is not complex. Setting works only if type is complex.
flat	Yes	return a one-dimensional, indexable iterator object that acts somewhat like a 1-d array.
_array_interface	No	A dictionary with keys (data, typestr, descr, shape, strides) for compliance with Python side of array protocol.
_array_struct_	No	array interface on C-level.
_array_priority	No	always 0.0 for base type ndarray.

**CONTIGUOUS (C)** the data layout is a C-style contiguous segment;

FORTRAN (F) the data layout is a Fortran-style contiguous segment;

**OWNDATA** (O) the array owns the memory it uses or if it borrows it from another object (if this is False, the base attribute retrieves a view of the object this array obtained its data from);

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

**ALIGNED (A)** the data and strides are aligned appropriately for the hardware;

**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.

Only the **UPDATEIFCOPY**, **WRITEABLE**, and **ALIGNED** flags can be changed by the user. This can be done using the special array-connected dictionary-like object that the flags attribute returns. By setting elements in this dictionary, the underlying array obect's flags are altered. Flags can also be changed using the method **setflags**(...). All flags in the dictionary can be accessed using their first (upper case) letter as well as the full name.

Certain logical combinations of flags can also be read using named keys to the special flags dictionary. These combinations are

FNC Returns FORTRAN and not CONTIGUOUS

FORC Returns FORTRAN or CONTIGUOUS (one-segment test).

BEHAVED (B) Returns ALIGNED and WRITEABLE

CARRAY (CA) Returns BEHAVED and CONTIGUOUS

**FARRAY\_(FA)** Returns BEHAVED and FORTRAN and not CONTIGUOUS



### NOTE

The array flags cannot be set arbitrarily. UPDATEIFCOPY can only be set False. the ALIGNED flag can only be set True if the data is truly aligned. The flag 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). The exception for string is made so that unpickling can be done without copying memory.

Flags can also be set and read using attribute access with the lower-case key equivalent (without first letter support). Thus, for example, self.flags.contiguous returns whether or not the array is contiguous, and self.flags.writeable=True changes the array to be writeable (if possible).

### shape

The shape of the array is a tuple giving the number of elements in each dimension. The shape can be reset for single-segment arrays by setting this attribute to another tuple. The total number of elements cannot change. However, a -1 may be used in a dimension entry to indicate that the array length in that dimension should be computed so that the total number of elements does not change. a.shape=x is equivalent to a=a.reshape(x) except the latter can be used even if the array is not single-segment and even if a is an array scalar.



### NOTE

Setting the shape attribute to () for a 1-element array will turn self into a 0-dimensional array. This is the one of Three ways to get a 0-dimensional array in Python. All other operations will return an array scalar (an instance of the array data type). The other ways to get a 0-dimensional array in Python are to call the \_array\_() method of an array scalar, or to call array() with a scalar argument. 0-dimensional arrays should normally not be needed directly.

### strides

The strides of an array is a tuple showing for each dimension how many bytes must be skipped to get to the next element in that dimension. Setting this attribute to another tuple will change the way the memory is viewed. This attribute can only be set to a tuple that will not cause the array to access unavailable memory. If an attempt is made to do so, ValueError is raised.

### ndim

The number of dimensions of an array is sometimes called the rank of the array. Getting this attribute reveals the length of the shape tuple and the strides tuple.

#### data

A buffer object referencing the actual data for this array if this array is single-segment. Otherwise, an AttributeError is raised. The buffer object is writeable depending on the status of self.flags.writeable.

#### size

The total number of elements in the array.

#### itemsize

The number of bytes each element of the array requires.

### nbytes

The total number of bytes used by the array. This is equal to self.itemsize\*self.size.

#### base

If the array does not own its own memory, then this attribute returns the object whose memory this array is referencing. The returned object may not be the original allocator of the memory, but may be borrowing it from still another object. If this array does own its own memory, then None is returned unless the UPDATEIFCOPY flag is True in which case self.base is the array that will be updated when self is deleted. (UPDATEIFCOPY gets set for an array that is created as a behaved copy of a general array. The intent is for the misaligned array to get any changes that occur to the copy).

### 3.1.2 Data Type attributes

There are several ways to specify the kind of data that the array is composed of. The fullest description that preserves field information is always obtained using an actual dtype object. See Chapter 7 for more discussion on data-type objects and acceptable arguments to construct data-types. Three commonly-used attributes of the data-type object returned are also documented here.

### dtype

A data-type object that fully describes (including any defined fields) each fixed-length item in the array. Whether or not the data is in machine byte-order is also determined by the data-type. This attribute can be set to anything that can be interpreted as a data-type (see Chapter 7 for more information). Setting this attribute allows you to change the interpretation of the data in the array. The new data-type must be compatible with the array's current data-type. The new data-type is compatible if it has the same itemsize as the current data-type descriptor, or (if the array is a single-segment array) if the the array with the new data-type fits in the memory already consumed by the array.

### dtype.type

A Python type object gives the typeobject whose instances represent elements of the array. This type object can be used to instantiate a scalar of that type.

### dtype.char

A typecode character unique to each of the 21 builtin types.

### dtype.str

This string consists of a required first character giving the "endianness" of the data ("<" for little endian, ">" for big endian, and "|" for irrelevant), the second character is a code for the kind of data ('b' for boolean, 'i' for signed integer, 'u' for unsigned integer, 'f' for floating-point, 'c' for complex floating point, 'O' for object, 'S' for ASCII string, 'U' for unicode, and 'V' for void), the final characters give the number of bytes each element uses.

### 3.1.3 Other attributes

### real

The real part of an array. For arrays that are not complex this attribute returns the array itself. Setting this attribute allows setting just the real part of an array. If the array is already real then setting this attribute is equivalent to self[...] = values.

#### imag

A view of the imaginary part of an array. For arrays that are not complex, this returns a read-only array of zeros. Setting this array allows in-place alteration of the complex part of an imaginary array. If the array is not complex, then trying to set this attribute raises an Error.

### flat

Return an iterator object (numpy.flatiter) that acts like a 1-d version of the array.

1-d indexing works on this array and it can be passed in to most routines as an array wherein a 1-d array will be constructed from it (using the \_\_array\_ protocol). The new 1-d array will reference this arrays data if this array is C-style contiguous, otherwise, new memory will be allocated for the 1-d array, the UPDATEIFCOPY flag will be set for the new array, and this array will have its WRITEABLE flag set FALSE until the the last reference to the new array disappears. When the last reference to the new 1-d array disappears,

the data will be copied over to this discontiguous array. This is done so that a.flat effectively references the current array regardless of whether or not it is contiguous or discontiguous. As an example, consider the following code:

The numpy.flatiter object has two methods: \_\_array\_\_(<dtype, None>) and copy() and one attribute: base. The base attribute returns a reference to the underlying array.

### \_array\_priority\_\_

The array priority attribute is a floating point number useful in mixed operations involving two subtypes to decide which subtype is returned. The base ndarray object has priority 0.0 and 1.0 is the default subtype priority.

### 3.1.4 Array Interface attributes

The array interface was created in 2005 as a means for array-like Python objects to re-use each other's data buffers intelligently whenever possible. The ndarray object supports the array interface. The system is able to consume objects that expose the array interface, and array objects can expose their inner workings to other objects that support the array interface. Four attributes of the array interface are exposed:

### \_array\_interface\_\_

The python-side of the array protocol. It is a dictionary with the following attributes:

data A 2-tuple (dataptr, read-only flag). The dataptr is a string giving the address (in hexadecimal format) of the array data. The read-only flag is True if the array memory is read-only.

**strides** The strides tuple. Same as **strides** attribute except None is returned if the array is C-style contiguous.

**shape** The shape tuple. Same as **shape** attribute.

typestr A string giving the format of the data. Same as dtype.str attribute.

**descr** A list of tuples providing the detailed description of this data type. This information is obtained from the arrdescr attribute of the dtypedescr object associated with each array. For arrays with fields, this will return a valid array-protocol descriptor list. For arrays without defined fields, this returns [(",typestr)].

### \_array\_struct\_

A PyCObject that wraps a pointer to a PyArrayInterface structure. This is only useful on the C-level for rapid implementation of the array interface, using a single attribute lookup.

### 3.2 ndarray Methods

In NumPy, the ndarray object has many methods which operate on or with the array in some fashion, typically returning a result. In Numeric, many of these methods were library calls whereas now they are also methods. These methods are explained in this chapter. Whenever the array itself needs to be referenced it will be referred to as "this array," or "self." Keyword arguments will be shown. Methods that only take one argument do not have keyword arguments. Default values for one argument methods will be shown in brackets "<default>".



### WARNING

If you are converting code from Numeric, then you will need to make the following (search and replace) conversions: .typecode() --> .dtype.char; .iscontiguous() --> .flags.contiguous; .byteswapped() --> .byteswap(); .toscalar() -> .item(); and .itemsize() --> .itemsize. The convertcode module can automate this for you.

### 3.2.1 Array conversion

### tolist ()

The contents of self as a nested list.

```
>>> a = array([[1,2,3],[4,5,6]]); a.tolist()
```

### item ()

This method only works for arrays with one element (a.size == 1). In this case, it returns a standard Python scalar object (if possible) copied from the first element of self. When the data type of self is longdouble or clongdouble, this 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.

```
>>> asc = a[0,0].item()
>>> type(asc)
<type 'int'>
>>> asc
1
>>> type(a[0,0])
<type 'int32scalar'>
```

### tostring ()

A Python string showing the raw contents of data memory. For single-segment arrays, this is equivalent to self.data[:], for other arrays this is equivalent to self.copy().data[:].

```
tofile (file=, sep=", format=")
```

Write the contents of self to the open file object, file. If file is a string, then open a file of that name first. If sep is the empty string, then write the file in binary mode. If sep is any other string, write the array in simple text mode separating each element with the value of the sep string. When the file is written in text mode, the format string can be used to alter the appearance of each entry. If format is the empty string, then it is equivalent to ''%s''. Each element of the array will be converted to a Python scalar, o, and written to the file as ''format'', o. Note that writing an array to a file does not store any information about the shape, type, or endianness of an array. When written in binary mode, to file is functionally equivalent to fid.write(self.tostring()).

```
>>> a.tofile('myfile.txt',sep=':',format='%03d')
Contents of myfile.txt
001:002:003:004:005:006
```

### dump (file)

Pickle the contents of self to the file object represented by file. Equivalent to cPickle.dump(self, file, 2)

### dumps ()

Return pickled representation of self as a string. Equivalent to cPickle.dumps(self, 2)

### astype (<None>)

Force conversion of this array to an array with the data type provided as the argument. If the argument is None, or equal to the data type of self, then return a copy of the array.

### byteswap (<False>)

Byteswap the elements of the array and return the byteswapped array. If the argument is True, then byteswap in-place and return a reference to self. Otherwise, return a copy of the array with the elements byteswapped. The data-type descriptor is not changed so the array will actually act different.

### copy ()

Return a copy of the array (which is always single-segment, and ALIGNED). However, the data-type is preserved (including whether or not the data is byteswapped).

### view (<None>)

Return a new array using the same memory area as self. If the optional argument is given, it can be either a typeobject that is a sub-type of the ndarray or an object that can be converted to a data-type descriptor. If the argument is a typeobject then a new array of that Python type is returned that uses the information from self. If the argument is a data-type descriptor, then a new array of the same Python type as a is returned using the given data-type.

```
_array__ (<None>)
```

Returns a reference to self. Present largely for consistency and for subclasses. The \_array\_ method of an object should return an ndarray. If the optional data-type argument is given, an array converted to that data-type will be returned.

```
getfield (dtype=, offset=0)
```

Return a *field* of the given array as an array of the given data type. A field is a view of the array's data at a certain byte offset interpreted as a given data type. The returned array is a reference into self, therefore changes made to the returned array will be reflected in self. This method is particularly useful for record arrays that use a void data type, but it can also be used to extract the low (high)-order bytes of other array types as well. For example, using getfield, you could extract fixed-length substrings from an array of strings.

```
>>> a = array(['Hello','World','NumPy'])
>>> a.getfield('S2',1)
array([el, or, um],
dtype='|S2')
```

```
setflags (write=None, align=None, uic=None)
```

Set array flags WRITEABLE, ALIGNED, and UPDATEIFCOPY, respectively. 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 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.

```
fill (scalar)
```

Method	Arguments	Description
astype	(dtype < None >)	Cast to another data type
byteswap	(inplace <false>)</false>	Byteswap array elements
copy	()	Copy array
dump	(file)	Pickle to stream or file
dumps	()	Get pickled string
fill	(scalar)	Fill an array with scalar value
getfield	(dtype=, offset=0)	Return a field of the array
setflags	(write=None,	Set array flags
	align=None, uic=None)	
tofile	(file=, sep=",	Raw write to file
	format=")	
tolist	()	Array as a nested list
item	()	Python scalar from first element
tostring	()	String of raw memory
view	(dtype < None >)	View as another data type

Table 3.2: Array conversion methods

Fill an array with the scalar value (appropriately converted to the type of self). If the scalar value is an array or a sequence, then only the first element is used. This method is usually faster than a[...]=scalar or self.flat=scalar, and always interprets its argument as a scalar.

### 3.2.2 Array shape manipulation

For reshape, resize, and transpose, the single tuple argument may be replaced with n integers which will be interpreted as an n-tuple.

### reshape (newshape)

Return an array that uses the same data as this array but has a new shape given by the newshape tuple (or a scalar to reshape as 1-d). The new shape must define an array with the same total number of elements. If one of the elements of the new shape tuple is -1, then that dimension will be determined such that the overall number of items in the array stays constant. If self is a single-segment array then the new array will reference the data of the old one. If self is not single-segment, then the new array will contain a copy of the data in self.

resize (newshape, refcheck=1)

Resize an array in-place. This changes self (in-place) to be an array with the new shape, reallocating space for the data area if necessary. If the data memory must be changed because the number of new elements is different than self.size, then an error will occur if this array does not own its data or if another object is referencing this one. Only a single-segment array can be resized. The method returns None. To bypass the reference count check, then set refcheck=0. The purpose of the reference count check is to make sure you don't 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 to another Python object, then you may safely set refcheck=0.

### transpose (<None>)

Return an array view with the shape transposed according to the argument. An argument of None is equivalent to range(self.ndim)[::-1]. The argument can either be a tuple or multiple integer arguments. This method returns a new array with permuted shape and strides attributes using the same data as self.

```
>>> a = arange(40).reshape((2,4,5))
>>> b = a.transpose(2,0,1)
>>> print a.shape, b.shape
(2, 4, 5) (5, 2, 4)
>>> print a.strides, b.strides
(80, 20, 4) (4, 80, 20)
>>> print a
[[[0 1 2 3 4]
[5 6 7 8 9]
[10 11 12 13 14]
[15 16 17 18 19]]
[[20 21 22 23 24]
[25 26 27 28 29]
[30 31 32 33 34]
[35 36 37 38 39]]]
>>> print b
[[[ 0 5 10 15]
[20 25 30 35]]
[[ 1 6 11 16]
[21 26 31 36]]
[[ 2 7 12 17]
[22 27 32 37]]
[[ 3 8 13 18]
[23 28 33 38]]
[[ 4 9 14 19]
[24 29 34 39]]]
```

### swapaxes (axis1, axis2)

Return an array view with axis1 and axis2 swapped. This is a special case of the transpose method with argument equal to arg=range(self.ndim); arg[axis1], arg[axis2] = arg[axis2], arg[axis1].

### flatten ()

Return a new 1-d array with data copied from self. Equivalent to but slightly faster then a.flat.copy().

### ravel ()

Return a 1-d version of self. If self is single-segment, then the new array references self, otherwise, a copy is made.

### squeeze ()

Return an array with all unit-length dimensions squeezed out.

### 3.2.3 Array item selection and manipulation

For array methods that take an axis keyword, it defaults to None. If axis is None, then the array is treated as a 1-D array. Any other value for axis represents the dimension along which the operation should proceed.

```
take (indices=, axis=None)
```

The functionality of this method is available using the advanced indexing ability of the ndarray object. However, for doing selection along a single axis it is usually faster to use take. If axis is not None, this method is equivalent to self[indxobj] preceded by indxobj=[slice(None)]\*self.ndim; indxobj[axis] = indices. It returns the elements or sub-arrays from self indicated by the index numbers in indices. If axis is None, then this method is equivalent to self.flat[indices].

```
put (values=, indices=)

Performs the equivalent of

for n in indices:
    self.flat[n] = values[n]

Values is repeated if it is too short.

putmask (values=, mask=)

Performs the equivalent of

for n, obj in enumerate(mask.flat):
    if obj:
        self.flat[n] = values[n]
```

The values array is repeated if it is too short.

```
repeat (repeats=, axis=None)
```

Copy elements (or sub-arrays selected along axis) of self **repeats** times. The repeats argument must be a sequence of length self.shape[axis] or a scalar. The repeats argument dictates how many times the element (or sub-array) will be repeated in the result array.

```
choose (choices)
```

The array must be an integer (or bool) array with entries from 0 to n. Choices is a tuple of n choice arrays:  $b0, b1, \ldots, bn$ . (Alternatively, choices can be replaced with n arguments where each argument is a choice array). The return array will be formed from the elements of the choice arrays occording to the value of the elements of self.

```
>>> a = array([0,3,2,1])
>>> a.choose([0,1,2,3],[10,11,12,13],
... [20,21,22,23],[30,31,32,33])
array([0,31,22,13])
```

```
sort (axis=-1, kind='quick')
```

Sort the array in-place and return None. The sort takes place over the given axis using an underlying sort algorithm specified by kind. The sorting algorithms available are 'quick', 'heap', and 'merge'. For flexible types only the quicksort algorithm is available.

```
>>> a=array([[0.2,1.3,2.5],[1.5,0.1,1.4]]);

>>> b=a.copy(); b.sort(0); print b

[[ 0.2  0.1  1.4]

[ 1.5  1.3  2.5]]

>>> b=a.copy(); b.sort(1); print b

[[ 0.2  1.3  2.5]

[ 0.1  1.4  1.5]]
```

```
argsort (axis=-1, kind='quick')
```

Return an index array of the same size as self showing which indices along the given axis should be selected to sort self along that axis. Uses an underlying sort algorithm specified by kind. The sorting algorithms available are 'quick', 'heap', and 'merge'.

```
>>> b=a.copy(); print b.argsort(0)
[[0 1 1]
[1 0 0]]
>>> b=a.copy(); print b.argsort(1)
[[0 1 2]
[1 2 0]]
```



### TIP

Complex valued arrays sort by comparing first the real parts and then the imaginary parts if the real parts are the same.

### searchsorted (values)

Return an index array (dtype=intp) of the same shape as values showing the index where the value would fit in self. The index is such that self[index-1] < value <= self[index]. In this formula self[self.size]= $\infty$  and self[-1]= $-\infty$ . Therefore, if value is larger than all elements of self, then index is self.size. If value is smaller than all elements of self, then index is 0. Self must be a sorted 1-d array. If elements of self are repeated, the index of the first occurrence is used.

```
>>> b=a.ravel(); b.sort()
>>> b.searchsorted([0.0, 1.35, 2.0, 3.0])
array([0, 3, 5, 6])
```

#### nonzero ()

Return the indices for elements of self that are nonzero. If self.ndim > 1, then return a tuple of index arrays of equal length giving the n-dimensional index to the non-zero elements of self.

```
>>> x = arange(15); y=x.reshape(3,5)
>>> (x>8).nonzero()
(array([ 9, 10, 11, 12, 13, 14]),)
>>> (y>8).nonzero()
(array([1, 2, 2, 2, 2]), array([4, 0, 1, 2, 3, 4]))
```

### compress (condition=, axis=None)

This method expects condition to be a one-dimensional mask array of the same length as self.shape[axis]. If the array is less than self.shape[axis], then False is assumed for the missing elements. The method returns the elements (or sub-arrays along the given axis) of self where condition is true. The shape of the return array is self.shape with the axis dimension replaced by the number of True elements of condition. The same effect can often be accomplished using array indexing.

```
>>> x=array([0,1,2,3])
>>> x.compress(x > 2)
array([3])
>>> x[x>2]
array([3])
```

### diagonal (offset=0, axis1=0, axis2=1)

If self is 2-d, return the **offset** (from the main diagonal) diagonal of self. If self is larger than 2-d, then return an array constructed from all the diagonals created from all the 2-d sub-arrays formed using all of axis1 and axis2. The offset parameter is with respect to axis2. The shape of the returned array is found by removing the axis1 and axis2 entries from self.shape and then appending the length of the offset diagonal of each 2-d sub-array.

### 3.2.4 Array calculation

Many of these methods take an argument named axis. In such cases, if axis is None (the default), the array is treated as a 1-d array and the operation is performed over the entire array. This behavior is also the default if self is a 0-dimensional array or array scalar. If axis is an integer, then the operation is done over the given axis (for each 1-d subarray that can be created along the given axis). The parameter dtype specifies the data type over which a reduction operation (like summing) should take place. The default reduce data type is the same as the data type of self. To avoid overflow, it is sometimes useful to perform the reduction using a larger data type.

```
max (axis <None>)
```

Return the largest value in self. This is a better way to compute the maximum over an array, than using max(self). The latter uses the generic sequence interface to self. This will be slower, and will try to get an answer by comparing whole sub-arrays of self. This will be incorrect for arrays larger than 1-d.

```
argmax (axis <None>)
Return the (first, 1-d) index of the largest value in self.
```

```
min (axis <None>)
```

Return the smallest value in self. This is a better way to compute the minimum over an array, than using min(self). The latter uses the generic sequence interface to self. This will be slower, and will try to get an answer by comparing whole sub-arrays of self. This will be incorrect for arrays larger than 1-d.

Table 3.3: Array item selection and shape manipulation methods. If axis is an argument, then the calculation is performed along that axis. An axis value of None means the array is flattened before calculation proceeds.

Method	Arguments	Description
argsort	argsort (axis <none>)</none>	Indices showing how to sort array.
choose	choose (c0,c1,,cn)	Choose from different arrays based on value of self.
compress	(condition=, axis=None)	Elements of self where condition is true.
diagonal	(offset=0, axis1=0, axis2=1)	Return a diagonal from self.
flatten	()	A 1-d copy of self.
nonzero	()	True where self is not zero.
put	(indices=, values=)	Place values at 1-d index locations of self.
putmask	(mask=, values=)	Place values in 1-d index locations where mask is True.
ravel	()	1-d version of self (no data copy if self is C-style contiguous).
repeat	(repeats=, axis=None)	Repeat elements of self.
reshape	(d1,d2,,dn)	Return reshaped version of self.
resize	(d1,d2,,dn,refcheck=1)	Resize self in-place.
searchsorted	(values)	Show where values would be placed in self (assumed sorted).
sort	(axis <none>)</none>	Copy of self sorted along axis.
squeeze	()	Squeeze out all length-1 dimensions.
swapaxes	(axis1, axis2)	Swap two dimensions of self.
take	(indices=, axis=None)	Select elements of self along axis according to indices.
transpose	(permute <none>)</none>	Rearrange shape of self according to permute.

argmin (axis <None>)

Return the (first, 1-d) index of the smallest value in self.

ptp (axis <None>)

Return the difference of the largest to the smallest value in self. Equivalent to self.max(axis) - self.min(axis)

clip (min=,max=)

Return a new array where any element in self less than min is set to min and any element less than max is set to max. Equivalent to self[self<min]=min; self[self>max]=max.

conj ()

conjugate ()

Return the conjugate of elements of the array.

round (decimals=0)

Round the elements of the array to the nearest decimal. For decimals < 0, the rounding is done to the nearest tens, hundreds, etc. Rounding of exactly the half-interval is to the nearest even integer. This is the only difference with standard Python rounding.

trace (offset=0, axis1=0, axis2=1, dtype=None)

Perform a summation along each diagonal specified by offset, axis1, and axis2. Equivalent to diagonal (offset, axis1, axis2).sum(axis=-1, dtype=dtype)

sum (axis=None, dtype=None)

Return the sum

$$\sum_{i=0}^{N-1} \operatorname{self}[\underbrace{:,\ldots,:}_{\operatorname{axis}}, i]$$

where axis ':' objects are placed before the i.

**cumsum** (axis=None, dtype=None)

Return the cumulative sum. If ret is the return array of the same shape as self, then

$$\operatorname{ret}[\underbrace{:,\ldots,:}_{\operatorname{axis}},j] = \sum_{i=0}^{j} \operatorname{self}[\underbrace{:,\ldots,:}_{\operatorname{axis}},i].$$

mean (axis=None, dtype=None)

Return the average value caculated as

$$\frac{1}{N} \sum_{i=0}^{N-1} \operatorname{self}[\underbrace{:,\dots,:}_{\text{axis}}, i]$$

where N is self.shape[axis] and axis ':' objects are placed before the i.

var (axis=None, dtype=None)

Return the variance of the data calculated as

$$\frac{1}{N} \sum_{i=0}^{N-1} \left( \text{self}[\underbrace{:,\dots,:}_{\text{axis}}, i] - \mu \right)^{2}$$

where N is self.shape[axis] and  $\mu$  is the mean (restored to the same number of dimensions as self with  $\mu$  copied along the axis dimension). This is equivalent to (self\*\*2).mean - self.mean()\*\*2 and ((self-self.mean())\*\*2).mean(). The value of N-1 was not chosen for normalization because while it gives an "unbiased" estimate, it is not prudent to return unbiased estimates as they may have larger mean-square error.

std (axis=None, dtype=None)

Return the standard deviation calculated as

$$\sqrt{\frac{1}{N}\sum_{i=0}^{N-1} \left( \text{self}[\underbrace{:,\dots,:}_{\text{axis}}, i] - \mu \right)^{2}}$$

where N is self.shape[axis] and  $\mu$  is the mean (restored to the same number of dimensions as self with  $\mu$  copied along the axis dimension).

**prod** (axis=None, dtype=None)

Return the product calculated as

$$\prod_{i=0}^{N-1} \operatorname{self}[\underbrace{:,\ldots,:}_{\operatorname{axis}}, i].$$

**cumprod** (axis=None, dtype=None)

Return the cumulative product so that the return array, ret, is the same shape as self and

$$\operatorname{ret}[\underbrace{:,\ldots,:}_{\operatorname{axis}},j] = \prod_{i=0}^{j} \operatorname{self}[\underbrace{:,\ldots,:}_{\operatorname{axis}},i].$$

all (axis <None>)

Return True if all entries along axis evaluate True, otherwise return False.

Return True if any entries along axis evaluate True, otherwise return False.

### 3.3 Array Special Methods

Methods in this chapter are not generally meant to be called directly by the user. They are called by Python and are used to customize behavior of the ndarray object as it interacts with the Python language and standard library.

### 3.3.1 Methods for standard library functions

To allow copy.copy(a) to perform a shallow copy of an array. Exactly the same as self.copy() (contents of object arrays are not copied).

```
_deepcopy_ (memodict)
```

To allow copy.deepcopy(a) to perform a deep copy. This is the same as a shallow copy unless self is an object array. Then, after the shallow copy is made, a copy.deepcopy(item) is called for every item in the object array.

```
_reduce__ ()
_setstate__ (shape, typestr, isfortran, data)
```

Pickling support for arrays is provided by these two methods. When an array needs to be pickled, the \_reduce\_() method is called to provide a 3-tuple of already-pickleable objects. To construct a new object from the pickle, the first two elements of the 3-tuple are used to construct a new (0-length) array of the correct type and the last element of the 3-tuple, which is itself a 4-tuple of (shape, typestr, isfortran, data) is passed to the \_setstate\_ method of the newly created array to restore its contents.

Table 3.4: Array object calculation methods. If axis is an argument, then the calculation is performed along that axis. An axis value of None means the array is flattened before calculation proceeds.

Method	Arguments	Description
all	(axis <none>)</none>	true if all entries are true.
any	(axis < None >)	true if any entries are true.
argmax	(axis < None >)	index of largest value.
argmin	(axis < None >)	index of smallest value.
clip	(min=, max=)	$self[self{>}max]{=}max; self[self{<}min]{=}min$
conj	()	complex conjugate
cumprod	(axis=None, dtype=None)	cumulative product
cumsum	(axis=None, dtype=None)	cumulative sum
max	(axis < None >)	maximum of self
mean	(axis=None, dtype=None)	mean of self
min	(axis < None >)	minimum of self
prod	(axis=None, dtype=None)	multiply elements of self together
ptp	(axis <none>)</none>	self.max(axis)-self.min(axis)
var	(axis=None, dtype=None)	variance of self
$\operatorname{std}$	(axis=None, dtype=None)	standard deviation of self
sum	(axis=None, dtype=None)	add elements of self together
trace	(offset, axis1=0, axis2=0, dtype=None)	sum along a diagonal

The reduce method returns a 3-tuple consisting of (callable, args, state) where callable is a simple constructor function that handles subclasses of the ndarray. Also, args is a 3-tuple of arguments to pass to this constructor function (type(self), (0,), self.dtypechar), and state is a 4-tuple of information giving the object's state (self.shape, self.dtypedescr, isfortran, string\_or\_list). In this tuple, isfortran is a Bool stating whether the following flattened data is in Fortran order or not, and string\_or\_list is a string formed by self.tostring() if the data type is not object. If the data type of self is an object array, then string\_or\_list is a flat list equivalent to self.ravel().tolist().

On load from a pickle, the pickling code uses the first two elements from the tuple returned by reduce to construct an empty 0-dimensional subclass of the correct type. The last element is then passed to the \_setstate\_ method of the newly created array to restore its contents.



### NOTE

When data is a string, the \_setstate\_ method will directly use the string memory as the array memory (new.base will point to the string). The typestr contains enough information to decode how the memory should be interpreted.

### 3.3.2 Basic customization

\_new\_\_ (subtype, shape=, dtype=long\_, buffer=None, offset=0, strides=None, order=None)

This method creates a new ndarray. It is typically only used in the \_\_new\_\_ method of a subclass. This method is called to construct a new array whenever the object name is called, a=ndarray(...). It supports two basic modes of array creation:

- 1. a single-segment array of the specified shape and data-type from newly allocated memory;
  - (a) uses shape, dtype, strides, and order arguments; others are ignored;
  - (b) The order argument allows specification of a Fortran-style contiguous memory segment (order='Fortran');
  - (c) If strides is given, then it specifies the new strides of the array (and the order keyword is ignored). The strides will be checked for consistency with the dimension size so that steps outside of the memory won't occur.

- 2. an array of the given shape and data type using the provided object, buffer, which must export the buffer interface.
  - (a) all arguments can be used:
  - (b) strides can be given and will be checked for consistency with the shape, data type, and available memory in buffer;
  - (c) order indicates whether the data buffer should be interpreted as Fortranstyle contiguous (order='Fortran') or not;
  - (d) offset can be used to start the array data at some offset in the buffer.



### NOTE

The ndarray uses the default no-op \_\_init\_\_ function because the array is completely initialized after \_\_new\_\_ is called.

```
_array__ (dtype <None>)
```

This is a special method that should always return an object of type ndarray. Useful for subclasses that need to get to the ndarray object.

```
_array_wrap__ (arr)
```

This is a special method that always returns an object of the same Python type as self using the array passed as an argument. This is mainly useful for subclasses as it is an easy way to get the subclass back from an ndarray.

```
_lt_ (other)
```

**\_le\_** (other)

**\_gt**\_\_ (other)

**\_ge**\_ (other)

**\_eq**\_ (other)

**\_ne**\_ (other)

Defined to support rich comparisons (<, <=, >=, ==, !=) on ndarrays using universal functions.

\_str\_\_ ()

\_repr\_ ()

- These functions print the array when called by str(self) and repr(self) respectively.

  Array printing can be changed using set\_string\_function(..). Default array printing has been borrowed from numerray whose printing code was written by Perry Greenfield and J. Todd Miller. By default, arrays print such that
- 1. The last axis is always printed left to right.
- 2. The next-to-last axis is printed top to bottom.
- 3. Remaining axes are printed top to bottom with increasing numbers of separators.

Five parameters of the printing can be set using keyword arguments with set\_printoptions(...)

The parameters can all be retrieved using get\_printoptions(). These printing options are

**precision** the number of digits of precision for floating point output (default 8);

threshold total number of array elements which triggers summarization rather than full representation (default 1000);

**edgeitems** number of array items in summary at beginning an end of each dimension (default 3);

**linewidth** the number of characters per line for the purpose of inserting line breaks (default 71);

suppress boolean indicating whether or not to supress printing of small floating point values using scientific notation (default False).

### \_nonzero\_\_ ()

Truth-value testing for the array as a whole. It is called whenever the truth value of the ndarray as a whole object is required. This raises an error if the number of elements in the the array is larger than 1 because the truth value of such arrays is ambiguous. Use .any() and .all() instead to be clear about what is meant in such cases. If the number of elements is 0 then False is returned. If there is one element in the array, then the truth-value of this element is returned.

### 3.3.3 Container customization

\_len\_\_ ()

Returns self.shape[0]. It is called in response to len(self). Use self.size to get the total number of elements in the array.

Notice that the default Python iterator for sequences is used when arrays are used in places that expect an iterator. This iterator returns successively self[0], self[1], ..., self[self.\_\_len\_\_()]. Use self.flat to get an iterator that walks through the entire array one element at a time.

```
_getitem_ (key)
```

Called when evaluating self[key] construct. Items from the array can be selected using this customization. This construct has both standard and extended indexing abilities which are explained in Section 3.3.4.4. A named field can be retrieved if key is a string and fields are defined in the dtypedescr object associated with this array.

```
_setitem_ (key, value)
```

Called when evaluating self[key]=value. Items in the array can be set using this construct. This construct is explained in Section 3.3.4.4. A named field can be set if key is a string and fields are defined in the dtypedescr object associated with this array.

```
\_getslice\_ (i, j)
```

Equivalent to self.\_getitem\_(slice(i,j)) but defined mainly so that C code can use the sequence interface. Called to evaluate self[i:j]

```
__setslice__ (i, j, value)
```

Equivalent to self.\_setitem\_(slice(i,j), value) but defined mainly so C code can use the sequence interface. Called to evaluate self[i:j] = value.

```
_contains_ (item)
```

Called to determine truth value of the item in self construct. Returns the equivalent of (self==item).any()

### 3.3.4 Arithmetic customization

### 3.3.4.1 Binary

```
_add__ (other)
```

\_sub\_ (other)

```
_mul_ (self, other)
_div_ (other)
_truediv_ (other)
_floordiv_ (other)
_mod_ (other)
_divmod_ (other)
_pow_ (other[,modulo])
_lshift_ (other)
_rshift_ (other)
_and_ (other)
_or_ (other)
_xor_ (other)
```

These methods are defined for ndarrays to implement the operations (+, -, \*, /, /, //, %, divmod(), \*\* or pow(), <<, >>, &, ^, |). This is done using calls to the corresponding universal function object (add, subtract, multiply, divide, true\_divide, floor\_divide, remainder, divide and remainder, power, left\_shift, right\_shift, bitwise\_and, bitwise\_xor, bitwise\_or). These implement element-by-element operations for arrays that are broadcastable to the same shape.

- any third argument to pow() is silently ignored as the underlying ufunc (power) only takes two arguments.
- the three division operators are all defined, div is active by default, truediv is active when \_future\_\_division is in effect.



### NOTE

Because it is a builtin type (written in C), the \_r<op>\_ special methods are not directly defined for the ndarray.

In-place

```
__iadd__ (other)
__isub__ (other)
__imul__ (other)
__idiv__ (other)
__itruediv__ (other)
```

3.3.4.2

**\_ifloordiv**\_ (other)

**\_imod\_** (other)

**\_ipow\_** (other)

**\_ilshift**\_ (other)

**\_irshift**\_ (other)

**\_\_iand\_\_** (other)

**\_ixor\_** (other)

**\_\_ior\_\_** (other)

These methods are implemented to handle the inplace operations (+=, -=, \*=, /=, /=, //=, %=, \*\*=, <<=, >>=, &=, ^=, |=). The inplace operators are implemented using the corresponding ufunc and its ability to take an output argument (which is set as self). Using inplace operations can save space and time and is therefore encouraged whenever appropriate.



### WARNING

In place operations will perform the calculation using the precision decided by the data type of the two operands, but will silently downcast the result (if necessary) so it can fit back into the array. Therefore, for mixed precision calculations, a  $\langle op \rangle = B$  can be different than  $a = a \langle op \rangle B$ . For example, suppose a=ones((3,3)). Then a+=3j is different than a=a+3j While they both perform the same computation, a+=3j casts the result to fit back in a, while a=a+3j re-binds the name a to the result.

### 3.3.4.3 Unary operations

```
__neg__ (self)
__pos__ (self)
__abs__ (self)
__invert__ (self)
```

These functions are called in response to the unary operations (-, +, abs(), ~). With the exception of \_pos\_, these are implemented using ufuncs (negative, absolute, invert). The unary + operator, however simply calls self.copy(), and can therefore be used to get a copy of an array.

```
__complex__ (self)
__int__ (self)
__long__ (self)
__float__ (self)
__oct__ (self)
__hex__ (self)
```

These functions are aslo defined for the ndarray object to handle the operations complex(), int(), long(), float(), oct(), and hex(). They work only on arrays that have one element in them and return the appropriate scalar.



### TIP

The function called to implement many arithmetic special methods for arrays can be modified using the function set\_numeric\_ops. This function is called with keyword arguments indicating which operation(s) to replace. A dictionary is returned containing showing the old functions. By default, these functions are set to the corresponding ufunc.

### 3.3.4.4 Array indexing

More powerful array indexing was an important extension introduced by numarray, and was therefore an important part of the development of numpy.core. In particular, the desire to select elements arbitrary elements based on their position in

the array, and according to a mask was desireable. There are two kinds of indexing available using the X[obj] syntax: basic slicing, and advanced indexing. For the description of this syntax given below, X is the array to-be-sliced and obj is the selection object. Furthermore, define  $N \equiv X$ .ndim. These two methods of slicing have different behavior and are triggered depending on obj. Adding additional functionality yet remaining compatible with old uses of slicing complicated the rules a little. Hopefully, after studying this section, you will have a firm grasp of what kind of selection will be initiated depending on the selection object.



### TIP

in Python  $X[(\exp 1, \exp 2, ..., \exp N)]$  is equivalent to  $X[\exp 1, \exp 2, ..., \exp N]$  as the latter is just syntatic sugar for the former.

### 3.3.5 Basic Slicing

Basic slicing extends Python's basic concept of slicing to N dimensions. Basic slicing occurs when obj is a slice object (constructed by start:stop:step notation inside of brackets), an integer, or a tuple of slice objects and integers. Ellipsis and newaxis objects can be interspersed with these as well. In order to remain backward compatible with a common usage in Numeric, basic slicing is also initiated if the selection object is any sequence (such as a list) containing slice objects, the Ellipsis object, or the newaxis object, but no integer arrays or other embedded sequences.

The standard rules of sequence slicing applies to basic slicing on a per-dimension basis (including using a step index). Some useful concepts to remember include:

- The basic slice syntax is i: j: k where i is the starting index, j is the stopping index, and k is the step  $(k \neq 0)$ . This selects the m elements (in the corresponding dimension) with index values  $i, i+k, \ldots, i+(m-1)k$  where  $m=q+(r\neq 0)$  where q and r are the quotient and remainder obtained by dividing j-i by k: j-i=qk+r, so that i+(m-1)k < j.
- Assume n is the number of elements in the dimension being sliced. Then, if i is not given it defaults to 0 for k > 0 and n for k < 0. If j is not given it defaults to n for k > 0 and -1 for k < 0. If k is not given it defaults to 1. Note that '::' is the same as ':' and means select all indices along this axis.
- If the number of objects in the selection tuple is less than N, then ':' is assumed for any remaining dimensions.

- Ellipsis expand to the number of ':' objects needed to make a selection tuple of the same length as X.ndim. Only one ellipsis is expanded, any others are interpreted as more ':'
- Each newaxis object in the selection tuple serves to expand the dimensions of the resulting selection by one unit-length dimension. The added dimension is the position of the newaxis object in the selection tuple.
- An integer, i, returns the same values as i:i+1 except the dimensionality of the returned object is reduced by 1. In particular, a selection tuple with the  $p^{\text{th}}$  element an integer (and all other entries ':') returns the corresponding sub-array with dimension N-1. If N=1, then the returned object is an Array Scalar. These objects are explained in Chapter 6.
- If the selection tuple has all entries ':' except the  $p^{\text{th}}$  entry which is a slice object i:j:k, then the returned array has dimension N formed by concatenating the sub-arrays returned by integer indexing of elements i, i+k, i+(m-1)k < j,
- Basic slicing with more than one non-':' entry in the slicing tuple, acts like repeated application of slicing using a single non-':' entry, where the non-':' entries are successively taken (with all other non-':' entries replaced by ':'). Thus, X[ind1,...,ind2,:] acts like X[ind1][...,ind2,:] under basic slicing. Note this is NOT true for advanced slicing.
- You may use slicing to set values in the array, but (unlike lists) you can never grow the array. The size of the value to be set in X[obj] = value must be (broadcastable) to the same shape as X[obj].

Basic slicing always returns another *view* of the array. In other words, the returned array from a basic slicing operation uses the same data as the original array. This can be confusing at first, but it is faster and can save memory. A copy can always be obtained if needed using the unary + operator (which has lower precedence than slicing) or the .copy() method.



### TIP

Remember that a slicing tuple can always be constructed as obj and used in the x[obj] notation. Slice objects can be used in the construction in place of the [start:stop:step] notation. For example, x[1:10:5,::-1] can also be implemented as obj=(slice(1,10,5), slice(None,None,-1)); X[obj]. This can be useful for constructing generic code that works on arrays of arbitrary dimension.

### 3.3.6 Advanced selection

Advanced selection is triggered when the selection object, obj, is a non-tuple sequence object, an indurray (of data type integer or bool), or a tuple with at least one sequence object or indurray (of data type integer or bool). There are two types of advanced indexing: integer and boolean. Advanced selection always returns a copy of the data (contrast with basic slicing that returns a view).

### 3.3.6.1 Integer

Integer indexing allows selection of arbitrary items in the array based on their N-dimensional index. This kind of selection occurs when advanced selection is triggered and the selection object is not an array of data type bool. For the discussion below, when the selection object is not a tuple, it will be referred to as if it had been promoted to a 1-tuple, which will be called the selection tuple. The rules of advanced integer-style indexing are:

- if the length of the selection tuple is larger than N(=X.ndim) an error is raised.
- all sequences and scalars in the selection tuple are converted to intp indexing arrays.
- all selection tuple objects must be convertible to intp arrays, or slice objects, or the Ellipsis (...) object.
- Exactly one Ellipsis object will be expanded, any other Ellipsis objects will be treated as full slice (':') objects. The Ellipsis object is replaced with as many full slice (':') objects as needed to make the length of the selection tuple N.
- If the selection tuple is smaller than N, then as many ':' objects as needed are added to the end of the selection tuple so that the modified selection tuple has length N.

- The shape of all the integer indexing arrays must be broadcastable to the same shape. Arrays are broadcastable if any of the following are satisfied
  - 1. The arrays all have exactly the same shape.
  - 2. The arrays all have the same number of dimensions and the length of each dimensions is either a common length or 1.
  - 3. The arrays that have too few dimensions can have their shapes prepended with a dimension of length 1 to satisfy property 2.
- The shape of the output (or the needed shape of the object to be used for setting) is the broadcasted shape.
  - **Example:** If a.shape is (5,1), b.shape is (1,6), c.shape is (6,) and d.shape is () so that d is a scalar, then a, b, c, and d are all broadcastable to dimension (5,6). The array "a" acts like a (5,6) array where a[:,0] is broadcast to the other columns, "b" acts like a (5,6) array where b[0,:] is broadcast to the other rows, "c" acts like a (1,6) array and therefore a (5,6) where c[:] is broadcast to every row, and finally "d" acts like a (5,6) array where the single values is repeated.
- After expanding any ellipses and filling out any missing (':') objects in the selection tuple, then let  $N_t$  be the number of indexing arrays, and let  $N_s = N N_t$  be the number of slice objects. Note that  $N_t > 0$  (or we wouldn't be doing advanced integer indexing).
- If  $N_s = 0$  then the M-dimensional result is constructed by varying the index tuple  $(i_1, \ldots, i_M)$  over the range of the result shape and for each value of the index tuple setting:

result
$$[i_1, \ldots, i_M] = X[\operatorname{ind}_1[i_1, \ldots i_M], \operatorname{ind}_2[i_1, \ldots, i_M], \operatorname{etc.}, \operatorname{ind}_N[i_1, \ldots, i_M].$$

- **Example:** Suppose the shape of the broadcasted indexing arrays is 3-dimensional and N is 2. Then the result is found by letting i, j, k run over the shape found by broadcasting  $\operatorname{ind}_1$ , and  $\operatorname{ind}_2$ , and for each i, j, k setting  $\operatorname{result}[i, j, k] = X[\operatorname{ind}_1[i, j, k], \operatorname{ind}_2[i, j, k]].$
- If  $N_s > 0$ , then partial indexing is done. This can be somewhat mind-boggling to understand, but if you think in terms of the shapes of the arrays involved, it can be easier to grasp what happens. In simple cases (i.e. one indexing array and N-1 slice objects) it does exactly what you would expect (concatenation of repeated application of basic slicing). The rule for partial indexing is that

the shape of the result (or the interpreted shape of the object to be used in setting) is the shape of X with the indexed subspace replaced with the broadcasted indexing subspace. If the index subspaces are right next to each other, then the broadcasted indexing space directly replaces all of the indexed subspaces in X. If the indexing subspaces are separated (by slice objects), then the broadcasted indexing space is first, followed by the sliced subspace of X.

**Example 1:** Suppose X.shape is (10,20,30) and ind is a (2,3,4) indexing intp array, then result=X[...,ind,:] has shape (10,2,3,4,30) because the (20,)-shaped subspace has been replaced with a (2,3,4)-shaped broadcasted indexing subspace. If we let i, j, k loop over the (2,3,4)-shaped subspace then result[...,i,j,k,:] = X[...,ind[i,j,k],:]. This example produces the same result as X.take(ind,axis=-2).

Example 2: Now let X.shape be (10,20,30,40,50) and suppose ind<sub>1</sub> and ind<sub>2</sub> are broadcastable to the shape (2,3,4). Then X[:,ind<sub>1</sub>,ind<sub>2</sub>] has shape (10,2,3,4,40,50) because the (20,30)-shaped subspace from X has been replaced with the (2,3,4) subpace from the indices. However, X[:,ind<sub>1</sub>,:,ind<sub>2</sub>,:] has shape (2,3,4,10,30,50) because there is no unambiguous place to drop in the indexing subpace, thus it is tacked-on to the beginning. It is always possible to use .transpose() to move the supspace anywhere desired. This example cannot be replicated using take.

### 3.3.6.2 Boolean

This advanced selection occurs when obj is an array object of boolean type (such as may be returned from comparison operators). It is always equivalent to (but faster than) X[obj.nonzero()] where as described above obj.nonzero() returns a tuple (of length obj.ndim) of integer index arrays showing the True elements of obj.

The special case when obj.ndim == X.ndim is worth mentioning. In this case X[obj] returns a 1-dimensional array filled with the elements of X corresponding to the True values of obj. It The search order will be C-style (last index varies the fastest). If obj has True values at entries that are outside of the bounds of X, then an index error will be raised.



#### WARNING

the definition of advanced selection means that X[(1,2,3),] is fundamentally different than X[(1,2,3)]. The latter is equivalent to X[1,2,3] which will trigger basic selection while the former will trigger advanced selection. Be sure to understand why this is True. You should also recognize that x[[1,2,3]] will trigger advanced selection, but X[[1,2,slice(None)]] will trigger basic selection.

#### 3.3.7 Flat Iterator indexing

As mentioned previously, X.flat returns an iterator that will iterate over the entire array (in C-contiguous style with the last index varying the fastest). This iterator object can also be indexed using basic slicing or advanced indexing as long as the selection object is not a tuple. This should be clear from the fact that X.flat is a 1-dimensional view. X.flat can be used for integer indexing using 1-dimensional C-style-flat indices. The shape of any returned array is therefore the shape of the integer indexing object.

# Chapter 4

## **Basic Routines**

### 4.1 Creating arrays

array (object=, dtype=None, copy=True, order=None, subok=False, ndmin=0)

Create a new array of data type, dtype (or determined from object if dtype is None). The shape of the new array will be determined from object. If copy is True, then ensure a copy of the object is made. If copy is False, then the returned object is a copy of the array only if dtype is not equivalent to the data type of object. If order is 'Fortran' then the resulting array will be in Fortran order, otherwise it is in C order. If subok (subclasses are O.K.) is True then pass through subclasses of the array object if possible. If subok is False then only ndarray objects may be returned. The ndmin parameter specifies that the returned array must have at least the given number of dimensions.

asarray (object=, dtype=None, order=None)

Exactly the same as array(...) except the default copy argument is False, and subok is always False. Using this function always returns the base class ndarray.

asanyarray (object, dtype=None, order=None)

Thin wrapper around array(...) with subok=1. You should use this routine if you are only making use of the array attributes, and believe the calculations that will follow would work with any subclass of the array. Use of this routine increases the chance that array subclasses will interact seamlessly with your function — returning the same subclasses.

arange (start=, stop=None, step=1, dtype=None)

Function similar to Python's builtin range() function except it returns an ndarray object. Return a 1-d array of data type, dtype (or determined from the start, stop, and step objects if None), that starts at start, ends before stop and is incremented by step. The returned array has length n where

$$n = \left\lceil \frac{\text{stop} - \text{start}}{\text{step}} \right\rceil$$

with element i equal to start +i step. If stop is None, then the first argument is interpreted as stop and start is 0.



#### NOTE

By defintion of the ceiling function (denoted by  $\lceil x \rceil$ ), we know that  $x \leq \lceil x \rceil < x+1$ , therefore this defintion of the length of arange guarantees that  $\text{start}+n\cdot\text{step} \geq \text{stop}$  as well as  $\text{start}+(n-1)\cdot\text{step} < \text{stop}$ .

#### isfortran (arr)

Equivalent to arr.flags.fnc

Return an uninitalized array of data type, dtype, and given shape. The memory layout defaults to C-style contiguous, but can be made Fortran-style contiguous with a 'Fortran' order keyword.

#### empty\_like (arr)

Syntatic sugar for empty(a.shape, a.dtype, isfortran(arr))

Return an array of data type dtype and given shape filled with zeros. The memory layout may be altered from the default C-style contiguous with the order keyword.

#### zeros\_like (arr)

Syntatic sugar for zeros(a.shape, a.dtype, isfortran(arr))

Syntatic sugar for a = zeros(shape, dtype, order); a+= 1.

from string (string=,dtype=int, count=-1, sep=")

If sep is ", then return a new 1-d array with data-type descriptor given by dtype and with memory initialized (copied) from the raw binary data in string. If count is non-negative, the new array will have count elements (with a ValueError raised if count requires more data than the string offers), otherwise the size of the string must be a multiple of the itemsize implied by dtype, and count will be the length of the string divided by the itemsize.

If sep is not ", then interpret the string in ASCII mode with the provided separator and convert the string to an array of numbers. Any additional whitespace will be ignored.

fromfile (file=, dtype=int, count=-1, sep=")

Return a 1-d array of data type, dtype, from a file (open file object or string with the name of a file to read). The file will be read in binary mode if sep is the empty string. Otherwise, the file will be read in text mode with sep providing the separator string between the entries. If count is -1, then the size will be determined from the file, otherwise, up to count items will be read from the file. If fewer than count items are read, then a RunTimeWarning is issued indicating the number of items read.

frombuffer (buffer, dtype=intp, count=-1, offset=0)

Very similar to (binary-mode) from tring in interpretation of the arguments, except buffer can be any object exposing the buffer interface (or any object with a \_buffer\_ attribute that returns a buffer exposing the buffer protocol). The new array shares memory with the buffer object. The new array will be read-only if the buffer does not expose a writeable buffer.

fromiter (iterator\_or\_generator, dtype=None)

Construct an array from an iterator or a generator. Only handles 1-dimensional cases. By default the data-type is determined from the objects returned from the iterator.

load (file)

Load a pickled array from an open file. If file is a string, then open a file with that name first. Except for the automatic file opening equivalent to cPickle.load(file)

loads (str)

Load a pickled array from a string. Equivalent to cPickle.loads(str).

```
indices (dimensions, dtype=intp)
```

Return an array of dtype representing n(=len(dimensions)) grids of indices each with variation in a single direction. The returned array has shape (n,)+dimensions. Compare with mgrid.

```
>>> indices((2,3))
array([[[0, 0, 0],
[1, 1, 1]],

[[0, 1, 2],
[0, 1, 2]]])
```

#### fromfunction (function, dimensions, \*\*kwargs)

Construct an array from a function called on a tuple of index grids. The function should be able to take array arguments and process them like ufuncs (use vectorize if it doesn't). The function should accept as many arguments as there are dimensions which is a sequence of numbers indicating the length of the desired output for each axis. Keyword arguments to function may also be passed in as keywords to fromfunction.

```
>>> print fromfunction(lambda i,j: i+j, (2,3))
[[0 1 2]
[1 2 3]]
```

```
identity (n, dtype=intp)
```

Return a 2-d array of shape (n,n) and data type, dtype with ones along the main diagonal.

```
where (condition[, x, y])
```

Returns an array shaped like condition, that has the elements of x and y respectively where condition is respectively true or false. If x and y are not given, then it is equivalent to nonzero(condition).

```
lexsort (keys=, axis=-1)
```

Return an array of indices similar to argsort except sorting is done using all of the provided keys. First a sort is computed using key[0], then the indices are further altered by sorting on key[1]. This is repeated until sorting has been performed on all of the keys. This is a useful function for multiple-field sorting.

```
>>> a = [1,2,1,3,1,5]; b = [0,4,5,6,2,3]
>>> ind = lexsort((b,a))
>>> print take(a,ind)
[1 1 1 2 3 5]
>>> print take(b,ind)
[0 2 5 4 6 3]
```

Notice the order the keys had to be used in order to get a lexicographical sorting order. To clarify, suppose three equal-length sequences are fields of an underlying data-type: (f1,f2,f3). If we want to sort first on f1 and then on f2 and then on f3, the indices that would accomplish that sort are obtained as lexsort((f3,f2,f1)).

### 4.2 Operations on two or more arrays

concatenate (seq=, axis=0)

Construct a new array from elements of the sequence object seq concatenated along the given axis. The elements of the sequence object must have compatible types and be the same shape. If axis is None, then flatten each element of seq before concatenating together to construct a 1-d array.

```
correlate (x, y, mode='valid')
```

Compute the 1-d cross correlation of x and y keeping portions determined by mode which may be 'valid' (0), 'same' (1), or 'full' (2). The 'full' cross-correlation between two 1-d arrays is computed as

$$z\left[n\right] = \sum_{i=\max(n-M,0)}^{\min(n,K)} x\left[i\right] y\left[n+i\right],$$

for n=0...K+M where  $K=\operatorname{len}(x)$ -1 and  $M=\operatorname{len}(y)$ -1, and we assume  $K\geq M$  (without loss of generality because we can interchange the roles of x and y without effect). For this formula to work, we assume that x[i]=0 when  $i\notin [0,K-1]$  and y[j]=0 when  $j\neq [0,M-1]$ .

If mode is 'same' then only the K middle values are returned starting at  $n = \lfloor \frac{M-1}{2} \rfloor$ . If the flag has a value of 'valid' then only the middle K - M + 1 = (K+1) - (M+1) + 1 output values are returned starting at n = M.

convolve (x, y, mode='valid')

Convolution is very similar to correlation except it is defined with one sequence reversed:

$$z[n] = \sum_{i} x[i]y[n-i].$$

The mode keyword has the same effect as it does for correlation. Convolution ('full') between two 1-d arrays implements polynomial multiplication where the array entries are viewed as coefficients for polynomials.

**Example:** Consider that  $(x^3+4x^2+2)(x^4+3x+1)=x^7+4x^6+5x^4+13x^3+4x^2+6x+2$ . This can be determined by using the code convolve([1,4,0,2], [1,0,0,3,1]) which returns [1,4,0,5,13,4,6,2]. Notice the one-to-one alignment between the elements of the arrays and the coefficients on powers of x in the polynomial.

outer (a, b)

compute an outerproduct which is syntatic sugar for a.ravel() [:,newaxis] \* b.ravel() [newaxis,:] (after first converting a and b to ndarrays).

```
>>> print outer([1,2,3],[10,100,1000])
[[ 10 100 1000]
[ 20 200 2000]
[ 30 300 3000]]
```

inner (a, b)

Computes the inner product between two arrays. This is an array that has shape a.shape[:-1] + b.shape[:-1] with elements computed as the sum of the product of the elements from the last dimensions of a and b. In particular, let I and J be the super<sup>1</sup> indices selecting the 1-dimensional arrays a[I,:] and b[J,:], then the resulting array, r, is

$$r[I,J] = \sum_k a[I,k]b[J,k].$$

 $<sup>^1\</sup>mathrm{A}$  super index is 0 or more integer indices used to index into an N-dimensional array. How many indices a super index represents should be implied by context.

dot (a, b)

Computes the dot (matrix) product between two arrays. The product-sum is over the last dimension of a and the second-to-last dimension of b. Specifically, if I and J are super indices for a[I,:] and b[J,:,j] so that j is the index of the last dimension of b. Then, the shape of the resulting array is a.shape[:-1] + b.shape[:-2] + (b.shape[-1],) with elements.

$$r[I, J, j] = \sum_{k} a[I, k]b[J, k, j],$$

vdot (a, b)

This computes the dot product between two arrays (flattened into one-dimensional vectors) after conjugating the first vector. This is an inner-product following the physicists convention of conjugating the first argument.

$$r = \sum_{k} \overline{\operatorname{a.flat}[k]} \operatorname{b.flat}[k].$$

**cross** (a, b, axisa=-1, axisb=-1, axisc=-1, axis=None)

Returns the cross product of two (arrays of) vectors. The cross product is performed over the axes of the input arrays indicated by the axisa, and axisb arguments. For both arrays, the axis used must have dimension either 2 or 3. If both axes used have dimension 2, then only the z-component of the equivalent 3-d cross product is returned. Otherwise, the entire vector is returned. The axisc argument gives the axis of the vectors in the returned cross-product result. If axis is not None, then it is assumed that axisa=axisb=axisc=axis (regardless of what else is specified).

**allclose** (a, b, rtol=
$$10^{-5}$$
, atol= $10^{-8}$ )

Returns true if all components of a and b are equal subject to the given relative and absolute tolerances. This returns true if every element of a and b satisfy

$$|a-b| < \operatorname{atol} + \operatorname{rtol} |b|$$
.

### 4.3 Printing arrays

array2string (a)

The default printing mechanism uses this function to produce a string from an array.

**set\_printoptions** (precision=None, the shold=None, edge items=None, linewidth=None, suppress=None)

Set options associated with representing an array.

**precision** the default number of digits of precision for floating point output (default 8);

threshold total number of array elements which triggers summarization rather than full representation (default 1000);

**edgeitems** number of array elements in summary at beginning and end of each dimension (default 3);

**linewidth** the number of characters per line (default 75);

**suppress** Boolean value indicating whether or not to suppress printing of small floating point values using scientific notation (default False).

#### get\_printoptions ()

Returns the values of precision, threshold, edgeitems, linewidth, and suppress that control printing of arrays.

```
set_string_function (func, repr=1)
```

Set the function to use in response to str(array) or repr(array). By default this function is array2string. The function passed in must take an array argument and return a string.

### 4.4 Functions redundant with methods

Several functions are available for purposes of backward compatibility with old Numeric, and are therefore redundant. The functions are all simple wrappers for asarray(a).<function>(\*args, \*\*kwds), or are replaceable by attribute access. The following list documents them. It is not recommended that these functions be used in new programs, but there are no plans for removing them as in functional form they work with arbitrary sequences. The functions that mirror methods and attributes are: take, reshape, squeeze, choose, repeat, put, putmask, swapaxes, transpose, real, imag, sort, argsort, amax, argmax, amin, argmin, ptp, alen, searchsorted, diagonal, trace, ravel, nonzero, shape, compress,

clip, sum, cumsum, product, cumproduct, sometrue (method is .any), all-true (method is .all), around (method is .round), rank (attribute is .ndim), shape, size (.size or .shape[axis]), and copy.

### 4.5 Dealing with data types

```
dtype (obj, align=0)
```

Return a data-type object from any object. See Chapter 7 for a more detailed explanation of what can be interpreted as a data-type object and the meaning of the align keyword.

#### maximum\_sctype (arg)

Returns the array-scalar type of highest precision of the same general kind as arg which can be any recognized form for describing a data-type.

#### issctype (obj)

Returns True if obj is an array data type (or a recognized alias for one)

```
obj2sctype (obj, default=None)
```

Returns the array type object corresponding to obj which can be an array type already, a python type object, an actual array, or any recognized alias for an array type object. If no suitable data type object can be determined, return default.

#### sctype2char (sctype)

Return the typecode character associated with an array-scalar type dtype. The first argument is first converted to a dtype if it needs to be.



#### TIP

the type attribute of data-type objects are actual Python type objects subclassed in a hierarchy of types. This can often be useful to check data types generically. For example, issubclass(dtype.type, integer) can check to see if the data type is one of the 10 different integer types. The issubclass function, however, raises an error if either argument is not an actual type object. NumPy defines issubclass\_(arg1, arg2) that will return false instead of raise an error. Alternatively, dtype.kind is a character describing the class of the data-type so dtype.kind in 'iu' would also check to see if the data-type is an integer type.

can\_cast (from=d1, to=d2)

Return Boolean value indicating whether or not data type d1 can be cast to data type d2 safely (without losing precision or information).

# Chapter 5

# Additional Convenience Routines

### 5.1 Shape functions

atleast\_1d (a1,a2,...,an)

Force a sequence of arrays (including array scalars) to each be at least 1-d.

 $atleast\_2d$  (a1,a2,...,an)

Force a sequence of arrays (including array scalars) to each be at least 2-d. Dimensions of length 1 are pre-pended to reach a two-dimensional array.

 $atleast\_3d$  (a1,a2,...,an)

Force a sequence of arrays (including array scalars) to each be at least 3-d. Dimensions of length 1 are pre-pended to reach a two-dimensional array.

vstack (seq)

Stack a sequence of arrays along the first axis (row wise). Arrays in seq must have the same shape along all dimensions but the first. Rebuilds array divided by vsplit. All 1-d arrays will be stacked row-wise.

hstack (seq)

Stack a sequence of arrays along the second axis (column wise). Arrays in seq must have the same shape along all dimensions but the second. Rebuilds array divided by hsplit. Notice that 1-d arrays will be appended into a new

1-d array. Use column\_stack to get a 2-d array from 1-d arrays. If some arrays are already 2-d, then the 1-d arrays need to have a dimension added to the end (e.g. y[:,newaxis]) in order to stack correctly.

#### column\_stack (seq)

Stack a sequence of 1-d arrays as columns into a 2-d array. All arrays must have the same length. Compare with hstack(seq).

#### row\_stack (seq)

Stack a sequence of 1-d arrays as rows into a 2-d array (alias for vstack).

#### dstack (seq)

Stack a sequence of arrays along the third axis (depth wise). Arrays in seq must have the same shape along all dimensions but the third. Rebuilds array divided by vsplit.

#### array\_split (ary, i\_or\_s, axis=0)

Divide ary into a list of sub-arrays along the specified axis. The i\_or\_s argument stands for indices\_or\_sections. If i\_or\_s is an integer, ary is divided into that many equally-sized arrays. If it is impossible to make an even split, each of the leading arrays in the returned list have one additional member. If i\_or\_s is a list of sorted integer, its entries define the indexes where ary is split. An empty list for i\_or\_s results in a single sub-array equal to the original array.

#### split (ary, i\_or\_s, axis=0)

The same as array\_split() except if i\_or\_s is an integer and it is impossible to make an even split, an error is raised.

#### hsplit (ary, i\_or\_s)

Split a single array into multiple columns of sub-arrays (along the first axis if 1-d or along the second second if >1-d). Only works on arrays of 1 or more dimension.

#### vsplit ()

Split a single array into multiple rows of sub-arrays (along the first axis). Only works on arrays of 2 or more dimensions.

#### dsplit ()

Split a single array into multiple sub-arrays along the third axis (depth). Only works on arrays of 3 or more dimensions.

```
apply_along_axis (func1d, axis, arr, *args)
```

Execute func1d(arr[sel\_i], \*args) where func1d takes 1-d arrays and arr is an N-d array, where sel\_i is a selection object sufficient to select a 1-d sub-array along the given axis. The function is executed for all 1-d arrays along axis in arr.

```
apply_over_axes (func, a, axes)
```

For each axis in the axes sequence, call func as res=func(a, axis). If res is the same shape as a then set a=res and continue. if res.ndim = a.ndim -1, then insert a dimension before axis and continue.

```
expand_dims (a, axis)
```

Expand the shape of array a by including newaxis **before** the given axis.

```
resize (a, new_shape)
```

Returns a new array with the specified shape which can be any size. The new array is filled with repeated copies of a. This function is similar in spirit to a.resize(new\_shape) except in how it fills the new array and in what it returns.

### 5.2 Basic functions

```
average (a, axis=0, weights=None, returned=0)
```

Computes the average along the indicated axis. If axis is None, average over the entire array. Inputs can be integer or floating types; result is type float. If weights are given, the result is sum(a\*weights)/sum(weights). Therefore, weights must have shape equal to a.shape or be 1-d with length a.shape[axis]. Integer weights are converted to float. If returned is True, then return a tuple showing both the result and the sum of the weights (or count of the values). The shape of these two results will be the same.

```
cov (x, y=None, rowvar=1, bias=0)
```

Compute the covariance matrix of data in x. If x is a vector and y is None, then this function is equivalent to asarray(x).var(). Otherwise, x is interpreted as observations of several random variables. If rowvar is True (default), then the variables are in the rows and the observations of the variables are in the columns. Otherwise, the variables are in the columns and the observations

are in the rows. If y is given then it is treated as another variable or set of variables to be added to x. By default, an unbiased estimate of the covariance matrix is made, if bias is non-zero, then a biased normalization factor is used instead. If X is a random vector, then the covariance matrix is defined as

$$\mathbf{C} = E\left[\mathbf{X}\mathbf{X}^H\right].$$

It can be approximated as

$$\mathbf{C} \approx \frac{1}{P} \sum_{i=0}^{N-1} \mathbf{x}_i \mathbf{x}_i^H$$

where  $\mathbf{x}_i$  is an observation of  $\mathbf{X}$  (as a column-vector), N is the number of observations made and P = N - 1 for an unbiased estimate or P = N for a biased (but lower mean-squared error) estimate.

**corrcoef** (x, y=None, rowvar=1, bias=0)

Estimate the correlation coefficient of x. By default, each row of x contains a random variable with observations of the random variable in the columns of x. (If rowvar is False, the each column is a random variable with observations in the rows). The y argument can be used to append additional variables to x. The  $i^{\rm th}$  row and  $j^{\rm th}$  column of the correlation coefficient matrix is defined as

$$\rho_{ij} = \frac{C_{ij}}{\sqrt{C_{ii}C_{jj}}}$$

where C is the covariance matrix. The rowvar and bias arguments are passed on to the cov function to estimate C.

#### msort (a)

Return a new array, sorted along the first axis. Equivalent to b=a.copy(); b.sort(0) median (m)

Returns the median of m along its first dimension.

The list argument is a 1-d integer array. Let r be the returned 1-d array whose length is (list.max()+1). If weights is None, then r[i] is the number of occurrences of i in list. If weight is present, then the i<sup>th</sup> element is

$$r[i] = \sum_{j: \text{list}[j]=i} \text{weights}[j].$$

Notice that if weights is None, it is equivalent to a weights array of all 1. The length of weights must be the same as the length of list.

#### **digitize** (x=,bins=)

Return an array of integers the same length as x with values i such that bins  $[i-1] \le x < \text{bins}[i]$  if bins is monotonically increasing, or  $\text{bins}[i] \le x < \text{bins}[i-1]$  if bins is monotonically decreasing. When x is beyond the bounds of bins, return either i = 0 or i = len(bins) as appropriate.

**histogram** (x=, bins=None, range=None, normed=0)

Construct a histogram for the data in x (treated as one-dimensional array of type float). If bins is not a sequence, then bins should be the number of bins which will be constructed ranging range[0] to range[1] or x.min() to x.max() if range is None. If normed is True, then the histogram will be normalized and comparable with a probability density function, otherwise it will be a count of the number of items in each bin. The return value is the tuple (n, bins) where n is the histogram.

logspace (start, stop, num=50, endpoint=1)

Evenly spaced samples on a logarithmic scale. Returns num evenly spaced (in logspace) samples from  $10^{**}$ start to  $10^{**}$ stop. If endopint is 1, then the last sample is  $10^{**}$ stop.

**linspace** (start, stop, num=50, endpoint=1, retstep=0):

Evenly spaced samples. Returns num evenly spaced samples from start to stop. If endpoint is True, then the last sample is stop. If retstep is True, then return the computed step size.

**select** (condlist, choicelist, default=0)

Returns an array comprised from different elements of choicelist depending on the list of conditions. The condlist argument is a list of boolean condition arrays. The choicelist argument is a list of choice arrays (of the same size as the arrays in condlist). The result has the same size as the arrays in choicelist. If condlist is  $[c_0, \ldots, c_{N-1}]$ , then choicelist must be of length N. The elements of choicelist can then be represented as  $[v_0, \ldots, v_{N-1}]$ . The default choice if none of the conditions are met is given as the default argument. The conditions are tested in order and the first one satisfied is used to select the choice. In other words, the elements of the output array are found from the following tree (evaluated on an element-by-element basis)

```
\begin{array}{ll} \text{if } c_0 \colon v_0 \\ \\ \text{elif } c_1 \colon v_1 \\ \\ \dots \\ \\ \text{elif } c_{N-1} \colon v_{N-1} \\ \\ \text{else: default} \end{array}
```

piecewise (x, condlist, funclist, \*args, \*\*kw)

Compute a piecewise-defined function. A piecewise defined function is

$$f(x) = \begin{cases} f_1(x) & x \in S_1, \\ f_2(x) & x \in S_2, \\ \vdots & \vdots \\ f_n(x) & x \in S_n. \end{cases}$$

where  $S_1$  are sets. Thus, the function is defined differently over different subdomains of the input. Such a function can be computed using **select** but such an implementation means calling each  $f_i$  over the entire region of x. The piecewise call guarantees that each function  $f_i$  will only be called over those values of x in  $S_i$ .

Arguments: x is the array of values over which to call the function; condlist is a sequence of boolean (indicator) arrays (or a single boolean array) of the same shape as x that defines the sets (True indicates that element of x is in the set). If needed, to match the length of funclist, an "otherwise" set will be added to condlist. This otherwise set is defined as  $S_n = \overline{\bigcup S_i}$ . The argument funclist is a list of functions to be called (or items to be inserted) corresponding to the conditions. Each of these functions can take extra arguments and key-word arguments which are passed in as \*args, and \*\*kw using standard Python syntax. Each of these functions should return vector output for vector input. If the function is a scalar, then it will simply be inserted where appropriate into the output. It is the equivalent of a constant function.

**Example:** Suppose we want to compute  $f(x) = x^2 \Pi\left(\frac{x}{3}\right) + u(x-5)$  where  $\Pi(x) = 1$  only when  $|x| \le 1$  and u(x) = 1 only when  $x \ge 0$ . This could be done using the code:

```
trim_zeros (filt, trim='fb'):
```

Trim the leading ('f' in trim) and trailing ('b' in trim) zeros from a sequence according to the trim keyword.

```
trapz (y, x=None, dx=1.0, axis=-1)
```

If  $\mathbf{y}$  contains samples of a function:  $y_i = f\left(x_i\right)$  then trapz can be used to approximate the integral of the function using the trapezoidal rule. If the sampling is not evenly spaced use  $\mathbf{x}$  to pass in the sample positions. Otherwise, only the sample-spacing is needed in dx. The trapz function can work with many functions at a time stored in an N-dimensional array. The axis argument controls which axis defines the sampling axis (the other dimensions are different functions). The number of dimensions of the returned result is y-ndim - 1.

```
diff (x, n=1, axis=-1)
```

Calculates the  $n^{\text{th}}$  order, discrete difference along the given axis.

```
gradient (f, *varargs)
```

Cacluate the gradient of an N-d scalar function, f. Uses central differences on the interior and first differences on boundaries to give the same shape for each component of the gradient. varargs can contain 0, 1, or N scalars corresponding to the sample distances in each direction (default 1.0). If f is N-d, then N arrays are returned each of the same shape as f, giving the derivative of f with respect to each dimension.

```
angle (z, deg=0)
```

Return the angle of a complex number z (in degrees if deg is True).

```
unwrap (p, discont=pi, axis=-1)
```

Unwraps raidan phase p by changing absolute jumps greater than discont to their  $2\pi$  complement along the given axis.

#### sort\_complex (x)

This is syntatic sugar for asarray(x).sort().astype(<cmplx\_type>) where cmplx\_type is csingle if x.dtype is integral with fewer bits than intp, clongfloat if x.dtype is longfloat, and cdouble for all other types. The sorting is done by comparing the real part of the array, and then the imaginary part if the real parts are the same.

```
disp (mesg, device=None, linefeed=1)
```

Display a message to device (defaults to sys.stdout) with or without a closing linefeed.

```
unique (seq)
```

Returns unique items in the 1-dimensional seq.

```
extract (condition, arr)
```

Equivalent to arr.compress(condition.flat) and arr.flat[abool(condition.flat)] which extracts the elements of (flattened) arr according to the elements of (flattened) condition that are True.

```
insert (arr, mask, vals)
```

Inverse of extract. Equivalent to arr[abool(mask)] = vals but it uses a different algorithm. Not clear which is faster, yet.

```
nansum (x, axis=-1)
nanmax (x, axis=-1)
nanargmax (x, axis=-1)
nanargmin (x, axis=-1)
nanmin (x, axis=-1)
```

These functions perform their respective operations over the given axis (or the entire array if axis is None), after replacing any nans with appropriate values so as not to affect the calculation.

#### vectorize

```
<u>__init__</u> (pyfunc, otypes=None, doc=None)
```

This is a class with whose \_\_init\_\_ signature is given. Instances of this class have a call method that invokes a ufunc that has been dynamically built to call the python function pyfunc internally. The output types can be controlled by the otypes argument. If it is None, then the output types will be determined upon first call to the function using the provided inputs. This can be reset, by re-setting the otypes attribute to "". The normal rules of array broadcasting are followed by the returned object.

#### asarray\_chkfinite (x)

Like asarray(x) except an error is raised if any of the values in x are not finite.

```
round_{-} (x, decimals=0)
```

Return an array with all the elements of x rounded to decimals places. Returns x if array is not floating point and rounds both the real and imaginary parts separately if array is complex. Rounds in the same way as standard python except for half-way values are rounded to the nearest *even* number.

#### add\_docstring (obj, doc)

Adds a docstring to a builtin object, obj, that does not have a docstring defined already. The obj can be a builtin function-or-method, a typeobject, a method descriptor, a getset descriptor, or a member descriptor. This is useful for improving the documentation of objects defined in C-compiled code without re-compiling.

#### add\_newdoc (place, obj, doc)

Adds a docstring to the 'obj' imported from 'place' using exec 'from %s import %s' % (place, obj). If doc is a string, then a single docstring is added to obj from place. If doc is a 2-tuple, then obj must be an object with attributes that need to be commented. The first element of the doc tuple is the attribute

to be commented on and the second element is the actual docstring. If doc is a list, then it must be composed of elements that are 2-tuples indicating that obj has several attributes that need to be documented.

### 5.3 Polynomial functions

There are two interfaces for dealing with polynomials: a class-based interface, and a collection of functions to deal with a polynomials represented as a simple list of coefficients. This latter representation results from the is a one-to-one correspondence between a length-(n+1) sequence of coefficients  $a_n \equiv a[n]$  and an  $n^{\text{th}}$  order polynomial:

$$p(x) = a_0 x^n + a_1 x^{n-1} + \dots + a_{n-1} x + a_n.$$

Most of the functions below operate on and return a simple sequence of coefficients representing a polynomial. There is, however, a simple polynomial class that provides some utility for doing simple algebra on polynomials.

This construction returns an instance of a simple polynomial class. It can take either a list of coefficients on polynomial powers, or a sequence of roots (if r=1). The returned polynomial can be added, subtracted, multiplied, divided, and taken to integer powers, resulting in new polynomials.

- .r roots of the polynomial
- .o order of the polynomial
- .c polynomial coefficients as an array (also \_\_array\_\_() )
- **\_\_call\_\_(x)** evaluate the polynomial at x (can be an array)
- **\_\_getitem\_(x)** p[k] returns the coefficient on the kth power of x (backwards from indexing the coefficient array)

```
>>> p=poly1d([2,5,7])
>>> print p
2
2 x + 5 x + 7
>>> print p*[1,3,1]
4 3 2
2 x + 11 x + 24 x + 26 x + 7
>>> print p([0.5,0.6,3])
[ 10.  10.72 40. ]
>>> print p.r
[-1.25+1.3919j -1.25-1.3919j]
```

#### poly (roots\_or\_matrix)

Return a sequence of coefficients representing a polynomial given the sequence of roots as an argument. Alternatively, if the argument is a 2-d array, then return the characteristic polynomial of the matrix.

#### roots (poly)

Return the roots of the polynomial represented by coefficients in poly

Return an exact  $m^{\text{th}}$ -order integral of the polynomial represented in poly. If k is None, then use 0 for the integrating constants. Otherwise, use the scalars in the sequence k as integrating constants. Also available as .integ (m=1,k=0) method of poly1d objects.

#### Example:

$$p(x) = x^{2} + 3x + 4$$

$$\int \int p(x) = \frac{1}{12}x^{4} + \frac{1}{2}x^{3} + 2x^{2} + k_{0}x + k_{1}$$

#### polyder (poly, m)

Return an exact  $m^{\text{th}}$ -order derivative of the polynomial represented in poly. Also available as .deriv(m=1) method of poly1d objects.

#### Example:

$$p(x) = x^3 + 2x^2 + 4x + 3$$
  
 $\frac{dp}{dx}(x) = 3x^2 + 4x + 4$ 

```
>>> polyder([1,2,4,3])
array([3, 4, 4])
```

#### polyadd (p1, p2)

Add the two polynomials represented by coefficients:  $p_1(x) + p_2(x)$ 

#### polysub (p1, p2)

Return coefficients for the polynomial found by subtracting the two polynomials represented by  $p_1$  and  $p_2$ :  $p_1(x) - p_2(x)$ 

#### polymul (p1, p2)

Return the coefficients for  $p_1(x) p_2(x)$ 

#### polydiv (p1, p2)

Return the quotient, q(x), and remainder, r(x), so that  $p_1(x) = q(x) p_2(x) + r(x)$ , with the order of r(x) less than the order of  $p_2(x)$ . This function requires the **lfilter** command in the signal processing library of full NumPy, which must be installed separately.

#### polyval (p, y)

Evaluate the polynomial p at y. The argument, y, can be a number or an array or a polynomial object. If x is a polynomial object, then polyval performs polynomial composition: p(y(x)), otherwise polyval computes the value of the polynomial at each y. Uses Horner's rule for evaluation, but this can still lead to numerical instabilities for wildly fluctuating coefficients.

#### polyfit (x,y,N)

Compute a best-fit polynomial in x of order N, to the data, y, in the sense of minimizing averaged-squared error between the measurement and the model. Useful for quick line-fitting.

### 5.4 Set Operations

The set operations were kindly contributed by Robert Cimrman. These set operations are based on sorting functions and all expect 1-d sequences with unique elements with the exception of unique1d and intersect1d\_nu which will flatten N-d nested-sequences to 1-d arrays and can handle non-unique elements.

#### unique1d (arr, retindx=False)

Return the unique elements of arr as a 1-d array. If retindx is True, then also return the indices, ind, such that arr.flat[ind] is the set of unique values.

#### intersect1d (a1, a2)

Return the (sorted) intersection of a1 and a2 which is an array containing the elements of a1 that are also in a2.

#### intersect1d\_nu (a1, a2)

Return the (sorted) intersection of a1 and a2 but allow a1 and a2 to be N-d arrays with non-unique elements. Equivalent to intersect1d(unique1d(a1), unique1d(a2)).

#### **union1d** (a1, a2)

Return the (sorted) union of a1 and a2 which is an array containing elements that are in either a1 or a2.

#### setdiff1d (a1, a2)

Return the set-difference of a1 and a2 which is an array containing the elements of a1 that are **not** in a2.

#### **setxor1d** (a1, a2)

Return the (sorted) set containing the exclusive-or of the arrays a1 and a2. The exclusive-or contains elements that are in a1 or in a2 as long as the element is not in both a1 and a2.

#### setmember1d (tocheck, set)

Return a Boolean 1-d array of the length of tocheck which is True whenever that element is contained in set and false when it is not. Equivalent to array([x in set for x in tocheck])

### 5.5 Array construction using index tricks

The functions and classes in this category make it simpler to construct arrays.

#### **ix**<sub>−</sub> (\*args)

This indexing cross function is useful for forming indexing arrays necessary to select out the cross-product of N 1-dimensional arrays. Note that the default indexing does not due a cross-product which someone coming from Matlab might initially expect. The default indexing is more general-purpose. Using the ix\_constructor can produce the indexing arrays necessary to select a cross-product.

#### mgrid [index expression]

This is an instance of a class. It can be used to construct a filled "mesh-grid" using slicing syntax.

#### ogrid [index expression]

This is similar to mgrid except it returns an open grid, so as to save space and time. The broadcasting rules will ensure that any universal function operating on the grid will act as if the ogrid had been the result of mgrid.

#### **r**\_ [index expression]

This is a simple way to build up arrays quickly. There are two use cases. 1) If the index expression contains comma separated arrays, then stack them along their first axis. 2) If the index expression contains slice notation or scalars then create a 1-d array with a range indicated by the slice notation. In otherwords the slice syntax start:stop:step is equivalent to arange(start, stop, step) inside of the brackets. However, if step is a complex number (i.e. 100j) then its integer portion is interpreted as a number-of-points desired and the start and stop are inclusive. In other words start:stop:stepj is interpreted as linspace(start, stop, step, endpoint=1) inside of the brackets. After expansion of slice notation, all comma separated sequences are concatenated together.

Optional character strings inserted into the index expression can be used to change the output. The strings 'r' or 'c' result in matrix output. If the result is 1-d and 'r' is specified a  $1 \times N$  (row) matrix is produced. If the result is 1-d and 'c' is specified, then a  $N \times 1$  (column) matrix is produced. If the result is 2-d then both provide the same result. A string integer specifies which axis to stack multiple comma separated arrays along.

```
>>> print r_{-1:1:9j,[0]*10,5,6]
       -0.75 -0.5 -0.25 0.
Γ-1.
                                 0.25 0.5
                                                                0.
      0.
            0.
                  0.
                         0.
                               0.
                                     0.
                                            0.
                                                  5.
                                                        6. 1
>>> print r_[1,2,5,6,'r']
[[1 2 5 6]]
>>> print r_[1,2,5,6,'c']
[[1]
[2]
[5]
[6]]
>>> a=arange(6).reshape(2,3)
>>> r_[a,a]
array([[0, 1, 2],
[3, 4, 5],
[0, 1, 2],
[3, 4, 5]])
>>> r_[a,a,'-1']
array([[0, 1, 2, 0, 1, 2],
[3, 4, 5, 3, 4, 5]])
```

### 5.6 Other indexing devices

index\_exp [index expression]

Return a tuple of Python objects that implements the index expression and can be modified and placed in any other index expression.

```
>>> index_exp[2:5,...,4,::-1]
(slice(2, 5, None), Ellipsis, 4, slice(None, None, -1))
```

#### $\mathbf{s}_{-}$ [index expression]

Translate index expressions into the equivalent Python objects. This is similar to index\_expression except a tuple is not always returned. For example:

```
>>> s_[1:10]
slice(1, 10, None)
>>> s_[1:10,-3:4:0.5]
(slice(1, 10, None), slice(-3, 4, 0.5))
```

This provides a standard way to construct index expressions to pass to functions and methods because Python does not allow slice expressions anywhere except for inside brackets.

#### ndindex (\*seq)

A sequence of N integers are passed in as separate arguments. These integers are used as the upper boundaries of an N-dimensional counter that starts at 0. The object returned is an iterator that implements the counter.

```
>>> for index in ndindex(3,3,2):
... print index
(0, 0, 0)
(0, 0, 1)
(0, 1, 0)
(0, 1, 1)
(0, 2, 0)
(0, 2, 1)
(1, 0, 0)
(1, 0, 1)
(1, 1, 0)
(1, 1, 1)
(1, 2, 0)
(1, 2, 1)
(2, 0, 0)
(2, 1, 1)
(2, 2, 0)
(2, 2, 1)
```

#### unravel\_index (indx, dims)

Convert a flat index, indx, into an index tuple for an array of the given shape. Keep in mind that it may be more convienient to use indx with a.flat, then to unravel the index.

### 5.7 Two-dimensional functions

These functions all deal with or return two dimensional arrays.

```
eye (N, M=None, k=0, dtype=float)
```

Return an  $N \times M$  array of the given type with ones down the  $k^{\text{th}}$  diagonal. If M is None, it defaults to N. Alternatively, if M is a valid data type, then it becomes the datatype used.

```
vander (x, N=None)
```

The Vandermonde matrix of vector, x. The  $i^{\text{th}}$  column of the return matrix is the  $m_i^{\text{th}}$  power of x where  $m_i = N - i - 1$ . If N is None, it defaults to the length of x.

```
>>> vander([1,2,3,4,5],3)
array([[1, 1, 1],
[4, 2, 1],
[9, 3, 1],
[16, 4, 1],
[25, 5, 1]])
```

#### $\mathbf{diag}$ (v, k=0)

Return the  $k^{\text{th}}$  diagonal if v is a 2-d array, or returns an array with v as the  $k^{\text{th}}$  diagonal if v is a 1-d array.

```
>>> diag(arange(12).reshape(4,3),k=1)
array([1, 5])
>>> diag([1,4,5,7],k=-1)
array([[0, 0, 0, 0, 0],
[1, 0, 0, 0, 0],
[0, 4, 0, 0, 0],
[0, 0, 5, 0, 0],
[0, 0, 7, 0]])
```

#### fliplr (m)

Return the array, m, with rows preserved and columns reversed in the left-right direction. For m.ndim > 2, this works on the first two dimensions (equivalent to m[:,::-1])

#### flipud (m)

Return the array, m, with columns preserved and rows reversed in the up-down direction. For m.ndim > 1, this works on the first dimension (equivalent to m[::-1])

```
rot90 (m, k=1)
```

Rotate the first two dimensions of an array, m, by k\*90 degrees in the counter-clockwise direction. Must have m.ndim >=2.

```
tri (N, M=N, k=0, dtype=aint)
```

Construct an  $N \times M$  array where all the diagonals starting from the lower left corner up to the  $k^{\text{th}}$  diagonal are all ones.

```
triu (m, k=0)
```

Return a upper-triangular 2-d array from m with all the elments below the  $k^{\text{th}}$  diagonal set to 0.

```
tril (m, k=0)
```

Return a lower-triangular 2-d array from m with all the elements above the  $k^{\text{th}}$  diagonal set to 0.

```
mat (data, dtype=None, copy=0)
```

Construct a matrix from data. Alias for numpy.core.defmatrix.matrix. The calling syntax is therefore the same as the \_new\_ method of the matrix object.

Note that data can be a string in which case the routine uses Matlab-style syntax to construct the matrix:

```
>>> mat('1 3 4; 5 6 9')
matrix([[1, 3, 4],
[5, 6, 9]])
```

```
bmat (obj, ldict=None, gdict=None)
```

Build a matrix from sub-blocks. This is similar to mat, except the items in the nested-sequence, or string, should be appropriately shaped 2-d arrays. If obj is a string, then ldict and gdict can be used to alter where the names represented in the string are found (default is current local and global namespace).

```
>>> A=mat('1 2; 3 4'); B=mat('5 6; 7 8')
>>> bmat('A, B; B, A')
matrix([[1, 2, 5, 6],
[3, 4, 7, 8],
[5, 6, 1, 2],
[7, 8, 3, 4]])
```

### 5.8 More data type functions

#### iscomplexobj (obj)

Return a single True or False value depending on whether or not obj would be interpreted as an array with complex-valued data type.

#### isrealobj (obj)

Return a single True or False value depending on whether or not obj would be interpreted as an array with real-valued data type.

#### isscalar (obj)

True if obj is a scalar (an instance of an array data type, or a standard Python scalar type). There is also a sequence of Scaladtypes defined in NumPy, so that this can also be tested as type(obj) in NumPy.Scaladtype.

#### nan\_to\_num (arr)

Returns an array with non-finite numbers changed to finite numbers. The mapping converts nan to 0, inf to the maximum value for the data type and -inf to the minimum value for the data type.

```
real_if_close (arr, tol=100)
```

Return a real arr if arr is complex with imaginary parts less than some tolerance. If tol > 1, then it represents a multiplicative factor on the value of epsilon for the data type of arr.

```
cast [dtype_or_alias] (obj)
```

Cast obj to an array of the given type. This is equivalent to array(obj, copy=0).astype(dtype\_or\_al When one type is cast to another in this fashion, a very low-level operation takes place. Typically, you get what your C-compiler produces for the cast.

Thus, for example, an integer array cast to a boolean type (which is internally

8-bits) will preserve low-order bits of the integer in memory. These will be interpreted as True (but you will be able to get back the integer on a re-cast).

```
>>> cast[bool]([1,2,3,4,5]).astype(int)
array([1, 1, 1, 1])
```

**asfarray** (a, dtype=afloat)

Return an array of inexact data type (floating or complexfloating).

mintypecode (typechars, typeset='GDFgdf', default='d')

Return a minimum data type character from typeset that handles all given typechars. The returned type character must correspond to the data type of the smallest size such that an array of the returned type can handle the data from an array of type t for each t in typechars. If the typechars does not intersect with the typeset, then default is returned. If an element of typechars is not a string, then t=asarray(t).dtypechar is applied.

#### finfo (dtype)

This class allows exploration of the details of how a floating point number is represented in the computer. It can be instantiated by an inexact data type object (or an alias for one). Complex-valued data types are acceptable and are equivalent to their real-valued counterparts. The attributes of the class are

**nmant** The number of bits in the floating point mantissa, or fraction.

**nexp** The number of bits in the floating point exponent

machep Exponent of the smallest (most negative) power of 2 that when added to 1.0 gives something different than 1.0.

eps Floating point precision: 2\*\*machep.

**precision** Number of decimal digits of precision: int(-log10(eps))

resolution 10\*\*(-precision)

**negep** Exponent of the smallest power of 2 that, subtracted from 1.0, gives domething different than 1.0.

**epsneg** Floating point precision: 2\*\*negep.

minexp Smallest (most negative) power of 2 producing "normal" numbers (no leading zeros in the mantissa).

tiny The smallest (in magnitude) usable floating point number equal to  $2^{**}$ minexp.

maxexp Smallest (positive) power of 2 that causes overflow.

max The largest usable floating value: (1-epsneg)\* (2\*\*maxep)

min The most negative usable floating value: -max

The most widely used attributes would probably be eps, max, min, and tiny.

#### 5.9 Functions that behave like usuncs

These functions are Python functions built on top of universal functions (ufuncs) and also take optional output arguments. They broadcast like ufuncs but do not have ufunc attributes.

```
fix (x, y=None)
```

Round x to the nearest integer towards zero.

```
isneginf (x, y=None)
```

True if  $x = -\infty$ . Should be the same as x==NumPy.NINF.

**isposinf** (x, y=None)

True if  $x = +\infty$ . Should be the same as x==NumPy.PINF.

log2 (x, y=None)

Compute the logarithm to the base 2 of x. An optional output array may be provided.

```
set_numeric_ops (<op1>=func1, <op2>=func2, ...)
```

This function can be used to alter the operations used for internal array calculations and array special methods. Replaceable operations (and possible entries for <opN>) are add, subtract, multiply, divide, remainder, power, sqrt, negative, absolute, invert, left\_shift, right\_shift, bitwise\_and, bitwise\_or, less, less\_equal, equal, not\_equal, greater, greater\_equal, floor\_divide, true\_divide, logical\_or, logical\_and, floor, ceil, maximum, and minimum. The example code below changes, then restores, the old Numeric behavior of remainder (which was changed because it was not consistent with Python).

### 5.10 Miscellaneous Functions

Some miscellaneous functions are available in NumPy which are included largely for compatibility with MLab of the old Numeric package. One notable difference, however, is that due to a separate implementation of the modified Bessel function, the kaiser window is available without needing a separate library.

#### sinc (x)

Compute the sinc function for x which can be a scalar or array. The sinc is defined as  $y = \text{sinc}(x) = \frac{\sin(\pi x)}{\pi x}$  with the caveat that the limiting value (1.0) of the ratio is taken for x = 0.

**i0** (x)

Modified Bessel function of the first kind of order 0. Needed to compute the kaiser window. The modified Bessel function is defined as

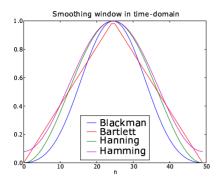
$$I_0(x) = \frac{1}{\pi} \int_0^{\pi} e^{x \cos \theta} d\theta = \sum_{k=0}^{\infty} \frac{x^{2k}}{4^k (k!)^2}.$$

#### blackman (M)

Construct an M-point Blackman smoothing window which is sequence of length M with values given for  $n=0\ldots M-1$  by

$$w[n] = 0.42 - 0.5 \cos\left(2\pi \frac{n}{M-1}\right) + 0.08 \cos\left(4\pi \frac{n}{M-1}\right).$$

#### bartlett (M)



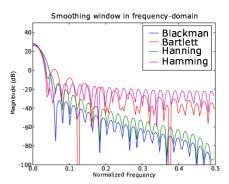


Figure 5.1: Blackman, Bartlett, Hanning, and Hamming windows in the time and frequency domain showing the trade-off between main-lobe width and side-lobe height (Figures made with matplotlib).

Construct an M-point Bartlett (triangular) smoothing window as

$$w[n] = \begin{cases} 2\frac{n}{M-1} & 0 \le n \le \frac{M-1}{2}, \\ 2 - 2\frac{n}{M-1} & \frac{M-1}{2} < n \le M-1. \end{cases}$$

#### hanning (M)

Construct an M-point Hanning smoothing window defined as

$$w\left[n\right] = \frac{1}{2} - \frac{1}{2}\cos\left(2\pi\frac{n}{M-1}\right).$$

#### hamming (M)

Construct an M-point Hamming smoothing window defined for  $n=0\ldots M-1$  as

$$w[n] = 0.54 - 0.46 \cos\left(2\pi \frac{n}{M-1}\right).$$

All of the windowing functions are smoothing windows that attempt to balance the inherent trade off between side-lobe height (ringing) and main-lobe width (resolution) in the frequency domain. A rectangular window has the smallest main-lobe width but the largest side-lobe height. A windowing (tapering) function tries to can help trade off main-lobe width By sacrficing a little in resolution using a windowing functio These windows can be used to smooth data using the convolve function. Figure 5.1 shows the windowing functions described so far and their time- and frequency-domain behavior.

The trade-off between main-lobe and side-lobe has been studied extensively. Solutions that maximize energy in the main-lobe compared to energy in the side-lobes can be found by finding an eigenvector which can be expensive to compute for large window sizes. A good approximation to these prolate-spheroidal windows is the Kaiser window.

#### kaiser (M, beta)

Construct an M-point Kaiser smoothing window. The  $\beta$  parameter controls the width of the window (and the frequency-domain side-lobe height and main-lobe width). The window is defined as

$$w[n] = \frac{1}{I_0(\beta)} I_0 \left( \beta \sqrt{1 - \frac{(2n - M - 1)^2}{(M - 1)^2}} \right).$$

There is an empirical relationship between  $\beta$  and the side-lobe height which can be used in FIR filter design. To achieve a side-lobe height of  $-\alpha dB$ , the  $\beta$  parameter is

$$\beta = \begin{cases} 0.1002 (\alpha - 8.7) & \alpha > 50, \\ 0.5842 (\alpha - 21)^{0.4} + 0.07886 (\alpha - 21) & 21 \le \alpha \le 50, \\ 0 & \alpha < 21. \end{cases}$$

The length M of the window determines the transition width. To obtain a transition width of  $\Delta\omega$ rad/s the window-length must be at least:

$$M = \frac{\alpha - 8}{2.285 \Delta \omega} + 1.$$

# Chapter 6

# Scalar objects

One important new feature of NumPy is the addition of a new scalar object for each of the 21 different data types that an array can have. In fact, the .dtype attribute of an array returns the type object of these new scalars. In addition, all data types are actual type objects for these new scalars. Five (or six) of these new scalar objects are essentially equivalent to fundamental Python types and therefore inherit from them as well as from the generic array scalar type. The bool\_ data type is very similar to the Python BooleanType but does not inherit from it because Python's BooleanType does not allow itself to be inherited from, and on the C-level the size of the actual bool\_ data is not the same as a Python Boolean scalar. Table 6.1 shows which array scalars inherit from basic Python types.

The array scalars have the same attributes and methods as arrays and live in a hierarchy of scalar types so they can be easily determined. However, because array scalars are immutable, and attributes change intrinsic properties of the object, the array scalar attributes are not settable.

Array scalars can be detected using the hierarchy of data types. For exam-

Table 6.1: Array scalar types that inherit from basic Python types. The intc array data type might also inherit from the IntType if it has the same number of bits as the int\_array data type on your platform.

array data type	Python type
int_	IntType
float_	FloatType
complex_	ComplexType
str_	StringType
unicode_	UnicodeType

ple, isinstance(val, generic) will return True if val is an array scalar object. Alternatively, what kind of array scalar is present can be determined using other members of the data type hierarchy. Thus, for example isinstance(val, complexfloating) will return True if val is a complex valued type, while isinstance(val, flexible) will return true if val is one of the flexible itemsize array types (string, unicode, void).



### WARNING

The bool\_type is not a subclass of the int\_type (the bool\_type is not even a number type). This is different than Python's default impelmentation of bool as a sub-class of int.

### 6.1 Attributes of array scalars

The array scalar objects do not have the <u>\_array\_priority\_</u> attribute because they do not inherit from the array type. Other than that, they share the same attributes as arrays.

### flags

Returns True for CONTIGUOUS, OWNDATA, FORTRAN, and ALIGNED. Always returns False for WRITEABLE, and UPDATEIFCOPY.

### shape

Returns ().

#### strides

Returns ().

#### ndim

Returns 0.

#### data

A read-only buffer object of size self.itemsize,

#### size

Return 1.

#### itemsize

The number of bytes this scalar requires.

#### base

Returns None.

### dtype

Returns data type descriptor corresponding to this array scalar.

#### real

The real part of the scalar.

### imag

The imaginary part of the scalar (or 0 if this is real).

#### flat

Return a 1-d iterator object (of size 1).

### \_array\_interface\_\_

The Python-side to the array interface.

#### \_array\_struct\_

The C-side to the array interface

### 6.2 Methods of array scalars

Array scalars have exactly the same methods as arrays. The default behavior of these methods is to internally convert the scalar to an equivalent 0-dimensional array and to call the corresponding array method. The exceptions to these rules are given below. In addition, math operations on array scalars default to the array math operations using 0-dimensional arrays for the array scalars.

```
_new__ (obj)
```

The default behavior is to return a new array or array scalar by calling array(obj) with the corresponding data type. There are two situations when this default behavior is delayed until another approach is tried. First, when the array scalar type inherits from a Python type, then the Python types new method is called first and the default method is called only if that approach fails. The second situation is for the void data type where a single integer-like argument will cause a void scalar of that size to be created and initialized to 0.

Notice that because array(obj) is called for new, if obj is a nested sequence, then the return object could actually be an ndarray. Thus, arrays of the correct type can also be created by calling the array data type name directly:

```
>>> uint32([[5,6,7,8],[1,2,3,4]])
array([[5, 6, 7, 8],
[1, 2, 3, 4]], dtype=uint32)
```

```
_array__ (<None>)
```

Returns a 0-dimensional array of the given data type, or of type(self) if argument is None.

```
_array_wrap_ (array)
```

Returns a scalar array object from the first-element of the array.

```
_squeeze_ ()
```

Returns self.

```
byteswap (<False>)
```

Trying to set the first (inplace) argument to True raises a ValueError. Otherwise, this returns a new array scalar with the data byteswapped.

```
_reduce_ ()
```

This is called to pickle an array scalar. It returns a tuple of (numpy.core.multiarray.scalar, self.dtypestr, obj or self.tostring()) which can be used to reconstruct the scalar on unpickling. Notice that no state is written, because the entire scalar can be constructed from just the string. Also, if this is an object array scalar, then the Python object being referenced is written.

```
_setstate_ ()
```

Does nothing but return Py\_None.

### setflags ()

Does nothing, as flags cannot be set for scalars.

### 6.3 Defining New Types

There are two ways to effectively define a new type of array. One way is to simply subclass the ndarray and overwrite the methods of interest. This will work to a degree, but internally certain behaviors are fixed by the data type of the array. To fully customize the data type of an array you need to define a new type that serves as the data type of the array, and register it with NumPy. This new type can only be defined in C. How to define a new data type in C will be discussed in the next part of the book.

# Chapter 7

# **Data-type Objects**

It is important to not confuse the "array-scalars" with the "data-type objects" even though an "array-scalar" can be interpreted as a data-type object and so can be used to refer to the data-type of an array. Every ndarray has an associated data-type object that completely defines the data in the array (including possible named fields). For every data-type object there is an associated type object whose instances are the array-scalars. Because of the association between each data-type object and a type-object of the corresponding array scalar, the array-scalar typeobjects can also be thought of as data-types. However, for the type objects of flexible array-scalars (string, unicode, and void), the type-objects alone are not enough to specify the full data-type because the length is not given. The data-type constructor, numpy.dtype, converts any object that can be considered as a datatype into a data-type object which is the actual object an ndarray looks to in order to interpret each element of its data region. Whenever a data-type is required in a NumPy function or method, supplying a dtype object is always fastest. If the object supplied is not a dtype object, then it will be converted to one using dtype (obj). Therefore, understanding data-type objects is the key to understanding how data types are really understood in NumPy.

### 7.1 Attributes

**type** The type object used to instantiate a scalar of this data-type.

kind A character code (one of 'biufcSUV') identifying the general kind of data.

char A unique character code for each of the 21 different built-in types.

- **num** A unique number for each of the 21 different built-in types roughly ordered from least-to-most precision.
- **str** The array-protocol typestring of this data-type object.
- **name** A bit-field-width name for this data-type (un-sized flexible data-type objects are missing the width).
- byteorder A character indicating the byte-order of this data-type object ('=' : native, '<' : little-endian, '>' : big-endian, '|' : not applicable). All builtin data-type objects have byteorder either '=' or '|'.
- itemsize The element size of this data-type object. For 18 of the 21 types this number is fixed by the data-type. For the flexible data-types, this number can be anything.
- **alignment** The required alignment (in bytes) of this data-type according to the compiler. More information is available in the C-API section.
- fields A dictionary showing any named fields that have been defined for this datatype (or None if there are no named fields). Fields can be assigned to any builtin data-type (e.g. using the tuple input to the dtypedescr constructor). However, fields are most useful for (subtypes of) void data-types which can be any size. Fields are a convenient way to keep track of fixed-size sub-parts of the total fixed-size array-element, or record. A field is defined in terms of another dtypedescr object and an offset (in bytes) into the current record.
  - The fields dictionary is indexed by keys that are the names of the fields. Each entry in the dictionary is a tuple fully describing the field: (dtypedescr, offset[, title]). If present, the optional title can be any string (or unicode), and it will also be a key in the fields dictionary (so that fields can be retrieved by title as well as by name). Notice also, that the first two elements of the tuple can be passed directly as arguments to the getfield and setfield attributes of an ndarray. If field names are not specified in a constructor, they default to 'f1', 'f2', ..., 'fn'.
  - An ordered (by offset) list of field names is also stored in the fields dictionary under the key -1. This can be used to walk through all of the named fields in offset order. Notice that the defined fields do not have to "cover" the record, but the itemsize of the container data-type object must always be at least as large as the itemsizes of the data-type objects in the defined fields.

subdtype Numarray introduced the concept of a fixed-length record having fields that were themselves arrays of another data-type. This is supported at a fundamental level in NumPy using this attribute which maintains the simplicity of defining a field by another data-type object. It either returns None or a tuple (base dtypedescr, shape) where shape is a tuple showing the size of the C-contiguous array and the base dtypedescr object indicates the data-type in each element of the subarray. If a field whose dtypedescr object has this attribute is retreived, then the extra dimensions implied by the shape are tacked on to the end of the retrieved array.

**descr** An array-interface-compliant full description of the data-type. The format is that required by descr \_array\_interface\_

**isbuiltin** A 1 if self is one of the builtin dtypedescr objects; a 2 if self is a user-defined data-type object; a 0, otherwise.

**isnative** True if this data-type object has a byteorder that is native to the platform; otherwise False.

hasobject A 1 if self is either an array of arbitrary objects ('O') or a data-type that includes fields or subdatatypes (at any depth) that are arbitrary objects ('O'). Recall that what is actually in the ndarray memory representing the Python object is the memory address of that object (a pointer). Special handling may be required and this attribute is useful for distinguishing data-types that may contain arbitrary Python objects and data-types that won't.

### 7.2 Construction

dtype (obj, align=0, copy=0)

Return a new data-type object from obj. The keyword argument, align, can only be nonzero if obj is a list, dictionary, or a comma-separated string. If it is nonzero in those cases it is used to add padding as needed to the fields to match what the compiler that compiled NumPy would do to a similar C-struct. The copy argument guarantees a new copy of the data-type object, otherwise, the result may just be a reference to a builtin data-type object.

Objects that can be converted to a data-type object are described in the following list. Because every object in this list can be converted to a data-type object it can also be used whenever a dtype is requested by a function or method in NumPy.

dtype Returns itself.

None Returns the default data-type descriptor object: currently int.

**type-object** Many Python type objects can be converted to data-type objects.

1. Array-scalar types: The type-objects of the 21 builtin array scalars all convert to an associated data-type object. This is true for subclasses as well. Not all data-type information can be supplied with a type-object. Flexible data-types with default itemsizes of 0, for example, require an itemsize to be useful.

Examples: int32, float64, uint16, complex128

- 2. Generic types: The generic hierarchial type objects convert to corresponding dtypedescr objects according to the associations: (numeric, inexact, floating) -> float; complexfloating -> cfloat; (integer, signedinteger) -> int\_; unsignedinteger -> uint; character -> string; (generic, flexible) -> void.
- 3. Builtin types: Several python types are equivalent to a corresponding array scalar when used to generate a dtypedescr object: int -> int\_; bool -> bool\_; float -> float\_; complex -> cfloat; str -> string; unicode -> unicode\_; buffer -> void; (all others) -> object\_.

Examples: object, str, float, int

4. Any type object with the dtype attribute: The attribute will be accessed and used directly. The attribute must return something that is convertible into a dtype object.

**string** Several kinds of strings can be converted. Recognized strings can be pre-pended with '>', or '<', to specify the byteorder.

1. One-character strings: Each builtin data-type has a character code (the updated Numeric typecodes), that uniquely identifies it.

**Examples:** 'b', 'H', 'f', 'd', 'F', 'D', Float64, Int32, UInt16

Array-protocol type strings: The first character specifies the kind of data and the remaining characters specify how many bytes of data.
 The supported kinds are 'b' -> boolean, 'i' -> (signed) integer, 'u' -> unsigned integer, 'f' -> floating-point, 'c' -> complex-floating point, 'S', 'a' -> string, 'U' -> unicode, 'V' -> anything (void).

Examples: 'i4', 'f8', 'c16', 'b1', 'S10', 'a25'

3. Comma-separated field formats: numarray introduced a short-hand notation for specifying the format of a record as a comma-separated

string of basic formats. A basic format in this context is an optional shape specifier followed by an array-protocol type string. Parenthesis are required on the shape if it is greater than 1-d. NumPy allows a modification on the format in that any string that can uniquely identify the type can be used. This data-type defines fields named 'f1', 'f2', ..., 'fN' where N (>1) is the number of comma-separated basic formats in the string. If the optional shape specifier is provided, then the data-type for the corresponding field contains a subdescr attribute providing the shape.

**Examples:** "i4, (2,3)f8, f4"; "a3, 3u8, (3,4)a10"

4. Any string in NumPy.typeDict.keys():

Examples: 'uint32', 'Int16', 'Uint64', 'Float64', 'Complex64'

**tuple** Three kinds of tuples each of length 2 can be converted into a data-type object:

(flexible dtype, itemsize): The first argument must be an object that
is converted to a flexible data-type object (one whose element size is
0), the second argument is an integer providing the desired itemsize.

**Examples:** (void, 10); (str, 35), ('U', 10)

2. (fixed dtype, shape): The first argument is any object that can be converted into a fixed-size data-type object. The second argument is the desired shape of this type. If the shape parameter is 1, then the data-type object is equivalent to fixed dtype.

**Examples:** (int32, (2,5)); ('S10', 1)=='S10'; ('i4, (2,3)f8, f4', (2,3))

3. (base dtype, new dtype): Both arguments must be convertible to data-type objects in this case. The base dtype is the data-type object that the new data-type builds on. This is how you could assign named fields to any builtin data-type object.

Examples: (int32, {'real':(int16,0), 'imag':(int16,2)}); (int32, (int8, 4)); ('i4', [('r','u1'),('g','u1'),('b','u1'),('a','u1')])

list Two styles of lists are accepted. These styles are unfortunately not uniquely separated. Ambiguity can occur if all of the field names correspond to recognizeable data-types. Therefore, you should recognize that the first format is tried first whenever the object is passed in to another routine using dtype=, but that the second format (array interface style) is tried first when calling the dtype constructor. In all cases, if one format fails the other is tried so the only real difficulty that could arise is

if you just happen to have all your field names corresponding to understandable types. Then you should create the data-type object explicitly using the dtype constructor.

1. [dtype1, dtype2, ..., dtypeN] where each element of the list is an object that can be converted to a data-type object. This constructs a data-type object with fields named 'f1', 'f2', ..., 'fN' with corresponding data-type objects dtype1, dtype2, ..., dtypeN, and automatically computed offsets (that are aligned if align is nonzero).

```
Examples: ['i4',('f8', (2,3)), 'S5']; [int32, (str, 10), 'u2, (2,3) u4, u1']
```

2. array interface style: This style is more fully described at this site http://numeric.scipy.org/array\_interface.html. It consists of a list of fields where each field is described by a tuple of length 2 or 3. The first element of the tuple is the field name (if this is "then a standard field name, 'fn', is assigned). The field name may also be a 2-tuple of strings where the first string is the "title" which may be any string or unicode string, and the second string is the "name" which must be a valid Python identifier. The second element of the tuple can be anything that can be interpreted as a data-type. The optional third element of the tuple contains the shape if this field represents an array of the data-type in the second element. This style does not accept align=1 as it is assumed that all of the memory is accounted for by the array\_descr protocol. See the web-page for more examples. Note that a 3-tuple with a third argument equal to 1 is equivalent to a 2-tuple.

dictionary There are also two dictionary styles. The first is a standard dictionary format while the second accepted format allows the fields attribute of dtypedescr objects to be interpreted as a data-type.

1. names and formats: This style has two required and two optional keys. The 'names' and 'formats' keys are required. Their respective values are equal-length lists with the field names and the field formats. The field names must be strings and the field formats can be any object accepted by dtypedescr constructor. The optional keys in the dictionary are 'offsets' and 'titles' and their values must each be lists of the same length as the 'names' and 'formats' lists. The

'offsets' value is a list of integer offsets for each field, while the 'titles' value is a list of titles for each field (None can be used if no title is desired for that field). The titles can be any string or unicode object and will add another entry to the fields dictionary keyed by the title and referencing the same field tuple which will contain the title as an additional tuple member.

```
Examples: {'names': ['r','g','b','a'], 'formats': [uint8, uint8, uint8]}; {'names':['r','b'], 'formats': ['u1', 'u1'], 'offsets': [0, 2], 'titles': ['Red pixel', 'Blue pixel']}
```

2. data-type object fields: This style is patterned after the format of the fields dictionary in a data-type object. It contains string or unicode keys that refer to (data-type, offset) or (data-type, offset, title) tuples. There is one special key (-1), present in data-type object fields, that refers to a list of the field names (not titles). It does not have to be present, however in order for the dtypedescr constructor to recognize the data-type.

**Examples:** {'col1': ('S10', 0), 'col2': (float32, 10), 'col3': (int, 14)}

### 7.3 Methods

```
newbyteorder (<'swap'>)
```

Construct a new copy of self with its byteorder changed according to the optional argument. All changes are also propagated to the data-type objects of all fields and sub-arrays. If a byteorder of '|' (meaning ignore) is encountered it is left unchanged. The default behavior is to swap the byteorder. Other possible arguments are 'big' ('>'), 'little' ('<'), and 'native' ('=') which recursively forces the byteorder of self (and it's field data-type objects and any sub-arrays) to the corresponding byteorder.

```
_reduce_ ()
_setstate_ (state)
```

Data-type objects can be pickled because of these two methods. The <u>reduce</u>() method returns a 3-tuple consisting of (callable object, args, state), where the callable object is numpy.core.multiarray.dtypedescr and args is (typestring, 0,

- 1) unless the data-type inherits from void in which case args is (typeobj, 0,
- 1). The state is a 5-tuple with (endian, self.subdescr, self.fields, self.itemsize, self.alignment). The last two entries are both -1 if the data-type object is

builtin and not flexible (because they are fixed on creation). The set state method takes the saved state and updates the date-type descr object.

# Chapter 8

## Standard Classes

The ndarray in NumPy is a "new-style" Python builtin-type. Therefore, it can be inherited from (in Python or in C) if desired. Therfore, it can form a foundation for many useful classes. Often whether to sub-class the array object or to simply use the core array component as an internal part of a new class is a difficult decision, and can be simply a matter of choice. NumPy has several tools for simplifying how your new object interacts with other array objects, and so the choice may not be significant in the end. One way to simplify the question is by asking yourself if the object you are interested can be replaced as a single array or does it really require two or more arrays at it's core. For example, in the standard NumPy distribution, the matrix and records classes inherit from the ndarray, while masked arrays use two ndarrays as objects of its internal structure.

Note that asarray(a) always returns the base-class ndarray. If you are confident that your use of the array object can handle any subclass of an ndarray, then asanyarray(a) can be used to allow subclasses to propagate more cleanly through your subroutine. In principal a subclass could redefine any aspect of the array and therefore, under strict guidelines, asanyarray(a) would rarely be useful. However, most subclasses of the arrayobject will not redefine certain aspects of the array object such as the buffer interface, or the attributes of the array. One of important example, however, of why your subroutine may not be able to handle an arbitrary subclass of an array is that matrices redefine the '\*' operator to be matrix-multiplication, rather than element-by-element multiplication.

# 8.1 Special attributes and methods recognized by NumPy

### \_array\_finalize\_ (obj)

This method is called whenever the system internally allocates a new array from obj, where obj is a subclass (subtype) of the (big)ndarray. It can be used to change attributes of self after construction (so as to ensure a 2-d matrix for example), or to update meta-information from the "parent." Subclasses inherit a default implementation of this method that does nothing.

### **\_array\_wrap\_** (array)

This method should return an instance of the class from the ndarray object passed in. For example, this is called after every ufunc for the object with the highest \_array\_priority\_. The ufunc-computed array object is passed in and whatever is returned is passed to the user. Subclasses inherit a default implementation of this method.

```
_array__ (dtype <None>)
```

This method is called to obtain an ndarray object when needed. You should always guarantee this returns an actual ndarray object. Subclasses inherit a default implementation of this method.

### \_array\_priority\_\_

The value of this attribute is used to determine what type of object to return in situations where there is more than one possibility for the Python type of the returned object. Subclasses inherit a default value of 1.0 for this attribute.

### 8.2 Matrix Objects

Matrix objects inherit from the ndarray and therefore, they have the same attributes and methods of ndarrays. There are six important differences of matrix objects, however that may lead to unexpected results when you use matrices but expect them to act like arrays:

1. Matrix objects can be created using a string notation to allow Matlab-style syntax where spaces separate columns and semicolons (';') separate rows.

- 2. Matrix objects are always two-dimensional. This has far-reaching implications, in that m.ravel() is still two-dimensional (with a 1 in the first dimension) and item selection returns two-dimensional objects so that sequence behavior is fundamentally different than arrays.
- 3. Matrix objects over-ride multiplication to be matrix-multiplication. Make sure you understand this for functions that you may want to receive matrices. Especially in light of the fact that asanyarray(m) returns a matrix when m is a matrix.
- 4. Matrix objects over-ride power to be matrix raised to a power. The same warning about using power inside a function that uses asanyarray(...) to get an array object holds for this fact.
- 5. The default \_array\_prioirty\_ of matrix objects is 10.0, and therefore mixed operations with ndarrays always produce matrices.
- 6. Matrices have special attributes which make calculations easier. These are
  - (a) .T return the transpose of self
  - (b) .H return the conjugate transpose of self
  - (c) .I return the inverse of self
  - (d) .A return a view of the data of self as a 2d array (no copy is done).

The matrix class is a Python subclass of the ndarray and can be used as a reference for how to construct your own subclass of the ndarray. Matrices can be created from other matrices, strings, and anything else that can be converted to an ndarray. The name "mat" is an alias for "matrix" in NumPy.

#### **Example 1:** Matrix creation from a string

```
>>> a=mat('1 2 3; 4 5 3')
>>> print (a*a.T).I
[[ 0.2924 -0.1345]
[-0.1345 0.0819]]
```

### **Example 2:** Matrix creation from nested sequence

```
>>> mat([[1,5,10],[1.0,3,4j]])
matrix([[ 1.+0.j, 5.+0.j, 10.+0.j],
[ 1.+0.j, 3.+0.j, 0.+4.j]])
```

### **Example 3:** Matrix creation from an array

```
>>> mat(rand(3,3)).T
matrix([[ 0.7224,  0.8579,  0.1112],
[ 0.2797,  0.6152,  0.2113],
[ 0.2002,  0.4644,  0.4951]])
```

```
matrix (data, dtype=None, copy=True)
```

The sequence to convert to a matrix is passed in as data. If dtype is None, then the data-type is determined from the data. If copy is True, then a copy of the data is made, otherwise, the same data buffer is used. If no buffer can be found for data, then a copy is also made. Note: The matrix object is actually a class and so using this syntax calls matrix.\_new\_(matrix, data, dtype, copy) which is what happens whenever you "call" any class object as a function.

#### mat

Just another name for matrix.

```
asmatrix (data, dtype=None)
```

Returns the data without copying. Equivalent to matrix(data, dtype, copy=False).

```
bmat (obj, ldict=None, gdict=None)
```

Build a matrix object from a string, nested sequence or an array. This command lets you build up matrices from other other objects. The ldict and gdict parameters are local and module (global) dictionaries that are only used when obj is a string. If they are not provided, then the local and module dictionaries present when bmat is called are used.

```
>>> A = mat('2 2; 2 2'); B=mat('1 1; 1 1');

>>> print bmat('A B; B A')

[[2 2 1 1]

[2 2 1 1]

[1 1 2 2]

[1 1 2 2]]
```

### 8.3 Memory-mapped-file arrays

Memory-mapped files are useful for reading and/or modifying small segments of a large file with regular layout, without reading the entire file into memory. A simple subclass of the ndarray uses a memory-mapped file for the data buffer of the array. For small files, the over-head of reading the entire file into memory is typically not significant, however for large files using memory mapping can save considerable resources.



### NOTE

Memory-mapped arrays use the Python memory-map object which (prior to Python 2.5) does not allow files to be larger than a certain size depending on the platform. This size is always  $< 2 \mathrm{GB}$  even on 64-bit systems.

The class is called memmap and is available in the NumPy namespace. The \_new\_ method of the class has been re-written to have the following syntax:

\_new\_ (cls, filename, dtype=uint8, mode='r+', offset=0, shape=None, order=0)

filename The file name to be used as the array data buffer

**dtype** A data-type object used to interpret the file contents (including byte-order).

mode The mode to open the file in. Valid modes are 'readonly' or 'r', 'copyonwrite' or 'c', 'readwrite' or 'r+', and 'write' or 'w+'. This mode determines the WRITEABLE flag of the returned array.

offset An offset into the file to start the array data.

**shape** The desired shape of the array. If this is None, then the returned array will be 1-d with the number of elements determined by the file size and data type.

**order** Either 'C' or 'Fortran' to indicate the order that an N-D array should be interpreted. This only has an effect if the shape is greater than 2-D.

Memory-mapped-file arrays have one additional method (besides those they inherit from the ndarray): self.sync() which must be called manually by the user to ensure that any changes to the array actually get written to disk.

#### Example:

```
>>> a = memmap('newfile.dat', dtype=float, mode='w+', shape=1000)
>>> a[10] = 10.0
>>> a[30] = 30.0
>>> del a
>>> b = fromfile('newfile.dat', dtype=float)
>>> print b[10], b[30]
10.0 30.0
>>> a = memmap('newfile.dat', dtype=float)
>>> print a[10], a[30]
10.0 30.0
```

### 8.4 Character arrays (numpy.char)

These are enhanced arrays of either string type or unicode\_ type. These arrays inherit from the ndarray, but specially-define the operations ==, <=, >=, !=, >, <, +, \*, and % on a (broadcasting) element-by-element basis. Because ufuncs do not support flexible-size arrays, these operations are not (yet) available on the basic ndarray of character type. In addition, the chararray has all of the standard string (and unicode) methods, excecuting them on an element-by-element basis. Perhaps the easiest way to creat a chararray is to use self.view(chararray) where self is an ndarray of string or unicode data-type. However, a chararray can also be created using the numpy.chararray.—new— method.

**\_\_new\_\_** (shape, itemsize, unicode=False, buffer=None, offset=0, strides=None, order=None)

Create a new character array of string or unicode type and itemsize characters. Create the array using buffer (with offset and strides) if it is not None. If buffer is None, then construct a new array with strides in Fortran order if len(shape) >=2 and order is 'Fortran' (otherwise the strides will be in 'C' order).

char.array (obj. itemsize=None, copy=True, unicode=False, order=None)

Create a chararray from the nested sequence obj. If obj is an ndarray of datatype unicode\_ or string, then its data is wrapped by the chararray object and converted to the desired type (string or unicode).

### 8.5 Record Arrays (numpy.rec)

NumPy provides a powerful data-type object that allows any ndarray to hold (arbitrarily nested) record-like items with named-field access to the sub-types. This is possible without any special record-array sub-class. Consider the example where each item in the array is a simple record of name, age, and weight. You could specify a data-type for an array of such records using the following data-type object:

```
>>> desc = dtype({'names': ['name', 'age', 'weight'], 'formats': ['S30', 'i2', '
>>> a = array([('Bill',31,260.0),('Fred', 15, 145.0)],dtype=desc)
>>> print a[0]
('Bill', 31, 260.0)
>>> print a['name']
[Bill Fred]
>>> print a['age']
[31 15]
>>> print a['weight']
[ 260. 145.]
>>> print a[0]['name'], a[0]['age'], a[0]['weight']
Bill 31 260.0
>>> print len(a[0])
3
```

This example shows how a general array can be assigned named fields and how these fields can be accessed. In this case the a[0] object is an array-scalar of type void. The void array-scalars are unique in that they contain references to (rather than copies of) the underlying data whenever fields are defined. Therefore, the record data can be modified in place:

```
>>> a[0]['name'] = 'George'; print a
[('George', 31, 260.0) ('Fred', 15, 145.0)]
```

The recarray subclass and its accompanying record item add the ability to access named fields through attribute lookup. A quick way to get a record array is to use the view method of the ndarray.

```
>>> r = a.view(recarray)
>>> print r.name
[George Fred]
```

The numpy.core.records module (aliased to nump.rec when numpy is imported) contains additional convenience functions for constructing record arrays.

**fromarrays** (array\_list, formats=None, names=None, titles=None, shape=None, aligned=0)

Create a record array from a (flat) list of ndarrays. The data from the arrays will be copied into the fields. If formats is None, then formats are determined from the arrays. The names and titles arguments can be a list, tuple or a (comma-separated) string specifying the names and/or titles to use for the fields. If aligned is True, then the structure will be padded according to the rules of the compiler that NumPy was compiled with.

```
>>> x1 = array([21,32,14])
>>> x2 = array(['my','first','name'])
>>> x3 = array([3.1, 4.5, 6.2])
>>> r = rec.fromarrays([x1,x2,x3], names='id, word, number')
>>> print r[1]
(32, 'first', 4.5)
>>> r.number
array([ 3.1,  4.5,  6.2])
>>> r.word
chararray([my, first, name],
dtype='|S5')
```

**fromrecords** (rec\_list, formats=None, names=None, titles=None, shape=None, aligned=0)

Construct a record array from a (nested) sequence of tuples that define the records. If formats are not given, they are deduced from the records, but this is slower. The field names and field titles can be specified. If aligned is non-zero, then the record array is padded so that fields are aligned as the platform compiler would do if the fields represented a C-struct.

```
>>> recs = [('Bill', 31, 260.0), ('Fred', 15, 145.0)]
>>> r = rec.fromrecords(recs, formats='S30,i2,f4', names='name, age, weight')
>>> print r.name
[Bill Fred]
>>> print r.age
[31 15]
>>> print r.weight
[ 260. 145.]
```

**fromstring** (datastring, formats, shape=None, names=None, titles=None, byte-order=None, aligned=0, offset=0):

Construct a record array using the provided datastring (at the given offset) as the memory. The record array will be read-only. The byteorder argument may be used to specify the byteorder of all of the fields at the same time. A True aligned argument causes padding fields to be added as needed so that the fields are aligned on boundaries determined by the compiler. The shape of the returned array can also be specified.

**fromfile** (fd, formats, shape=None, names=None, titles=None, byteorder=None, aligned=0, offset=0)

Construct a record array from the (binary) data in the given file object, fd. This object may be an open file or a string to indicate a file to read from. If offset is non-zero, then data is read from the file at offset bytes from the current position.

array (obj, formats=None, names=None, titles=None, shape=None, byteorder=None, aligned=0, offset=0, strides=None)

A general-purpose record array constructor that is a front-end to recarray, fromstring, fromarrays, fromrecords, and fromfile. If obj is None, then call the recarray constructor. If obj is a string, then call the fromstring constructor. If obj is a list or a tuple then if the first object is an ndarray, the call fromarrays, otherwise call fromrecords. If obj is a recarray, then make a copy of the data in recarray and use the new formats, names, and titles. If obj is a file then call fromfile. Finally, if obj is an ndarray, then return obj.view(recarray). The formats argument must be given if obj is None, a string, or a file, and if obj is None so the recarray constructor will be called, then shape must be given as well. The following classes are also available in the numpy.core (and therefore the numpy) namespace

**record** A subclass of the void array scalar type that allows field access using attributes.

recarray A subclass of the ndarray that allows field access using attributes

**\_\_new\_\_** (subtype, shape, formats, names=None, titles=None, buf=None, offset=0, strides=None, byteorder=None, aligned=0)

Construct an array of the given subtype and shape with data-type (record, dtype) where dtype is constructed from formats, names, and titles. If buf is None, then create new memory. Otherwise, use the memory of buf exposed through the buffer protocol.

**format\_parser** A class useful for creating a data-type descriptor from formats, names, titles, and aligned arguments. This is used by several of the record array constructors for consistency in behavior.

**\_\_init\_\_** (self, formats, names, titles, aligned=False)

Construct a data-type object from formats, names, titles, and aligned argument. Upon completion the constructed data-type object is in self.\_descr.

### 8.6 Masked Arrays (numpy.ma)

These are adapted from the masked arrays provided with Numeric. Masked Arrays do not inherit from the ndarray, they simply use two ndarray objects in their internal representation. Fortunately, as I have not used masked arrays in my work, Paul Dubois (the original author of MA for Numeric) adapted and modified the code for use by NumPv.

Masked arrays are created using the masked array creation function:

### 8.7 UserArray

For backward compatibility and as a standard "container" class, the UserArray from Numeric has been brought over to NumPy. The UserArray is a Python class whose self-array attribute is an ndarray. Multiple inheritance might be easier with UserArray than with the ndarray and so it is included by default. It is not documented here beyond mentioning its existence because you are encouraged to use the ndarray class directly if you can.

### 8.8 Array Iterators

Iterators are a powerful concept for array processing. Essentially, iterators implement a generalized for-loop. If myiter is an iterator object, then the Python code

```
for val in myiter:
...
some code involving val
```

calls val=myiter.next() repeatedly until StopIteration is raised by the iterator. There are several ways to iterate over an array that may be useful: default iteration, flat iteration, and N-dimensional enumeration.

### 8.8.1 Default iteration

The default iterator of an ndarray object is the default Python iterator of a sequence type. Thus, when the array object itself is used as an iterator. The default behavior is equivalent to:

```
for i in arr.shape[0]:
    val = arr[i]
```

This default iterator selects a sub-array of dimension N-1 from the array. This can be a useful construct for defining recursive algorithms. To loop over the entire array requires N for-loops.

```
>>> a = arange(24).reshape(3,2,4)+10
>>> for val in a:
... print 'item:', val
item: [[10 11 12 13]
[14 15 16 17]]
item: [[18 19 20 21]
[22 23 24 25]]
item: [[26 27 28 29]
[30 31 32 33]]
```

### 8.8.2 Flat iteration

As mentioned previously, the flat attribute of ndarray objects returns an iterator that will cycle over the entire array in C-style contiguous order.

```
>>> for i, val in enumerate(a.flat):
... if i%5 == 0: print i, val
0 10
5 15
10 20
15 25
20 30
```

Here, I've used the builtin enumerate iterator to return the iterator index as well as the value.

### 8.8.3 N-dimensional enumeration

Sometimes it may be useful to get the N-dimensional index while iterating. The ndenumerate iterator can achieve this.

```
>>> for i, val in ndenumerate(a):
... if sum(i)%5 == 0: print i, val
(0, 0, 0) 10
(1, 1, 3) 25
(2, 0, 3) 29
(2, 1, 2) 32
```

### 8.8.4 Iterator for broadcasting

The general concept of broadcasting is also available from Python using the **broadcast** iterator. This object takes N objects as inputs and returns an iterator that returns tuples providing each of the input sequence elements in the broadcasted result.

```
>>> for val in broadcast([[1,0],[2,3]],[0,1]):
... print val
(1, 0)
(0, 1)
(2, 0)
(3, 1)
```

The methods and attributes of the broadcast object are:

nd the number of dimensions in the broadcasted result.

**shape** the shape of the broadcasted result.

**size** the total size of the broadcasted result.

index the current (flat) index into the broadcasted array

iters a tuple of (broadcasted) NumPy.flatiter objects, one for each array.

reset ()

Reset the multiter object to the beginning.

next ()

Get the next tuple of objects from the (broadcasted) arrays

# Chapter 9

# **Universal Functions**

Computers make it easier to do a lot of things, but most of the things they make it easier to do don't need to be done.

—Andy Rooney

People think computers will keep them from making mistakes. They're wrong. With computers you make mistakes faster.

—Adam Osborne

### 9.1 Description

Universal functions are wrappers that provide a common interface to mathematical functions that operate on scalars, and can be made to operate on arrays in an element-by-element fashion. All universal functions (ufuncs) wrap some core function that takes  $n_i$  (scalar) inputs and produces  $n_o$  (scalar) outputs. Typically, this core function is implemented in compiled code but a Python function can also be wrapped into a universal function using the basic method frompyfunc in the umath module.

### frompyfunc (func, nin, nout)

This function returns a new universal function wrapping a Python function func with nin inputs and nout outputs. The resulting universal function works using Object arrays for both input and output. The vectorize class makes use of frompyfunc internally. You can view the source code in numpy/core/function\_base.py.

### 9.1.1 Broadcasting

Each universal function takes array inputs and produces array outputs by performing the core function element-wise on the inputs. The standard broadcasting rules are applied so that inputs without exactly the same shapes can still be usefully operated on. Broadcasting can be understood by four rules:

- 1. All input arrays with ndim smaller than the input array of largest ndim have 1's pre-pended to their shapes.
- 2. The size in each dimension of the output shape is the maximum of all the input shapes in that dimension.
- 3. An input can be used in the calculation if it's shape in a particular dimension either matches the output shape or has value exactly 1.
- 4. If an input has a dimension size of 1 in its shape, the first data entry in that dimension will be used for all calculations along that dimesion. In other words, the stepping machinery of the ufunc will simply not step along that dimension when otherwise needed (the stride will be 0 for that dimension).

While perhaps somewhat difficult to explain, broadcasting can be quite useful and becomes second nature rather quickly.

### 9.1.2 Output type determination

The output of the ufunc (and its methods) does not have to be an ndarray. All output arrays will be passed to the \_array\_wrap\_ method of any input (besides ndarrays, and scalars) that defines it and has the highest \_array\_priority\_ of any other input to the universal function. The default \_array\_priority\_ of the ndarray is 0.0, and the default \_array\_priority\_ of a subtype is 1.0. Matrices have \_array\_priority\_ equal to 10.0.

The ufuncs can also all take output arguments. The output will be cast if necessary to the provided output array. If a class with an \_array\_ method is used for the output, results will be written to the object returned by \_array\_. Then, if the class also has an \_array\_wrap\_ method, the returned ndarray result will be passed to that method just before passing control back to the caller.

### 9.1.3 Use of internal buffers

Internally, buffers are used for misaligned data, swapped data, and data that has to be converted from one data type to another. The size of the internal buffers is settable on a per-thread basis. There can be up to  $2(n_i + n_o)$  buffers of the specified

size created to handle the data from all the inputs and outputs of a ufunc. The default size of the buffer is 10,000 elements. Whenever buffer-based calculation would be needed, but all input arrays are smaller than the buffer size, those misbehaved or incorrect typed arrays will be copied before the calculation proceeds. Adjusting the size of the buffer may therefore alter the speed at which ufunc calculations of various sorts are completed. A simple interface for setting this variable is accessible using the function

### setbufsize (size)

Set the buffer size to the given number of elements in the current thread. Return the old buffer size (so that it can be reset later if desired).

### 9.1.4 Error handling

Universal functions can trip special floating point status registers (such as divideby-zero) which will be regularly checked during calculation. The user can determine what should be done if such errors are encountered. Error handling is controlled on a per-thread basis. Four errors can be individually configured: divide-by-zero, overflow, underflow, and invalid. The errors can each be set to ignore, warn, raise, or call. The easiest way to configure the error mask is using the function

seterr (divide=None, over=None, under=None, invalid=None)

This will set the current thread so that errors can be handled if desired. If one of the errors is set to 'call', then a function must be provided using the seterrcall() routine. If any of the arguments are None, then that error mask will be unchanged. The return value of this function is a dictionary with the old error conditions. Thus, you can restore the old condition after you are finished with your function by calling seterr(\*\*old).

### seterrcall (callable)

This sets the function to call when an error is triggered for an error condition configured with the "call" handler. This function should take two arguments: a string showing the type of error that triggered the call, and an integer showing the state of the floating point status registers. Any return value of the call function will be ignored, but errors can be raised by the function. Only one error function handler can be specified for all the errors. The status argument shows which errors were raised. The return value of this routine is the old callable. The argument passed in to this function must be any callable object with the right signature or None.



### NOTE

FPE\_DIVIDEBYZERO, FPE\_OVERFLOW, FPE\_UNDERFLOW, and FPE\_INVALID, are all defined constants in NumPy. The status flag returned for a 'call' error handling type shows which errors were raised by adding these constants together.

### 9.2 ufunc attributes

There are some informational attributes that universal functions possess. None of the attributes can be set.

#### $\_doc\_$

A docstring for each ufunc. The first part of the docstring is dynamically generated from the number of outputs, the name, and the number of inputs. The second part of the doc string is provided at creation time and stored with the ufunc.

#### \_name\_

The name of this ufunc.

#### nin

The number of inputs

#### nout

The number of outputs

#### nargs

The total number of inputs + outputs

### ntypes

The total number of different types for which this usunc is defined.

#### types

A list of length ntypes containing strings showing the types for which this ufunc is defined. Other types may still be used as inputs (and as output arrays), they will just need casting. For inputs, standard casting rules will be used

Name	Description
_doc_	Dynamic docstring.
_name	Name of ufunc
nin	Number of input arguments
nout	Number of output arguments
nargs	Total number of arguments
ntypes	Number of defined inner loops.
types	List showing types for which inner loop is defined.
identity	Identity for this ufunc.

Table 9.1: Universal function (ufunc) attributes.

to determine which of the supplied internal functions that will be used (and therefore the default type of the output). Results will always be force-cast to any array provided to hold the output.

### identity

A 1, 0, or None to show the identity for this universal function. This identity is used for reduction on zero-sized arrays (arrays with a shape that includes a 0).

### 9.3 Casting Rules

At the core of every ufunc is a one-dimensional strided loop that implements the actual function for a specific type combination. When a ufunc is created, it is given a static list of inner loops and a corresponding list of type signatures over which the ufunc operates. The ufunc machinery uses this list to determine which inner loop to use for a particular case. You can inspect the .types attribute for a particular ufunc to see which type combinations have a defined inner loop and which output type they produce (the character codes are used in that output for brevity).

Casting must be done on one or more of the inputs whenever the ufunc does not have a core loop implementation for the input types provided. If an implementation for the input types cannot be found, then the algorithm searches for an implementation with a type signature to which all of the inputs can be cast "safely." The first one it finds in its internal list of loops is selected and performed with types cast. Recall that internal copies during ufuncs (even for casting) are limited to the size of an internal buffer which is user settable.



### NOTE

Universal functions in NumPy are flexible enough to have mixed type signatures. Thus, for example, a universal function could be defined that works with floating point and integer values. See Idexp for an example.

By the above description, the casting rules are essentially implemented by the question of when a data type can be cast "safely" to another data type. The answer to this question can be determined in Python with a function call: can\_cast (fromtype, totype). Figure shows the results of this call for my 32-bit system on the 21 internally supported types. You can generate this table for your system with code shown in that Figure.

You should note that, while included in the table for completeness, the 'S', 'U', and 'V' types cannot be operated on by ufuncs. In addition, there is currently no mechanism for user-defined types to participate in this automatic casting processes. Also, note that on a 64-bit system the integer types may have different sizes resulting in a slightly altered table.

Mixed scalar-array operations use a different set of casting rules that ensure that a scalar cannot upcast an array unless the scalar is of a fundamentally different kind of data (*i.e.* under a different hierarchy in the data type hierarchy) then the array. This rule enables you to use scalar constants in your code (which as Python types are interpreted accordingly in ufuncs) without worrying about whether the precision of the scalar constant will cause upcasting on your large (small precision) array.

### 9.4 ufunc Object methods

All ufuncs have 4 methods. However, these methods only make sense on ufuncs that take two input arguments and return one output argument. Attempting to call these methods on other ufuncs will cause a ValueError. The reduce-like methods all take an axis keyword and an dtype keyword, and the arrays must all have dimension >= 1. The axis keyword specifies which axis of the array the reduction will take place over and may be negative, but must be an integer. The dtype keyword allows you to manage a very common problem that arises when naively using <op>.reduce. Sometimes you may have an array of a certain data type and wish to add up all of its elements, but the result does not fit into the data type of the array. This commonly happens if you have an array of single-byte integers. The dtype keyword allows you to alter the data type that the reduction takes place over (and therefore the type of the output). Thus, you can ensure that the output is a data type with

```
>>> def print_table(ntypes):
    print 'X',
    for char in ntypes: print char,
    print
    for row in ntypes:
      print row,
       for col in ntypes:
         print int(can_cast(row, col)),
       print
>>> print_table(typecodes['All'])
X?bhilqpBHILQPfdgFDGSUVO
i 0 0 0 1 1 1 1 0 0 0 0 1 0 0 1 1 0 1 1 1 1 1 1
1 0 0 0 1 1 1 1 0 0 0 0 1 0 0 1 1 0 1 1 1 1 1 1
q 0 0 0 0 0 1 0 0 0 0 0 0 0 1 1 0 1 1 1 1 1 1
p 0 0 0 1 1 1 1 0 0 0 0 1 0 0 1 1 0 1 1 1 1 1 1
I 0 0 0 0 0 1 0 0 0 1 1 1 1 0 1 1 0 1 1 1 1 1 1
L 0 0 0 0 0 1 0 0 0 1 1 1 1 0 1 1 0 1 1 1 1 1 1
Q 0 0 0 0 0 0 0 0 0 0 0 1 0 0 1 1 0 1 1 1 1 1 1
P 0 0 0 0 0 1 0 0 0 1 1 1 1 1 0 1 1 0 1 1 1 1 1 1
d 0 0 0 0 0 0 0 0 0 0 0 0 0 1 1 0 1 1 1 1 1 1
g 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 1 0 0 1 1 1 1 1 1
D 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 1 1 1 1 1 1 1
G O O O O O O O O O O O O O O O O O 1 1 1 1 1 1
```

Figure 9.1: Code segment showing the can cast safely table for a 32-bit system.

large-enough precision to handle your output. The responsibility of altering the reduce type is mostly up to you. If no dtype is given, then if the input type is an integer type and smaller than size of the int\_ data type, it will be internally upcast to the int\_ (or uint) data type. Boolean arrays will use an dtype of int\_ for "add" and "multiply" operations only. For other array types, the reduction takes place in the array type.



### WARNING

A reduce-like operation on an array with a data type that has range "too small" to handle the result will silently wrap. You should use dtype to increase the data type over which reduction takes place.

### **9.4.1** Reduce

```
<op>.reduce (array=, axis=0, dtype=None)
```

For each one-dimensional sequence along the axis dimension of the array, return a single number resulting from recursively applying the operation to succesive elements along that dimension. If the input array has N dimensions, then the returned array has N-1 dimensions. This produces the equivalent of the following Python code:

```
>>> indx = [index_exp[:]]*array.ndim
>>> indx[axis] = 0; N=array.shape[axis]
>>> result = array[indx].astype(dtype)
>>> for i in range(1,N):
... indx[axis] = i
... <op>(result, array[indx], result)
```

Studying the above code can also help you gain an appreciation for how to do generic indexing in Python using index\_exp. For example, if <op> is add, then <op>.reduce produces a summation along the given axis. If <op> is prod, then a repeated multiply is performed.

### 9.4.2 Accumulate

```
<op>.accumulate (array=, axis=0, dtype=None)
```

This method is similar to reduce, except it returns an array of the same shape as the input, and keeps intermediate calculations. The operation is still performed along the access. This method underlies the operations of the cumsum and cumprod methods of arrays. The following Python code implements an equivalent of the accumulate method.

```
>>> i1 = [index_exp[:]]*array.ndim
>>> i2 = [index_exp[:]]*array.ndim
>>> i1[axis] = 0; N=array.shape[axis]
>>> result = array.astype(dtype)
>>> for i in range(1,N):
... i1[axis] = i
... i2[axis] = i-1
... <op>(result[i1], array[i1], result[i2])
```

### 9.4.3 Reduceat

```
<op>.reduceat (array=, indices=, axis=0, dtype=None)
```

This method is a generalization of both reduce and accumulate. It offers the ability to reduce along an axis but only between certain indices. The indices input must be a one dimensional (index) sequence. Then, if  $I_k$  is the  $k^{\text{th}}$  element of indices, the reduceat method computes <op>.reduce(array[ $I_k:I_{k+1}$ ]). This formula assumes  $I_{k+1} > I_k$ , and also that  $I_{k+1}$  is the length of the input array when  $I_k$  is the laste element. There is no requirement that the indices be monotonic. If  $I_{k+1} < I_k$ , then reduceat simply returns array[ $I_k$ ] for that particular element of indices. In these formulas, we have assumed that array is one dimensional (or axis is 0). If the array is N-dimensional and axis>0, then the index expression needs axis ':' (full slice objects) inserted (i.e. array[:,...,:, $I_k:I_{k+1}$ ]). The effect is to slice along the axis dimension.

Equivalent Python code is

```
>>> i1 = [index_exp[:]]*array.ndim
>>> i2 = [index_exp[:]]*array.ndim
>>> outshape = list(array.shape)
>>> N = array.shape[axis]
>>> outshape[axis] = len(indices)
>>> result = zeros(outshape, dtype or array.dtype)
>>> for k,Ik in enumerate(indices):
... i1[axis] = k
... try:
... Ikp1 = indices[k+1]
... except IndexError:
```

The returned array has as many dimensions as the input array, and is the same shape except for the *axis* dimension which has shape equal to the length of indices (the number of reduce operations that were performed). If you ever have a need to compute multiple reductions over portions of an array, then (if you can get your mind around what it is doing) reduceat may be just what you were looking for.

```
Example: Suppose a is a two-dimensional array of shape 10 \times 20. Then, res=add.reduce (a, [0,3,1]) returns a 3 \times 20 array with res[0,:] = add.reduce(a[:,0:3]), res[1,:] = a[:,3], and res[2,:] = add.reduce(a[:,1:]).
```

### 9.4.4 Outer

```
<op>.outer (a, b)
```

This method computes an outer operation on <op>. It computes <op>(a2, b2) where a2 is 'a' with b.ndim 1's post-pended to it's shape and b2 is 'b' with a.ndim 1's pre-pended to its shape (broadcasting takes care of this automatically in the code below). The return shape has a.ndim + b.ndim dimensions. Equivalent Python code is

```
>>> a.shape += (1,)*b.ndim
>>> <op>(a,b)
>>> a = a.squeeze()
```

Arithmetic tables can be conveniently built using outer:

```
>>> multiply.outer([1,7,9,12],arange(5,12))
array([[ 5,  6,  7,  8,  9,  10,  11],
  [ 35,  42,  49,  56,  63,  70,  77],
  [ 45,  54,  63,  72,  81,  90,  99],
  [ 60,  72,  84,  96,  108,  120,  132]])
```

### 9.5 Available ufuncs

There are currently more than 60 universal functions defined on one or more types, covering a wide variety of operations. Some of these usuncs are called automatically on arrays when the relevant infix notation is used (i.e. add(a,b) is called internally when a + b is written and a or b is an ndarray). Nonetheless, you may still want to use the usunce call in order to use the optional output argument(s) to place the output(s) in an object (or in objects) of your choice.

Recall the each each usunce operates element-by-element. Therefore, each usunce will be described as if acting on a set of scalar inputs to return a set of scalar outputs.



### NOTE

The ufunc still returns its output(s) even If you use the optional output argument(s).

### 9.5.1 Math operations

**add**  $(x_1, x_2 [, y])$ 

 $y = x_1 + x_2$ . Called to implement x1+x2 for arrays

subtract  $(x_1, x_2 [, y])$ 

 $y = x_1 - x_2$ . Called to implement x1-x2 for arrays

multiply  $(x_1, x_2 [, y])$ 

 $y = x_1 \cdot x_2$ . Called to implement x1\*x2 for arrrays.

**divide**  $(x_1, x_2 [, y])$ 

 $y = x_1/x_2$  Integer division results in truncation. Floating-point does not. Called to implement x1/x2 for arrays (when \_future\_\_.division is not active).

true\_divide  $(x_1, x_2 [, y])$ 

This version of division always returns an inexact number so that integer division returns floating point. Called with \_\_future\_\_.division is active to implement x1/x2 for arrays.

floor\_divide  $(x_1, x_2 [, y])$ 

This version of division always results in truncation of an fractional part remaining. Called to implement x1//x2 for arrays.

**negative** (x [, y])

y = -x. Called to implement -x for arrays.

**power**  $(x_1, x_2 [, y])$ 

 $y = x_1^{x_2}$ . There is no three-term power ufunc defined. This two-term power function is called to implement pow(x1,x2,<any>) or x1\*\*x2 for arrays. Note that the third term in pow is ignored for array arguments.

remainder  $(x_1, x_2 [, y])$ 

Returns  $x - y^* \text{floor}(x/y)$ . Result has the sign of y. Called to implement x1\%x2.

 $mod (x_1, x_2 [, y])$ 

Same as remainder  $(x_1, x_2 [, y])$ .

**fmod**  $(x_1, x_2 [, y])$ 

 $x_1 = kx_2 + y$  where k is the largest integer satisfying this equation. Computes C-like  $x_1\%x_2$  element-wise. This was the behavior of x1%x2 in old Numeric.

absolute (x [, y])

y = |x|. Called to implement abs(x) for arrays.

 $\mathbf{rint}\ (x,\,[,\,y])$ 

Round x to the nearest integer. Rounds half-way cases to the nearest even integer.

sign (x [, y])

Sets y according to

$$sign(x) = \begin{cases} 1 & x :> 0, \\ 0 & x = 0, \\ -1 & x < 0. \end{cases}$$

 $\mathbf{conj}\ (x\ [,\ y])$ 

 $\mathbf{conjugate}\ (x\ [,\ y])$ 

 $y = \overline{x}$ ; in other words, the complex conjugate of x.

 $\exp (x [, y])$ 

 $y = e^x$ .

 $\mathbf{log}\ (x\ [,\ y])$ 

```
y = \log(x). In other words, y is the number so that e^y = x.

\exp \mathbf{m1} \ (x, [, y])

y = e^x - 1. Calculated so that it is accurate for small |x|.

\log \mathbf{1p} \ (x, [, y])

y = \log(1 + x) but accurate for small |x|. Returns the number y such that e^y - 1 = x

\log \mathbf{10} \ (x \ [, y])

y = \log \mathbf{10} \ (x). In other words, y is the number so that 10^y = x.

\operatorname{sqrt} \ (x \ [, y])

y = \sqrt{x}.

\operatorname{square} \ (x \ [, y])

y = x * x

\operatorname{reciprocal} \ (x \ [, y])

y = 1/x

\operatorname{ones\_like} \ (x, [, y])
```

y = 1 If an output argument is not given the returned data-type is the same as the input data type.



# TIP

The optional output arguments can be used to help you save memory for large calculations. If your arrays are large, complicated expressions can take longer than absolutely necessary due to the creation and (later) destruction of temporary calculation spaces. For example, the expression  $G=a^*b+c$  is equivalent to  $G=a^*B$ ; G=T+C; del t1; It will be more quickly executed as  $G=A^*B$ ; addG,C,G which is the same as  $G=A^*B$ ; G+C.

# 9.5.2 Trigonometric functions

All trigonometric functions use radians when an angle is called for. The ratio of degrees to radians is  $180^{\circ}/\pi$ .

$$\sin (x [, y])$$

$$\cos (x [, y])$$

The standard trignometric functions.  $y = \sin(x)$ ,  $y = \cos(x)$ , and  $y = \tan(x)$ .

$$\arctan (x [, y])$$

The inverse trigonometric functions:  $y=\sin^{-1}\left(x\right),\,y=\cos^{-1}\left(x\right),\,y=\tan^{-1}\left(x\right)$ . These return the value of y (in radians) such that  $\sin\left(y\right)=x$  with  $y\in\left[-\frac{\pi}{2},\frac{\pi}{2}\right]$ ;  $\cos\left(y\right)=x$  with  $y\in\left[0,\pi\right]$ ; and  $\tan\left(y\right)=x$  with  $y\in\left[-\frac{\pi}{2},\frac{\pi}{2}\right]$ , respectively.

**arctan2** 
$$(x_1, x_2 [, y])$$

Returns  $\tan^{-1}\left(\frac{x_1}{x_2}\right)$  but takes into account the sign on  $x_1$  and  $x_2$  to place the angle in the correct quadrant. The angle y is returned in the full range  $-\pi < y \le \pi$ . The angle is chosen so that  $\sin\left(y\right) = \frac{x_1}{\sqrt{x_1^2 + x_2^2}}$ , and  $\cos\left(y\right) = \frac{x_2}{\sqrt{x_1^2 + x_2^2}}$ . Particular values are showin in the following table:

$x_1$	$x_2$	$y = \arctan 2\left(x_1, x_2\right)$
0	1	0
1	0	$\frac{\pi}{2}$
0	-1	$\pi$
-1	0	$-\frac{\pi}{2}$

**hypot** 
$$(x_1, x_2 [, y])$$

Returns  $y = \sqrt{x_1^2 + x_2^2}$ . Given a complex number in cartesian form, arctan2 and hypot can be used to compute phase and magnitude, quickly.

Computes  $y = \sinh(x)$  which is defined as  $\frac{1}{2}(e^x - e^{-x})$ .

$$\cosh (x [, y])$$

```
Computes y = \cosh(x) which is defined as \frac{1}{2}(e^x + e^{-x}).

tanh (x [, y])

Computes y = \tanh(x) which is defined as (e^x - e^{-x}) / (e^x + e^{-x}).

arcsinh (x [, y])

arctanh (x [, y])
```

These compute the inverse hyperpolic functions.  $y = \operatorname{arc} func(x)$  is the (principal) value of y such that func(y) = x.

# 9.5.3 Bit-twiddling functions

These function all need integer arguments and they maniuplate the bit-pattern of those arguments.

```
bitwise_and (x_1, x_2 [, y])
```

Each bit in y is the result of a bit-wise 'and' operation on the corresponding bits in  $x_1$  and  $x_2$ . Called to implement x1&x2 for arrays.

```
bitwise_or (x_1, x_2 [, y])
```

Each bit in y is the result of a bit-wise 'or' operation on the corresponding bits in  $x_1$  and  $x_2$ . Called to implement  $x_1 \mid x_2$  for arrays.

```
bitwise_xor (x_1, x_2 [, y])
```

Each bit in y is the result of a bit-wise 'xor' operation on the corresponding bits in  $x_1$  and  $x_2$ . An xor operation sets the output to 1 if one and only one of the input bits is 1. Called to implement  $x1^x2$  for arrays. Using the bitwise\_xor operation and the optional output argument you can swap the values of two integer arrays of equivalent types without using temporary arrays.

```
>>> a=arange(10)
>>> b=arange(10,20)
>>> bitwise_xor(a,b,a); bitwise_xor(a,b,b);
>>> bitwise_xor(a,b,a)
array([10, 11, 12, 13, 14, 15, 16, 17, 18, 19])
>>> print a; print b
[10 11 12 13 14 15 16 17 18 19]
[0 1 2 3 4 5 6 7 8 9]
```

```
invert (x, [, y])
```

Each bit in y is the opposite of the corresponding bit in x. Called to implement "x for arrays.

```
left_shift (x_1, x_2 [, y])
```

Shifts the bits of  $x_1$  to the left by  $x_2$ . Called to implement x1<<x2 for arrays. Provided there is no overflow, the result is equal to  $y = x_1 2^{x_2}$ .

right\_shift 
$$(x_1, x_2 [, y])$$

Shifts the bits of  $x_1$  to the right by  $x_2$ . Called to implement x1>>x2 for arrays. Absent overflow, the result is equal to  $y=x_12^{-x_2}$ .

# 9.5.4 Comparison functions

All of these functions (except maximum, minimum, and sign) return Boolean arrays.

```
greater (x_1, x_2 [, y])
greater_equal (x_1, x_2 [, y])
less (x_1, x_2 [, y])
less_equal (x_1, x_2 [, y])
not_equal (x_1, x_2 [, y])
equal (x_1, x_2 [, y])
```

These functions are called to implement x1>x2, x1>=x2, x1<x2, x1<=x2, x1!=x2 (or x1<>x2), and x1==x2, respectively, for arrays.

The fact that these functions return Boolean arrays make them very useful in combination with advanced array indexing. Thus, for example, arr[arr>10] = 10 clips large values to 10. Used in conjunction with bitwise operators quite compilicated expressions are possible. For example, arr[~((arr<10)&(arr>5))] = 0 clips all values outside of the range (5, 10) to 0.



# WARNING

Do not use the Python keywords and and or to combine logical array expressions. These keywords will test the truth value of the entire array (not element-by-element as you might expect). Use the bitwise operators: & and | instead.

logical\_and  $(x_1, x_2 [, y])$ 

The output is the truth value of  $x_1$  and  $x_2$ . Numbers equal to 0 are considered False. Nonzero numbers are True.

 $logical\_or (x_1, x_2 [, y])$ 

The output, y, is the truth value of  $x_1$  or  $x_2$ .

 $logical\_xor (x_1, x_2 [, y])$ 

The output, y, is the truth value of  $x_1$  **xor**  $x_2$ , which is the same as  $(x_1$  and not  $x_2$ ) or (not  $x_1$  and  $x_2$ ).

 $logical\_not (x [, y])$ 

The output, y is the truth value of **not** x.



#### WARNING

The Bitwise operators (& and |) are the proper way to combine element-by-element array comparisons. Be sure to understand the operator precedence: (a>2) & (a<5) is the proper syntax because a>2 & a<5 will result in an error due to the fact that 2 & a is evaluated first.

maximum  $(x_1, x_2 [, y])$ 

The output, y, is the larger of  $x_1$  and  $x_2$ .

```
>>> maximum([1,0,5,10],[3,2,4,5])
array([3, 2, 5, 10])
>>> max([1,0,5,10],[3,2,4,5])
[3, 2, 4, 5]
```



#### TIP

The Python function max() will find the maximum over a onedimensional array, but it will do so using a slower sequence interface. The reduce method of the maximum ufunc is much faster. Also, the max() method will not give answers you might expect for arrays with greater than one dimension. The reduce method of minimum also allows you to compute a total minimum over an array.

```
minimum (x_1, x_2 [, y])
```

The output, y, is the smaller of  $x_1$  and  $x_2$ .

```
>>> minimum([1,0,5,10],[3,2,4,5])
array([1, 0, 4, 5])
>>> min([1,0,5,10],[3,2,4,5])
[1, 0, 5, 10]
```



# WARNING

the behavior of maximum(a,b) is than that of max(a,b). As a ufunc, maximum(a,b) performs an element-by-element comparison of a and b and choses each element of the result according to which element in the two arrays is larger. In contrast, max(a,b) treats the objects a and b as a whole, looks at the (total) truth value of a>b and uses it to return either a or b (as a whole). A similar difference exists between minimum(a,b) and min(a,b).

# 9.5.5 Floating functions

Recall that all of these functions work element-by-element over an array, returning an array output. The description details only a single operation.

# isreal (x)

True if x has an imaginary part that is 0.

#### iscomplex (x)

True if x has an imaginary part that is non-zero.

# isfinite (x)

True if x is a finite floating point number (not a NaN or an Inf).

#### isinf (x)

True if x is  $\pm \infty$ .

# isnan (x)

True if x is Not-a-Number. This represents invalid results. When a NaN is created, the invalid flag is set. If you set the error mode of invalid to 'warn', 'raise', or 'call', then the appropriate action will be performed on NaN creation.

# signbit (x)

True where the sign bit of the floating point number is set. This should correspond to x > 0 when x is a finite number. When, x is NaN or infinite, then this tests the actual signbit.

**modf** 
$$(x [, y_1, y_2])$$

Breaks up the floating point value x into its fractional,  $y_1$ , and integral,  $y_2$ , parts. Thus,  $x = y_1 + y_2$  with floor(y2)==y2.

$$\mathbf{ldexp}\ (x,\,n\ [,\,y])$$

Fast multiply of a floating point number by an integral power of 2:  $y = 2^n x$ .

frexp 
$$(x [, y, n])$$

Breaks up the floating point value x into a normalized fraction, y and an exponent, n which corresponds to how the value is represented in the computer. The results are such that  $x = y2^n$ . Effectively, the inverse of ldexp.

**fmod** 
$$(x_1, x_2 [, y])$$

Computes the remainder of dividing  $x_1$  by  $x_2$ . The result, y, is  $x_1 - nx_2$  where n is the quotient (rounded towards zero to an integer) of  $x_1/x_2$ .

floor 
$$(x [,y])$$

Return  $y = \lfloor x \rfloor$  where y is the nearest integer smaller-than or equal to x. Thus,  $\lfloor x \rfloor \leq x < \lfloor x \rfloor + 1$ .

**ceil** 
$$(x [,y])$$

Return  $y = \lceil x \rceil$  where y is the nearest integer greater-than or equal to x. Thus,  $x \leq \lceil x \rceil < x+1$ .

# Chapter 10

# Basic Modules

The NumPy distribution contains some basic functionality equivalent to what was available in the Numeric packages previously. This section documents the new interfaces. These are sub-packages of the NumPy namespace. The linalg and fft capabilities are useful but limited. You should install the full SciPy package to access more functionality. The numpy dual module contains functions that are defined in both SciPy and NumPy. If SciPy defines func, then numpy dual func will point to the SciPy version, otherwise it will point to the NumPy version. It must be imported specifically to be used. Table XXX shows the functions defined in numpy dual that are in both NumPy and SciPy.



# TIP

The fft, ifft, rand, and randn functions are also available in the NumPy namespace due to the frequency of their use.

# 10.1 Linear Algebra (linalg)

These functions are in the numpy.linalg sub-package.

Table 10.1: Functions in numpy.dual (both in NumPy and SciPy)

Family	Functions
Fourier Transforms	fft, ifft, fft2, ifft2, fftn, ifftn
Linear Algebra	inv, svd, solve, det, eig, eigvals, lstsq, pinv, cholesky
Special Functions	i0

#### inv (A)

Return the (matrix) inverse of the 2-d array A. The result, X, is such that dot(A,X) is equal to eye(\*A.shape) (to within machine precision).

# solve (A,b)

Find the solution to the linear equation  $\mathbf{A}\mathbf{x} = \mathbf{b}$ , where A is a 2-d array and b is a 1-d or 2-d array.

# cholesky (A)

Return,  $\mathbf{L}$ , the Cholesky decomposition of  $\mathbf{A}$ . Cholesky decomposition is applicable to a Hermitian, positive definite matrices. When  $\mathbf{A} = \mathbf{A}^H$  and  $\mathbf{x}^H \mathbf{A} \mathbf{x} \geq 0$  for all  $\mathbf{x}$ , then decompositions of  $\mathbf{A}$  can be found so that  $\mathbf{A} = \mathbf{L} \mathbf{L}^H$ , where  $\mathbf{L}$  is lower-triangular.

# eigvals (A)

Return all solutions ( $\lambda$ ) to the equation  $\mathbf{A}\mathbf{x} = \lambda \mathbf{x}$ .

# eig (A)

Return all solutions  $(\lambda, \mathbf{x})$  to the equation  $\mathbf{A}\mathbf{x} = \lambda \mathbf{x}$ . The first element of the return tuple contains all the eigenvalues. The second element of the return tuple contains the eigenvectors in the columns  $(\mathbf{x}[:,i]$  is the ith eigenvector).

#### eigvalsh (U)

#### eigh (U)

These functions are identical to eigvals and eig except they only work with Hermitian matrices where  $\mathbf{U}^H = \mathbf{U}$  (only the lower-triangular part of the array is used).

# svd (A)

Compute the singular value decomposition of the 2-d array  $\mathbf{A}$ . Every  $m \times n$  matrix can be decomposed into a pair of unitary matrices,  $\mathbf{U} = \mathbf{U}^H$   $(m \times m)$  and  $\mathbf{V} = \mathbf{V}^H$   $(n \times n)$  and an  $m \times n$  "diagonal" matrix  $\mathbf{\Sigma}$ , such that  $\mathbf{A} = \mathbf{U}\mathbf{\Sigma}\mathbf{V}^H$ . The only non-zero portion of  $\mathbf{\Sigma}$  is the upper  $r \times r$  block where  $r = \min(m, n)$ . The svd function returns three arrays as a tuple:  $(\mathbf{U}, \boldsymbol{\sigma}, \mathbf{V}^H)$ . The singular values are returned in the 1-d array  $\boldsymbol{\sigma}$ . If needed, the array  $\mathbf{\Sigma}$  can be found (if really needed) using the command  $\operatorname{diag}(\boldsymbol{\sigma})$  which creates the  $r \times r$  diagonal block and then inserting this into a zeros matrix:

**pinv** (A, rcond=
$$10^{-10}$$
)

Return the generalized, pseudo inverse, of A. For invertible matrices, this is the same as the inverse.

# det (A)

Return the determinant of the array. The determinant of an array is the product of its singular values.

**lstsq** (A, b, rcond=
$$10^{-10}$$
)

Return (x, resids, rank, s) where x minimizes resids= $\|\mathbf{A}\mathbf{x} - \mathbf{b}\|_2$ . The output rank is the rank of A and s is the singular values of a in descending order. Singular values less than s[0]\*rcond are treated as 0. If the rank of A is less than the number of columns of A or greater than the number of rows, resids will be returned as an empty array.

# 10.2 Discrete Fourier Transforms (dft)

All of the algorithms here are most efficient if the length of the data is a power of 2 (or decomposible into low prime factors).

**fft** 
$$(x, n=None, axis=-1)$$

Return, X, the N-point Discrete Fourier Transform (DFT) of x along the given axis using a fast algorithm. If N is larger than x.shape[axis], then x will be zero-padded. If N is smaller than x.shape[axis], then the first N items will be used. The result is computed for  $k=0\ldots n-1$  from the formula:

$$X[k] = \sum_{m=0}^{n-1} x[m] \exp\left(-j\frac{2\pi km}{n}\right).$$



# TIP

The fft returns values for  $k=0\ldots N-1$ . Because X[N-k]=X[-k] in the FFT formula, larger values of k correspond also to negative frequencies.

**ifft** (X, n=None, axis=-1)

Return the inverse of the fft so that (ifft(fft(a)) == a within numerical precision. The order of frequencies must be the same as returned by fft. The result is computed (using a fast algorithm) for  $m = 0 \dots n-1$  from the formula:

$$x[m] = \sum_{k=0}^{n-1} X[k] \exp\left(j\frac{2\pi km}{n}\right).$$

Sometimes having the "negative" frequencies at the end of the output returned by fft can be a little confusing. There are two ways to deal with this confusion. In my opinion, the most useful way is to get a collection of DFT sample frequencies and use them to keep track of where each frequency is. The function fftfreq provides these sample frequencies. Making an x-y plot, where the sample frequencies are along the "x"-axis and the result of the DFT is along the "y"-axis provides a useful visualization with the frequencies at the center. The advantage of this approach is that your data is still in proper order for using the ifft function. Some people, however, prefer to simply swap one-half of the output with the other. This is exactly what the function fftshift does. Of course, now the data is not in the proper order for the ifft function, but to each his own.

The reason that the "negative" frequencies are in the upper part of the return signal was given in the description of the DFT. The reason is that the output of the DFT is just one period of a periodic function (with period n). The traditional output of the FFT algorithm is to provide the portion of the function from from k=0 to k=n-1.

fftshift (x, axes=None)

Shift zero-frequency component to the center of the spectrum. This function swaps half-spaces for all axes listed (defaults to all of them).

**ifftshift** (x, axes=None)

Reverse the effect of the fftshift operation. Thus, it takes zero-centered data and shifts it into the correct order for the ifft operation.

fftfreq (n, d=1.0)

Provide the DFT sample frequencies. The returned float array contains the frequency bins in the order returned from the fft function. If given, d represents the sample-spacing. The units on the frequency bins are cycles / unit. For example, the following example computes the output frequencies (in Hz) of the fft of 256 samples of a voice signal sampled at 20000Hz.

```
>>> from numpy.dft import fftfreq; f=fftfreq(256,d=1./20e3)
>>> print f[0], f[1], f[2], f[128]
0.0 78.125 156.25 -10000.0
```

fft2 (x, s=None, axes=(-2,-1))

Return the two-dimensional fft of the array x for each 2-d array formed by axes. The 2-d fft is computed as

$$X\left[k_{0}, k_{1}\right] = \sum_{m_{0}=0}^{s[0]-1} \sum_{m_{1}=0}^{s[1]-1} x\left[m_{0}, m_{1}\right] \exp\left(-j\frac{2\pi k_{0}m_{0}}{s[0]}\right) \exp\left[-j\frac{2\pi k_{1}m_{1}}{s\left[1\right]}\right]$$

and can be realized by repeated application of the 1-d fft (first over the axes[0] and then over axes[1]). In other-words fft2(x,s,axes) is equivalent to fft(fft(x,s[0], axes[0]), s[1], axes[1]). The 2-d fft is returned for  $k_0 = 0 \dots s[0] - 1$  and  $k_1 = 0 \dots s[1] - 1$ . Symmetry  $(X[s[0] - k_0, s[1] - k_1] = X[-k_0, -k_1])$  ensures that higher values of  $k_i$  correspond to negative frequencies.

ifft2 (X, s=None, axes=(-2,-1))

Return the inverse of the two-dimension fft. Thus, ifft2(fft2(x)) == x to within numerical precision. Note that the "negative frequencies" must be

**fftn** (x, s=None, axes=None)

Return the N-dimensional fft of x. If s is not given, then if axes is given, then N=len(axes), otherwise N=x.ndim. If s is given, then N=len(s). Results are computed using a simliar formula as for the 1- and 2-d FFT with N summations.

**ifftn** (X, s=None, axes=None)

Return the N-dimensional inverse fft of X. Note that ifftn(fftn(x)) == x to within machine precision.

The Discrete Fourier transform returns complex-valued data (even for real-valued input). If the data was originally real-valued, then much of the output of the full DFT is redundant. Notice that if x[m] is real, then

$$X[n-k] = \sum_{m=0}^{n-1} x[m] \exp\left(-j\frac{2\pi(n-k)m}{n}\right)$$
$$= \left[\sum_{m=0}^{n-1} x[m] \exp\left(-j\frac{2\pi km}{n}\right)\right]^*$$
$$= X^*[k],$$

where  $a^*$  denotes the complex-conjugate of a. So, the upper half of the fft output (the negative frequencies) is determined exactly by the lower half of the output when the input is purely real. This kind of symmetry is called Hermitian symmetry. The real-valued Fourier transforms described next take advantage of Hermitian symmetry to compute only the unique outputs more quickly.

The symmetry in higher dimensions is always about the origin. If N is the number of dimensions, then:

$$X[n_1 - k_1, n_2 - k_2, \dots n_N - k_N] = X^* [k_1, k_2, \dots, k_N].$$

Thus, the data-savings remains constant at about 1/2 for higher dimensions as well.

**rfft** 
$$(x, n=None, axis=-1)$$

Compute the first n//2+1 points of the *n*-point discrete Fourier transform of the real valued data along the given axis. The returned array will be just the first half of the fft, corresponding to positive frequencies: rfft(x) = fft(x)[:n//2+1]

Compute the inverse *n*-point discrete Fourier transform along the given axis using the first n/(2+1) points. To within numerical precision, irfft(rfft(x)) = = x.

**rfft2** (x, s=None, axes=
$$(-2, -1)$$
)

Compute only the first half-plane of the two-dimensional discrete Fourier transform corresponding to unique values. s[0] and s[1]-point DFTs will be computed along axes[0] and axes[1] dimensions, respectively. Requires a real array. If s is None it defaults to the shape of x. The real fft will be computed along the last axis specified in axes while a full fft will be computed in the other dimension.

```
irfft2 (X, s=None, axes=(-2, -1))
```

Compute the inverse of the 2-d DFT using only the first quadrant. Returns a real array such that to within numerical precision irfft2(rfft2(x)) == x.

```
rfftn (x, s=None, axes=None)
```

Compute only the unique part of the N-dimensional DFT from a real-valued array. If s is None it defaults to the shape of x. If axes is not given it defaults to all the axes (-n, ..., -1). The length of axes provides the dimensionality of the DFT. The unique part of the real N-dimensional DFT is obtained by slicing the full fft along the last axis specified and taking n/2+1 slices. rfftn(x) == fft(x)[sliceobj] where sliceobj[axes[-1]] = slice(None,s[-1]/2+1,None).

```
irfftn (X, s=None, axes=None)
```

Compute the inverse DFT from the unique portions of the N-dimensional DFT provided by rfftn.

Occasionally, the situation may arise where you have complex-valued data with Hermitian symmetry (or real-valued symmetric data). This ensures that the Fourier transform will be real. The two functions below can calculate it without wasting extra space for the zero-valued imaginary entries of the Discrete Fourier transform, or the entire signal.

```
hfft (x, n=None, axis=-1)
```

Calculate the n-point real-valued Fourier transform from (the first half of Hermitian-symmetric data, x.

```
ihfft (X, n=None, axis=-1)
```

Return (the first half-of) Hermitian-symmetric data from the real-valued Fourier transform, X.

# 10.3 Random Numbers (random)

The random number capabilities surpass those that were available in Numeric. The random number facilities were generously contributed by Robert Kern, who has been a dedicated and patient help to many mailing list questioners. Robert built the random package using PyRex to build on top of his own code as well as that of randomkit by Jean-Sebastien Roy as well as code by Ivan Frohne. The fundamental random number generator is the Mersenne Twister based on code written by Makoto Matsumoto and Takuji Nishimura (and modified for Python by

Raymond Hettinger). Random numbers from discrete and continuous distributions are available, as well as some useful random-number-related utilities. Many of the random number generators are based on algorithms published by Luc Devroye in "Non-Uniform Random Variate Generation" available electronically at http://cgm.cs.mcgill.ca/~luc/rnbookindex.html

Each of the discrete and continuous random number generators take a size keyword. If this is None (default), then a single number is generated from the selected distribution. If this is an integer, then a 1-d array of that size is generated filled with random numbers from the selected distribution. Finally, if size is a tuple, then an array of that shape is returned filled with random numbers.

#### 10.3.1 Discrete Distributions

Discrete random numbers take on only a countable number of values (typically integers). Each distribution has associated with it a probability mass function (pmf),  $p_m(k;\cdot)$ , that is defined as the probability that the returned random number is k. The arguments after k represent possible parameters to the distribution. Thus, let  $X(\cdot)$  represent the random number generator for a particular distribution. Then,

$$p_m(k;\cdot) = \text{Probability} \{X(\cdot) = k\}.$$

It will be useful to define the discrete indicator function,  $I_S(k)$ , where S is a set of integers (often represented by an interval).  $I_S(k) = 1$  if  $k \in S$ , otherwise  $I_S(k) = 0$ . This convenient notation isolates the relavence of a particular functional form to a certain range. Also, the formulas below make use of the following definition:

$$\left(\begin{array}{c} n \\ k \end{array}\right) = \frac{n!}{k! \, (n-k)!}$$

where  $k! = k \cdot (k-1) \cdot \cdots \cdot 2 \cdot 1$ .

**binomial** (n, p, size=None)

This random number models the number of successes in n independent trials of a random experiment where the probability of success in each experiment is p.

$$p_m(k) = \binom{n}{k} p^k (1-p)^{n-k} I_{[0,n]}(k).$$

**geometric** (p, size=None)

This random number models the number of (independent) attempts required to obtain a success where the probability of success on each attempt is p.

$$p_m(k; p) = (1-p)^{k-1} p I_{[1,\infty)}(k).$$

hypergeometric (ngood, nbad, nsample, size=None)

Imagine a probability theorists favorite urn filled with  $ngood\ (n_g)$  "good" objects and  $nbad\ (n_b)$  "bad" objects. In other words there are two types of objects in a jar. The hypergeometric random number models how many "good" objects will be present when  $nsample\ (N)$  items are taken out of the urn without replacement.

$$p\left(k; n_{g}, n_{b}, N\right) = \frac{\left(\begin{array}{c}n_{g}\\k\end{array}\right)\left(\begin{array}{c}n_{b}\\N-k\end{array}\right)}{\left(\begin{array}{c}n_{g}+n_{b}\\N\end{array}\right)} I_{\left[N-n_{b}, \min\left(n, N\right)\right]}\left(k\right).$$

logseries (p, size=None)

A random number whose pmf with terms proportional to the Taylor series expansion of  $\log (1-p)$ . It has been used in biological studies to model the species abundance distribution.

$$p_m(k; p) = -\frac{p^k}{k \log(1-p)} I_{[1,\infty)}(k).$$

multinomial (n, pvals, size=None)

This generator produces random vectors of length N where N = len (pvals). The shape of the returned array is always the shape indicated by size + (N,). The multinomial distribution is a generalization of the binomial distribution. This time, n trials of an experiment are independently repeated but each trial results in N possible integers  $k_1, k_2, \ldots, k_N$  with  $\sum_{i=1}^{N} k_i = n$ .

$$p_{m}\left(k_{1}, k_{2}, \dots, k_{N}; \cdot\right) = \operatorname{Probability}\left\{X\left(\cdot\right) = \left[k_{1}, k_{2}, \dots, k_{N}\right]\right\}$$
$$= \frac{n!}{k_{1}! k_{2}! \cdots k_{N}!} p_{1}^{k_{1}} p_{2}^{k_{2}} \cdots p_{N}^{k_{N}}$$

where  $pvals = [p_1, p_2, ..., p_N]$ . It must be true that  $\sum_{i=1}^{N} p_i = 1$ . Therefore, as long as  $\sum_{i=1}^{N-1} p_i \leq 1$ , the last entry in pvals is computed as  $1 - \sum_{i=1}^{N-1} p_i$ .

negative\_binomial (n, p, size=None)

Models the number of extra independent trials (beyond n) required to accumulate a total of n successes where the probability of success on each trial is p. Equivalently, this random number models the number of failures encountered while accumulating n successes during independent trials of the experiment that succeeds with probability, p.

$$p_m(k; n, p) = {k+n-1 \choose n-1} p^n (1-p)^k I_{[0,\infty)}(k).$$

poisson (lam=1.0, size=None)

This random number counts the number of successes in n independent experiments (where the probability of success in each experiment is p) in the limit as  $n \to \infty$  and  $p \to 0$  gets very small such that  $\lambda = np \ge 0$  is a constant. It can be used, for example, to model how many typographical errors are on each page of a book.

$$p(k; \lambda) = e^{-\lambda} \frac{\lambda^k}{k!} I_{[0,\infty)}(k).$$

**zipf** (a, size=None)

The probability mass function of this random number (also called the zeta distribution) is

$$p_m(k;a) = \frac{1}{\zeta(a) k^a} I_{[1,\infty)}(k),$$

where

$$\zeta\left(a\right) = \sum_{n=1}^{\infty} \frac{1}{n^a}$$

is the Riemann zeta function. Zipf distributions have been shown to characterize use of words in a natural language (like English), the popularity of library books, and even the use of the web. The Zipf distribution describes collections that have a few items whose probability of selection is very high, a medium number of items whose probability of selection is medium, and a huge number of items whose probability of selection is very low.

# 10.3.2 Continuous Distributions

Continuous random numbers can take on an uncountable number of values. Therefore, the value returned by a continuous distribution is denoted x. Because there

is an uncountable number of possibilities for the random number<sup>1</sup>, a continuous distribution is modeled by a probability density function,  $f(x;\cdot)$ . To obtain the probability that the random number generated by  $X(\cdot)$  is in a certain interval, we integrate this density function:

$$\int_{-\infty}^{b} f(x) dx = \text{Probability} \{X(\cdot) \leq b\}.$$

To obtain a probability, we have to integrate f(x) which is why it is called a density function. Most continuous distributions are defined by their probability density functions (pdf). Some have basic origins, a few are derived from other distributions, and some are used mainly for modelling unknown distributions.

Some of the parameters of the distributions are labeled as location (loc) and scale parameters. These parameters are not shown in the equation for the pdf. because they affect the distribution in a known way. This is due to the fact that if X is a number drawn from a distribution with pdf  $f_X(x)$ , then Y = Sx + L is a number drawn from a distribution with pdf

$$f_Y(y) = \frac{1}{S} f_X\left(\frac{y-L}{S}\right).$$

Thus, from the standard from provided, the pdf of the actual random numbers generated by fixing the location and scale parameters can be quickly found.

In this section, the indicator function  $I_A(x)$  will be used where A is a set defined over all the real numbers. For clarity,

$$I_A(x) = \begin{cases} 1 & x \in A, \\ 0 & x \notin A. \end{cases}$$

Also, the following functions are used in the definitions:

$$\begin{split} &\Gamma\left(x\right) &= \int_{0}^{\infty}t^{x-1}e^{-t}dt = \left(x-1\right)\Gamma\left(x-1\right),\\ &B\left(a,b\right) &= \frac{\Gamma\left(a\right)\Gamma\left(b\right)}{\Gamma\left(a+b\right)}. \end{split}$$

**beta** (a, b, size=None)

<sup>&</sup>lt;sup>1</sup>A computer actually always generates a random number from a discrete distribution because there are only a finite set of numbers that can be represented by a computer. However, for continuous random number generators, the resulting random numbers usually approximate the continuous distribution well enough to ignore the subtlety.

$$f(x; a, b) = \frac{1}{B(a, b)} x^{a-1} (1 - x)^{b-1} I_{(0,1)}(x).$$

**chisquare** ( $\nu$ , size=None)

If  $Z_1, \ldots, Z_{\nu}$  are random numbers from standard normal distributions, then  $W = \sum_{k=1}^{\nu} Z_k^2$  is a random number from the chi-square  $(\chi^2)$  distribution with  $\nu$  degrees of freedom.

$$f\left(x;\nu\right) = \frac{1}{2\Gamma\left(\frac{\nu}{2}\right)} \left(\frac{x}{2}\right)^{\nu/2-1} e^{-x/2} I_{\left[0,\infty\right)}\left(x\right).$$

exponential (scale=1.0, size=None)

$$f(x) = e^{-x} I_{[0,\infty)}(x)$$
.

 $\mathbf{f}$  ( $\nu_1, \nu_2, \text{ size=None}$ )

The distribution of  $\frac{X_1/\nu_1}{X_2/\nu_2}$  where  $X_i$  is chi-squared with  $\nu_i$  degrees of freedom.

$$f(x;\nu_1,\nu_2) = \frac{\nu_2^{\nu_2/2} \nu_1^{\nu_1/2} x^{\nu_1/2-1}}{(\nu_2 + \nu_1 x)^{(\nu_1 + \nu_2)/2} B\left(\frac{\nu_1}{2}, \frac{\nu_2}{2}\right)} I_{[0,\infty)}(x).$$

gamma (a, scale=1.0, size=None)

$$f(x; a) = \frac{1}{\Gamma(a)} x^{a-1} e^{-x} I_{[0,\infty)}(x).$$

gumbel (loc=0.0, scale=1.0, size=None)

A right-skewed extreme value distribution.

$$f\left(x\right) = \exp\left[-x - e^{-x}\right].$$

laplace (loc=0.0, scale=1.0, size=None)

$$f(x) = \frac{1}{2}e^{-|x|}.$$

lognormal ( $\mu$ =0.0,  $\sigma$ =1.0, size=None)

$$f(x; \mu, \sigma) = \frac{1}{\sigma x \sqrt{2\pi}} \exp \left[ -\frac{1}{2} \left( \frac{\log x - \mu}{\sigma} \right)^2 \right] I_{[0, \infty)}(x),$$

The parameters,  $\mu$  and  $\sigma$  are not the mean and variance of this distribution, but the parameters of the underlying normal distribution. Random numbers from this distribution are generated as  $e^{\sigma Z + \mu}$  where Z is a standard normal random number.

logistic (loc=0.0, scale=1.0, size=None)

$$f(x) = \frac{e^{-x}}{[1 + e^{-x}]^2} I_{[0,\infty)}(x)$$

# multivariate\_normal ( $\mu$ , C, size=None)

Returns a vector of random numbers which are jointly drawn from a multivariate normal distribution. The last-dimension of the output array contains the sample vector, which is of length  $N = \text{len}\,(mean)$ . The covariance matrix must be  $N \times N$ . If  $\mu \equiv mean$  and  $\mathbf{C} = cov$ , then the joint-pdf representing the returned random vector(s) is

$$f(\mathbf{x}) = \frac{1}{\sqrt{(2\pi)^N |\mathbf{C}|}} \exp\left[-\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^T \mathbf{C}^{-1} (\mathbf{x} - \boldsymbol{\mu})\right].$$

# **noncentral\_chisquare** ( $\nu$ , $\lambda$ , size=None)

This is the distribution of  $\sum_{i=1}^{\nu} (Z_i + \delta_i)^2$  where  $Z_i$  are independent standard normal random numbers and  $\delta_i$  are constants. It is a generalized Rayleigh-Rice distribution:

$$f\left(x;\nu,\lambda\right)=e^{-(\lambda+x)/2}\frac{1}{2}\left(\frac{x}{\lambda}\right)^{(\nu-2)/4}I_{(\nu-2)/2}\left(\sqrt{\lambda x}\right)I_{(0,\infty)}\left(x\right),$$

where  $I_{\nu}\left(z\right)$  (a real-number in the subscript, not an interval) is the modified Bessel Function of the first kind.

# **noncentral\_f** ( $\nu_1$ , $\nu_2$ , $\lambda$ , size=None)

The pdf of this distribution is

$$f(x; \nu_1, \nu_2, \lambda) = \exp\left[\frac{\lambda}{2} + \frac{\lambda v_1 x}{2(\nu_1 x + \nu_2)}\right] \nu_1^{\nu_1/2} \nu_2^{\nu_2/2} x^{\nu_1/2 - 1}$$

$$\times (\nu_{2} + \nu_{1}x)^{-(\nu_{1} + \nu_{2})/2} \times \frac{\Gamma\left(\frac{\nu_{1}}{2}\right)\Gamma\left(1 + \frac{\nu_{2}}{2}\right)L_{n_{2}/2}^{n_{1}/2 - 1}\left(-\frac{\lambda\nu_{1}x}{2(\nu_{1}x + \nu_{2})}\right)}{B\left(\frac{\nu_{1}}{2}, \frac{\nu_{2}}{2}\right)\Gamma\left(\frac{\nu_{1} + \nu_{2}}{2}\right)}.$$

**normal** (loc=0.0, scale=1.0, size=None)

The normal, or Gaussian, distribution is the limiting distribution of independent samples from any sufficiently well-behaved distributions (this is the content of the celebrated central limit theorem). The normal distribution is also the distribution of maximum entropy when the mean and variance alone are fixed. These two facts account for its name as well as the wide variety of situations that can be usefully modelled using the normal distribution. Because it is so widely used, the full pdf with the location  $(\mu)$  and scale  $(\sigma)$  parameters is provided:

$$f(x) = \frac{1}{\sigma\sqrt{2\pi}} \exp\left[-\frac{(x-\mu)^2}{2\sigma^2}\right].$$

pareto (a, size=None)

$$f(x; a) = \frac{a}{x^{a+1}} I_{[1,\infty)}(x)$$
.

**power** (a, size=None)

A special case of the beta distribution with b = 1.

$$f(x; a) = ax^{a-1}I_{[0,1]}(x)$$
.

rand  $(d_1, d_2, ..., d_n)$ 

A convenient interface to obtain an array of shape  $(d_1, d_2, \ldots, d_n)$  of uniform random numbers in the interval [0,1). This function is available in the NumPy namespace. Notice the different convention for passing in the shape (as separate arguments intend of a tuple). The standard convention is used in the function numpy.random.random(shape) for which this function is merely a convenient short-hand. If you have a tuple named shape, then rand(\*shape) will work correctly.

randint (low, high=None, size=None)

Equally probably random integers in the range  $low \le x < high$ . If high is None, then the range is  $0 \le x < low$ . Similar to random\_integers, but check the difference on the bounds.

# randn (\*args)

A convenient interface to obtain an array of shape  $(d_1, d_2, \ldots, d_n)$  of standard normal  $(\mu = 0, \sigma = 1)$  random numbers. This function is available in the NumPy namespace as well. Notice the different convention for passing in the shape (as separate arguments intend of a tuple). The standard convention is used in the function numpy.random.standard\_normal(shape) for which this function is merely a convenient short-hand. If you have a tuple named shape, then randn(\*shape) will work correctly.

# random\_integers (low, high=None, size=None)

Equally probably random integers in the range  $low \le x \le high$ . If high is None, then the range is  $1 \le x \le low$ . Similar to randint, but check the difference on the bounds.

rayleigh (scale=1.0, size=None)

Rayleigh-distributed random numbers can be obtained as  $X = \sqrt{Z_1^2 + Z_2^2}$  where  $Z_i$  are independent standard normal random numbers. The scale parameter is also the mode of the distribution (the value of X with highest probability).

$$f(x) = xe^{-x^2/2}I_{[0,\infty)}(x)$$

standard\_cauchy (size=None)

A Cauchy distribution is a heavy-tailed distribution with no variance. It's distribution is that of the ratio of two standard normal distributions  $Z_1/Z_2$ .

$$f\left(x\right) = \frac{1}{\pi\left(1 + x^2\right)}.$$

#### standard\_exponential (size=None)

A standard exponetial random number with scale=1.0. The pdf was given under the description of random.exponential.

standard\_gamma (a, size=None)

A standard gamma random number with scale=1.0. The pdf was given under the description of random.gamma.

# standard\_normal (size=None)

A zero-mean, unit-variance, normally distributed random number often denoted Z.

$$f(x) = \frac{1}{\sqrt{2\pi}}e^{-x^2/2}.$$

standard\_t ( $\nu$ , size=None)

Often called Student's t distribution, this random number distribution arises in the problem of estimating the mean of normally distributed samples when the sample-size is small. The first parameter,  $\nu$ , is the number of degrees of freedom of the distribution.

$$f(x;\nu) \frac{\Gamma\left(\frac{\nu+1}{2}\right)}{\sqrt{\pi\nu}\Gamma\left(\frac{\nu}{2}\right)\left[1+\frac{x^2}{\nu}\right]^{\frac{\nu+1}{2}}}.$$

triangular (left, mode, right, size=None)

Returns random numbers according to a triangularly-shaped density that starts at left, peaks at mode, and ends at right.

uniform (low=0.0, high=1.0, size=None)

Returns random numbers that are equally probable over the range [low, high].

vonmises 
$$(\mu, \kappa, \text{size=None})$$

A continuous distribution that is well suited for circular attributes such as angles, time of day, day of the year, etc. The mean direction is  $\mu$  and concentration (or dispersion) parameter is  $\kappa$ . For small  $\kappa$  the distribution tends towards a uniform distribution over  $[-\pi, \pi]$ . For large  $\kappa$ , the distribution tends towards a normal distribution with mean  $\mu$  and variance  $1/\kappa$ .

$$f(x) = \frac{e^{\kappa \cos(x-\mu)}}{2\pi I_0(\kappa)} I_{[-\pi,\pi]}(x).$$

wald  $(\mu, \lambda, \text{size=None})$ 

This distribution is also called the inverse Gaussian distribution (and the Wald distribution considered as a special case when  $\mu = \lambda$ ). It can be generated by noticing that if X is a wald random number then  $\frac{\lambda(X-\mu)^2}{\mu^2X}$  is the square

of a standard normal random number (i.e. it is chi-square with one degree of freedom). The pdf is

$$f\left(x\right) = \sqrt{\frac{\lambda}{2\pi x^{3}}} e^{-\frac{\lambda(x-\mu)^{2}}{2\mu^{2}x}}.$$

**weibull** (a, size=None)

An extreme-value distribution:

$$f\left(x;c\right)=ax^{a-1}\exp\left(-x^{a}\right)I_{\left(0,\infty\right)}\left(x\right).$$

# 10.3.3 Miscellaneous utilities

bytes (length)

Return a string of random bytes of the provided length.

# get\_state ()

Return an object that holds the state of the random number generator (allows you to retstart simulations where you left off).

# set\_state (state)

Set the state of the random number generator. The argument should be the returned object of a previous get\_state command.

#### **shuffle** (sequence)

Randomly permute the items of any sequence. If sequence is an array, then it must be 1-d.

# permutation (n)

Return a permutation of the integers from 0 to n-1.

# Chapter 11

# Testing and Packaging

There are two additional sub-packages distributed with NumPy that simplify the process of distributing and testing code based on NumPy. The numpy distutils sub-package extends the standard distutils package to handle Fortran code along with providing support for the auto-generated code in NumPy. The numpy testing sub-package defines a few functions and classes for standardizing unit-tests in NumPy. These facilities can be used in your own packages that build on top of NumPy.

# 11.1 Testing

In this sub-package are two classes and some useful utilities for writing unit-tests

ScipyTestCase a subclass of unittest. TestCase which adds a measure method that can determine the elasped time to execute a code string and enhances the \_call\_ method

ScipyTest the test manager for NumPy which was extracted originally from the SciPy code base. This test manager makes it easy to add unit-tests to a package simply by creating a tests sub-directory with files named test\_<module>.py.

These test files should then define sub-classes of ScipyTestCase (or unittest.TestCase) named "test\*". These classes should then define functions named "test\*" or "bench\*" or "check\*" that contain the actual unit-tests. The first keyword argument should specify the level above which this test should be run.

To run the tests execute ScipyTest(<package>).test(level=1, verbosity=1) which will run all tests above the given level using the given verbosity. Here <package> can be either a string or a previously imported module. You can get the

level and verbosity arguments from sys.argv using ScipyTest(<package>).run() with -v or -verbosity and -l or -level as command-line arguments.

```
set_local_path (reldir=", level=1)
```

prepend local directory (+ reldir) to sys.path. The caller is responsible for removing this path using restore\_path().

```
set_package_path (level=1)
```

prepend package directory to sys.path. This should be called from a test\_file.py that satisfies the tree structure: <somepath>/<somedir>/test\_file.py. The, the first existing path name from the list <somepath>/build/lib.<platform>-<version>, <somepath>/.. is pre-pended to sys.path. The caller is responsible for removing this path using restore\_path().

# restore\_path ()

Remove the first entry from sys.path.

```
assert_equal (actual, desired, err_msg=", verbose=1)
```

Raise an assertion error if the two items are not equal. Automatically calls assert\_array\_equal if actual or desired is an ndarray.

```
assert_almost_equal (actual, desired, decimal=7, err_msg=", verbose=1)
```

Raise an assertion error if the two items are not equal within decimal places. Automatically calls assert\_array\_almost\_equal if actual or desired is an ndarray.

```
assert_approx_equal (actual, desired, significant=7, err_msg=", verbose=1)
```

Raise an assertion error if the two items are not equal to within the given significant digits. Does not work on arrays.

```
assert_array_equal (x, y, err_msg=")
```

Raise an error if the two arrays x and y are not equal at every element.

```
assert_array_less (x, y, err_msg=")
```

Raise an error if the two arrays x and y have different shapes or if x is not less than y at every element.

```
assert_array_almost_equal (x, y, decimal=6, err_msg=")
```

Raise an error if x and y are not equal to decimal places at every element.

# jiffies ()

Return a number of 1/100ths of a second that this process has been scheduled in user mode. Implemented using time.time() unless on Linux where the special /proc directory filesystem is used.

# memusage ()

Return the virtual memory size in bytes of the running python. If the operation is not supported on the platform, then return None. This works only on linux for now.

# rand (\*args)

Return an array of random numbers with the given shape using only the standard library random number generator.

# runstring (astr, dict)

Run the given string in the dictionary provided. Functional form for (exec astr in dict) that is useful for the failUnlessRaises method of unittest.TestCase class.

# 11.2 NumPy Distutils

NumPy provides enhanced distutils functionality to make it easier to build and install sub-packages, auto-generate code, and extension modules that use Fortrancompiled libraries. To use features of numpy distutils use the setup command from numpy.distutils.core. A useful Configuration class is also provided in numpy.distutils.misc\_util that can make it easier to construct keyword arguments to pass to the setup function (by passing the dictionary obtained from the todict() method of the class). More information is available in the NumPy Distutils Users Guide in <site-packages>/numpy/doc/DISTUT

# 11.2.1 misc\_util

Configuration (package\_name=None, parent\_name=None, top\_path=None, package\_path=None, \*\*attrs)

Construct a configuration instance for the given package name. If parent\_name is not None, then construct the package as a sub-package of the parent\_name package. If top\_path and package\_path are None then they are assumed equal to the path of the file this instance was created in. The setup.py files in the numpy distribution are good examples of how to use the Configuration instance.

#### self.todict ()

Return a dictionary compatible with the keyword arguments of distutils setup function. Thus, this method may be used as setup(\*\*config.todict()).

# self.get\_distribution ()

Return the distutils distribution object for self.

# self.get\_subpackage (subpackage\_name, subpackage\_path=None)

Return a Configuration instance for the sub-package given. If subpackage\_path is None then the path is assumed to be the local path plus the subpackage\_name. If a setup.py file is not found in the subpackage\_path, then a default configuration is used.

# self.add\_subpackage (subpackage\_name, subpackage\_path=None)

Add a sub-package to the current Configuration instance. This is useful in a setup.py script for adding sub-packages to a package. The sub-package is contained in subpackage\_path / subpackage\_name and this directory may contain a setup.py script or else a default setup (suitable for Python-code-only subpackages) is assumed. If the subpackage\_path is None, then it is assumed to be located in the local path / subpackage\_name.

# self.add\_data\_files (\*files)

Add files to the list of data\_files to be included with the package. The form of each element of the files sequence is very flexible allowing many combinations of where to get the files from the package and where they should ultimately be installed on the system. The most basic usage is for an element of the files argument sequence to be a simple filename. This will cause that file from the local path to be installed to the installation path of the self.name package (package path). The file argument can also be a relative path in which case the entire relative path will be installed into the package directory. Finally, the file can be an absolute path name in which case the file will be found at the absolute path name but installed to the package path.

This basic behavior can be augmented by passing a 2-tuple in as the file argument. The first element of the tuple should specify the relative path (under the package install directory) where the remaining sequence of files should be installed to (it has nothing to do with the file-names in the source distribution). The second element of the tuple is the sequence of files that should be installed. The files in this sequence can be filenames, relative paths, or absolute paths. For absolute paths the file will be

installed in the top-level package installation directory (regardless of the first argument). Filenames and relative path names will be installed in the package install directory under the path name given as the first element of the tuple. An example may clarify:

```
self.add_data_files('foo.dat',
    ('fun', ['gun.dat', 'nun/pun.dat', '/tmp/sun.dat']),
    'bar/cat.dat',
    '/full/path/to/can.dat')
will install these data files to:
    <package install directory>/
    foo.dat
    fun/
        gun.dat
        nun/
        pun.dat
    sun.dat
    bar/
        car.dat
    can.dat
```

where <package install directory> is the package (or sub-package) directory such as '/usr/lib/python2.4/site-packages/mypackage' ('C:\Python2.4\Lib\\site-packages/mypackage/mysubpackage ('C:\Python2.4\Lib\\site-packages\\mysubpackage\\mysubpackage').

An additional feature is that the path to a data-file can actually be a function that takes no arguments and returns the actual path(s) to the data-files. This is useful when the data files are generated while building the package.

#### self.add\_data\_dir (data\_path)

Recursively add files under data\_path to the list of data\_files to be installed (and distributed). The data\_path can be either a relative pathname, or an absolute path-name, or a 2-tuple where the first argument shows wher in the install directory the data directory should be installed to. For example suppose the source directory contains fun/foo.dat and fun/bar/car.dat

```
self.add_data_dir(('fun')
self.add_data_dir(('sun', 'fun'))
self.add_data_dir(('gun', '/full/path/to/fun'))
```

Will install data-files to the locations

```
<package install directory>/
  fun/
    foo.dat
    bar/
    car.dat
sun/
  foo.dat
  bar/
    car.dat
gun/
  foo.dat
  car.dat
```

# self.add\_include\_dirs (\*paths)

Add the given sequence of paths to the beginning of the include\_dirs list. This list will be visable to all extension modules of the current package.

```
self.add_headers (*files)
```

Add the given sequence of files to the beginning of the headers list. By default, headers will be installed under <python-include>/<self.name.replace('.','/')>/ directory. If an item of files is a tuple, then its first argument specifies the actual installation location relative to the <python-include> path.

```
self.add_extension (name, sources, **kw)
```

Create and add an Extension instance to the ext\_modules list. The first argument defines the name of the extension module that will be installed under the self.name package. The second argument is a list of sources. This method also takes the following optional keyword arguments that are passed on to the Extension constructor: include\_dirs, define\_macros, undef\_macros, library\_dirs, libraries, runtime\_library\_dirs, extra\_objects, swig\_opts, depends, language, f2py\_options, module\_dirs, and extra\_info.

The self.paths(...) method is applied to all lists that may contain paths. The extra\_info is a dictionary or a list of dictionaries whose content will be appended to the keyword arguments. The depends list contains paths to files or directories that the sources of the extension module depend on. If any path in the depends list is newer than the extension module, then the module will be rebuilt.

The list of sources may contain functions (called source generators) which must take an extension instance and a build directory as inputs and return a source file or list of source files or None. If None is returned then no sources are generated. If the Extension instance has no sources after processing all source generators, then no extension module is built.

# self.add\_library (name, sources, \*\*build\_info)

Add a library to the list of libraries. Allowed keyword arguments are depends, macros, include\_dirs, extra\_compiler\_args, and f2py\_options. The name is the name of the library to be built and sources is a list of sources (or source generating functions) to add to the library.

# self.add\_scripts (\*files)

Add the sequence of files to the beginning of the scripts list. Scripts will be installed under the cprefix/bin/ directory.

# self.paths (\*paths)

Applies glob.glob(...) to each path in the sequence (if needed) and pre-pends the local\_path if needed. Because this is called on all source lists, this allows wildcard characters to be specified in lists of sources for extension modules and libraries and scripts and allows path-names be relative to the source directory.

### self.get\_config\_cmd ()

Returns the numpy.distutils config command instance.

#### self.get\_build\_temp\_dir ()

Return a path to a temporary directory where temporary files should be placed.

# self.have\_f77c ()

True if a Fortran 77 compiler is available (because a simple Fortran 77 code was able to be compiled successfully).

#### self.have\_f90c ()

True if a Fortran 90 compiler is available (because a simple Fortran 90 code was able to be compiled successfully)

#### self.get\_version ()

Return a version string of the current package or None if the version information could not be detected. This method scans files named \_version\_py, <packagename>\_version.py, version.py, and \_svn\_version\_py for string variables version, \_version\_, and <packagename>\_version, until a version number is found.

# self.make\_svn\_version\_py ()

Appends a data function to the data\_files list that will generate \_\_svn\_version\_\_.py file to the current package directory. This file will be removed from the source directory when Python exits (so that it can be re-generated next time the package is built). This is intended for working with source directories that are in an SVN repository.

# self.make\_config\_py ()

Generate a package \_config\_py file containing system information used during the building of the package. This file is installed to the package installation directory.

# self.get\_info (\*names)

Return information (from system\_info.get\_info) for all of the names in the argument list in a single dictionary.

# get\_numpy\_include\_dirs ()

Return the include directory where the numpy/arrayobject.h and numpy/ufuncobject.h files are found. This should be added to the include\_dirs of any extension module built using NumPy. If numpy.distutils is used to build the extension, then this directory is added automatically.

# dict\_append (d, \*\*kwds)

Add the keyword arguments given as entries in the dictionary provided as the first argument. If the entry is already present, then assume it is a list and extend the list with the keyword value.

#### appendpath (prefix, path)

Platform-independent intelligence for appending path to prefix. It replaces '/' in the prefix and the path with the correct path-separator on the platform ad returns a full path name that will be valid for the platform.

#### allpath (name)

Convert a '/' separated pathname to one using the platform's path separator.

#### dot\_join (\*args)

Converts a sequence of string arguments to a string joined by '.' (removing any empty strings).

# **generate\_config\_py** (extension, build\_dir)

A suitable function that can be used in a source list. This constructs a python file that contains system\_info information used during building the package. Generally easier to use a Configuration instance and the config.make\_config\_py() method.

```
get_cmd (cmdname, _cache={})
```

Returns an instance of the distutils command object named cmdname if the setup distribution instance has been initialized. Caches the result in \_cache[cmdname] and gets it from there if present.

# terminal\_has\_colors ()

Tries to determine if the stdout terminal can be written to using ANSI colors. Returns 1 if it can be determined that ANSI colors are acceptable or 0 if not.

```
red_text (s)
```

green\_text (s)

yellow\_text (s)

blue\_text (s)

#### cyan\_text (s)

If terminal\_has\_colors() is true, then these commands return a string with the necessary codes prepended to display the given string argument in the specified color on an ANSI terminal. If terminal\_has\_colors() is false, then these functions simply return the input argument.

```
cyg2win32 (path)
```

Convert a cygwin path beginning with /cygdrive to a standard win32 path name.

```
all_strings (lst)
```

Return True if all items in the input list are string objects otherwise return False.

```
has_f_sources (sources)
```

Return True if any of the source files listed in the input argument are Fortran files because its name matches against the compiled regular expression fortran\_ext\_match.

```
has_cxx_sources (sources)
```

Return True if any of the source files listed in the input argument are C++ files because its name matches against the compiled regular expression cxx\_ext\_match.

# filter\_sources (sources)

From the provided list of sources, return four lists of filenames containing C, C++, Fortran, and Fortran 90 module sources respectively. The compiled regular expressions used in this search (which are also available in the misc\_util module) are cxx\_ext\_match, fortran\_ext\_match, f90\_ext\_match, and f90\_module\_name\_match.

# get\_dependencies (sources)

Scan the files in the sources list for include statements.

# is\_local\_src\_dir (directory)

Return True if the provided directory is the local current working directory.

# get\_ext\_source\_files (ext)

Get sources and any include files in the same directory from an Extension instance.

# get\_script\_files (scripts)

Returns the list scripts with all non-string arguments removed.

#### 11.2.2 Other modules

#### system\_info.get\_info (name)

For the given string representing a particular resource, return a dictionary that is compatible with the distutils.setup keyword arguments. If this is an empty dictionary, then the requested resource is not available. Some of the names that can be checked are 'lapack\_opt', 'blas\_opt', 'fft\_opt', 'fftw', 'fftw3', 'fftw2', 'djbfft', 'numpy', 'numarray', 'boost\_python', 'agg2', 'wx', 'gdk', 'xft', 'freetype2'.

# system\_info.get\_standard\_file (filename)

Return a list of length 0 to 3 containing the full-path filenames for the filename provided. The filename is searched for in three places in the following order 1) the system-wide location which is the directory that the system-info file is located in; 2) the directory specified by the environment variable HOME; and 3) the current local directory.

**cpuinfo.cpu** an instance of a cpuinfo class that defines methods for checking various aspects of the cpu. The info attribute is a list of length (# of CPUs). Each entry is a dictionary providing technical information about that CPU.

# log.set\_verbosity (level)

Set the distutils logging threshold and return the previously stored value. The level is an integer that corresponds to distutils.log thresholds: -1 <-> ER-ROR, 0 <-> WARN, 1 <-> INFO, and 2 <-> DEBUG.

#### exec\_command

exec\_command (command, execute\_in=", use\_shell=None, use\_tee=None,
 \_with\_python=1, \*\*env)

Return (status, output) of the executed command. The command input is a string of executable and arguments. The output contains both stderr and stdout messges. If execute\_in is given, then change to the provided directory prior to executing the command and afterwords restore to the current directory. On NT, and DOS systems the returned status is correct for external commands. However, wild cards will not work for non-posix systems.

# splitcmdline (line)

Inverse of ''.join(sys.argv)

find\_executable (exe, path=None)

Return full path of an executable using information from the PATH environment variable. Equivalent to the POSIX 'which' command.

#### get\_pythonexe ()

Return the full path to the python executable with some fixes for nt and dos to replace pythonw with python if it is encountered. A basic wrapper around sys.excecutable.

# 11.3 Conversion of .src files

NumPy distutils supports automatic conversion of source files named <somefile>.src. This facility can be used to maintain very similar code blocks requiring only simple changes between blocks. During the build phase of setup, if a template file named <somefile>.src is encountered, a new file named <somefile> is constructed from the template and placed in the build directory to be used instead. Two forms of template conversion are supported. The first form occurs for files named named <file>.ext.src where ext is a recognized Fortran extension (f, f90, f95, f77, for, ftn, pyf). The second form is used for all other cases.

#### 11.3.1 Fortran files

This template converter will replicate all **function** and **subroutine** blocks in the file with names that contain '<...>' according to the rules in '<...>'. The number of comma-separated words in '<...>' determines the number of times the block is repeated. What these words are indicates what that repeat rule, '<...>', should be replaced with in each block. All of the repeat rules in a block must contain the same number of comma-separated words indicating the number of times that block should be repeated. If the word in the repeat rule needs a comma, leftarrow, or rightarrow, then prepend it with a backslash '\'. If a word in the repeat rule matches '\\<index>' then it will be replaced with the <index>-th word in the same repeat specification. There are two forms for the repeat rule: named and short.

#### 11.3.1.1 Named repeat rule

A named repeat rule is useful when the same set of repeats must be used several times in a block. It is specified using <rule1=item1, item2, item3,..., itemN>, where N is the number of times the block should be repeated. On each repeat of the block, the entire expression, '<...>' will be replaced first with item1, and then with item2, and so forth until N repeats are accomplished. Once a named repeat specification has been introduced, the same repeat rule may be used in the current block by referring only to the name (i.e. <rule1>.

#### 11.3.1.2 Short repeat rule

A short repeat rule looks like <item1, item2, item3, ..., itemN>. The rule specifes that the entire expression, '<...>' should be replaced first with item1, and then with item2, and so forth until N repeats are accomplished.

#### 11.3.1.3 Pre-defined names

The following predefined named repeat rules are available:

- $\bullet$  cprefix=s,d,c,z>
- $\bullet$  <\_c=s,d,c,z>
- <\_t=real, double precision, complex, double complex>
- <ftype=real, double precision, complex, double complex>
- <ctype=float, double, complex\_float, complex\_double>

•  $\langle \text{ctypereal=float, double, } \backslash \langle 0, \rangle \rangle$ 

#### 11.3.2 Other files

Non-Fortran files use a separate syntax for defining template blocks that should be repeated using a variable expansion similar to the named repeat rules of the Fortran-specific repeats. The template rules for these files are:

- 1. "/\*\*begin repeat" on a line by itself marks the beginning of a segment that should be repeated.
- 2. Named variable expansions are defined using #name=item1, item2, item3, ..., itemN# and placed on successive lines. These variables are replaced in each repeat block with corresponding word. All named variables in the same repeat block must define the same number of words.
- 3. In specifying the repeat rule for a named variable, item\*N is short-hand for item, item, ..., item repeated N times. In addition, parenthesis in combination with \*N can be used for grouping several items that should be repeated. Thus, #name=(item1, item2)\*4# is equivalent to #name=item1, item2, item1, item2, item1, item2, item1, item2#
- 4. "\*/" on a line by itself marks the end of the the variable expansion naming. The next line is the first line that will be repeated using the named rules.
- 5. Inside the block to be repeated, the variables that should be expanded are specified as @name@.
- 6. "/\*\*end repeat\*\*/" on a line by itself marks the previous line as the last line of the block to be repeated.

Part II

C-API

# Chapter 12

# New Python Types and C-Structures

NumPy provides a C-API to enable users to extend the system and get access to the array object for use in other routines. The best way to truly understand the C-API is to read the source code. If you are unfamiliar with (C) source code, however, this can be a daunting experience at first. Be assured that the task becomes easier with practice, and you may be surpised at how simple the C-code can be to understand. Even if you don't think you can write C-code from scratch, it is much easier to understand and modify already-written source code then create it de novo.

Python extensions are especially straightforward to understand because they all have a very similar structure. Admittedly, NumPy is not a trivial extension to Python, and may take a little more snooping to grasp. This is especially true because of the code-generation techniques, which simplify maintenance of very similar code, but can make the code a little less readable to beginners. Still, with a little persistance, the code can be opened to your understanding. It is my hope, that this guide to the C-API can assist in the process of becoming familiar with the compiled-level work that can be done with NumPy in order to squeeze that last bit of necessary speed out of your code.

Several new types are defined in the C-code. Most of these are accessible from Python, but a few are not exposed due to their limited use. Every new Python type has an associated PyObject \* with an internal structure that includes a pointer to a "method table" that defines how the new object behaves in Python. When you receive a Python object into C code, you always get a pointer to a PyObject structure. Because a PyObject structure is very generic and defines only PyObject\_HEAD, by itself it is not very interesting. However, different objects contain

more details after the PyObject\_HEAD (but you have to cast to the correct type to access them — or use accessor functions or macros).

# 12.1 New Python Types Defined

Python types are the functional equivalent in C of classes in Python. By constructing a new Python type you make available a new object for Python. The ndarray object is an example of a new type defined in C. New types are defined in C by two basic steps:

- 1. creating a C-structure (usually named Py<Name>Object) that is binary-compatible with the PyObject structure itself but holds the additional information needed for that particular object;
- 2. populating the **PyTypeObject** table (pointed to by the ob\_type member of the PyObject structure) with pointers to functions that implement the desired behavior for the type.

Instead of special method names which define behavior for Python classes, there are "function tables" which point to functions that implement the desired results. Since Python 2.2, the PyTypeObject itself has become dynamic which allows C types that can be "sub-typed" from other C-types in C, and sub-classed in Python. The children types inherit the attributes and methods from their parent(s).

There are two major new types: the ndarray (PyArray\_Type) and the ufunc (PyUFunc\_Type). Additional types play a supportive role: the PyArrayIter\_Type, the PyArrayMultiIter\_Type, and the PyArrayDescr\_Type. The PyArrayIter\_Type is the type for a flat iterator for an ndarray (the object that is returned when getting the flat attribute). The PyArrayMultiIter\_Type is the type of the object returned when calling broadcast(). It handles iteration and broadcasting over a collection of nested sequences. Also, the PyArrayDescr\_Type is the data-type-descriptor type whose instances describe the data. Finally, there are 21 new scalar-array types which are new Python scalars corresponding to each of the fundamental data types available for arrays. An additional 10 other types are place holders that allow the array scalars to fit into a hierarchy of actual Python types.

# 12.1.1 PyArray\_Type

The Python type of the ndarray is **PyArray\_Type**. In C, every ndarray is a pointer to a **PyArrayObject** structure. The ob\_type member of this structure contains a pointer to the **PyArray\_Type** typeobject.

The **PyArrayObject** C-structure contains all of the required information for an array. All instances of an ndarray (and its subclasses) will have this structure. For future compatibility, these structure members should normally be accessed using the provided macros.

```
typedef struct PyArrayObject {
    PyObject_HEAD
    char *data;
    int nd;
    intp *dimensions;
    intp *strides;
    PyObject *base;
    PyArray_Descr *descr;
    int flags;
    PyObject *weakreflist;
} PyArrayObject;
```

- PyObject\_HEAD This is needed by all Python objects. It consists of (at least) a reference count member (ob\_refcnt) and a pointer to the typeobject (ob\_type). (Other elements may also be present if Python was compiled with special options see Include/object.h in the Python source tree for more information). The ob\_type member points to a Python type object.
- data A pointer to the first element of the array. This pointer can (and normally should) be recast to the data type of the array.
- nd An integer providing the number of dimensions for this array. When nd is 0, the array is sometimes called a rank-0 array. Such arrays have undefined dimensions and strides and cannot be accessed. MAX\_DIMS is the largest number of dimensions for any array.
- dimensions An array of integers providing the shape in each dimension as long as nd≥1. The integer is always large enough to hold a pointer on the platform, so the dimension size is only limited by memory.
- **strides** An array of integers providing for each dimension the number of bytes that must be skipped to get to the next element in that dimension.
- base This member is used to hold a pointer to another Python object that is related to this array. There are two use cases: 1) If this array does not own its own memory, then base points to the Python object that owns it (perhaps another array object), 2) If this array has the UPDATEIFCOPY flag set, then this

array is a working copy of a "misbehaved" array. As soon as this array is deleted, the array pointed to by base will be updated with the contents of this array.

descr A pointer to a data-type descriptor object (see below). The data-type descriptor object is an instance of a new builtin type which allows a generic description of memory. There is a descriptor structure for each data type supported. This descriptor structure contains useful information about the type as well as a pointer to a table of function pointers to implement specific functionality.

flags Flags indicating how the memory pointed to by data is to be interpreted. Possible flags are CONTIGUOUS, FORTRAN, OWNDATA, ALIGNED, WRITEABLE, and UPDATEIFCOPY.

weakreflist This member allows array objects to have weak references (using the weakref module).

## 12.1.2 PyArrayDescr\_Type

The PyArrayDescr\_Type is the builtin type of the data-type-descriptor objects used to describe how the bytes comprising the array are to be interpreted. There are 21 statically-defined PyArray\_Descr objects for the builtin data-types. While these participate in reference counting, their reference count should never reach zero. There is also a dynamic table of user-defined PyArray\_Descr objects that is also maintained. Once a data-type-descriptor object is "registered" it should never be deallocated either. The function PyArray\_DescrFromType(...) can be used to retrieve a PyArray\_Descr object from an enumerated typenumber (either builtin or user-defined). The format of the structure that lies at the heart of the PyArrayDescr\_Type is.

```
typedef struct {
    PyObject_HEAD
    PyTypeObject *typeobj;
    char kind;
    char type;
    char byteorder;
    char hasobject;
    int type_num;
    int elsize;
    int alignment;
```

```
PyArray_ArrayDescr *subarray;
PyObject *fields;
PyArray_ArrFuncs *f;
} PyArray_Descr;
```

- **typeobj** Pointer to a typeobject that is the corresponding Python type for the elements of this array. For the builtin types, this points to the corresponding array scalar. For user-defined types, this should point to a user-defined typeobject.
- **kind** A character code indicating the kind of array (using the array interface type-string notation).
- type A traditional character code indicating the data type.
- byteorder A character indicating the byte-order: '>' (big-endian), '<' (little-endian), '=' (native), '|' (irrelevant, ignore). All builtin data-types have byteorder '='.
- **hasobject** Non-zero if this data-type descriptor describes memory where any part of it
- **type\_num** A number that uniquely identifies the data type.
- **elsize** For data types that are always the same size (such as long), this holds the size of the data type. For flexible data types where different arrays can have a different elementsize, this should be 0.
- alignment A number providing alignment information for this data type. Specifically, it shows how far from the start of a 2-element structure (whose first element is a char), the compiler places an item of this type: offsetof(struct {char c; type v;}, v)
- subarray If this is non-NULL, then this data-type descriptor is a C-style contiguous array of another data-type descriptor. In other-words, each element that this descriptor describes is actually an array of some other base descriptor. This is most useful as the data-type descriptor for a field in another data-type descriptor. The fields member should be NULL if this is non-NULL (the fields member of the base descriptor can be non-NULL however). The PyArray\_ArrayDescr structure is defined using

```
typedef struct {
    PyArray_Descr *base;
```

```
PyObject *shape;
} PyArray_ArrayDescr;
```

The elements of this structure are:

base The data-type-descriptor object of the base-type.

**shape** The shape (always C-style contiguous) of the sub-array as a Python tuple.

- fields If this is non-NULL, then this data-type-descriptor has fields described by a Python dictionary whose keys are names (and also titles if given) and whose values are tuples that describe the fields. Recall that a data-type-descriptor always describes a fixed-length set of bytes. A field is a named sub-region of that total, fixed-length collection. A field is described by a tuple composed of another data-type-descriptor and a byte offset. Optionally, the tuple may contain a title which is normally a Python string. These tuples are placed in this dictionary keyed by name (and also title if given).
- f A pointer to a structure containing functions that the type needs to implement internal features. These functions are not the same thing as the universal functions (ufuncs) described later. Their signatures can vary arbitrarily. Not all of these function pointers must be defined for a given type. The required members are nonzero, copyswap, copyswapn, setitem, getitem, and cast. These are assumed to be non-NULL and NULL entries will cause a program crash. The other functions may be NULL which will just mean reduced functionality for that data-type. (Also, the nonzero function will be filled in with a default function if it is NULL when you register a user-defined data-type).

```
typedef struct {
    PyArray_VectorUnaryFunc *cast[PyArray_NTYPES];
    PyArray_GetItemFunc *getitem;
    PyArray_SetItemFunc *setitem;
    PyArray_CopySwapNFunc *copyswapn;
    PyArray_CopySwapFunc *copyswap;
    PyArray_CompareFunc *compare;
    PyArray_ArgFunc *argmax;
    PyArray_DotFunc *dotfunc;
    PyArray_DotFunc *scanfunc;
    PyArray_FromStrFunc *fromstr;
    PyArray_NonzeroFunc *nonzero;
```

```
PyArray_FillFunc *fill;
PyArray_FillWithScalarFunc *fillwithscalar;
PyArray_SortFunc *sort[PyArray_NSORTS];
PyArray_ArgSortFunc *argsort[PyArray_NSORTS];
PyArray_ScalarKindFunc *scalarkind;
int **cancastscalarkindto;
int *cancastto
}
PyArray_ArrFuncs;
```

The concept of a behaved segment is used in the description of the function pointers. A behaved segment is one that is aligned and in native machine byte-order for the data-type. The **nonzero**, **copyswap**, **copyswapn**, **getitem**, and **setitem** functions can all deal with mis-behaved arrays. The other functions require behaved memory segments.

```
cast (void) (void* from, void* to, intp n, void* fromarr, void* toarr)
```

An array of function pointers to cast from the current type to all of the other builtin types. Each function casts a contiguous, aligned, and notswapped buffer pointed at by from to a contiguous, aligned, and notswapped buffer pointed at by to The number of items to cast is given by n, and the arguments from arr and toarr are interpreted as PyArrayObjects for flexible arrays to get itemsize information.

```
getitem (PyObject*) (void* data, void* arr)
```

A pointer to a function that returns a standard Python object from a single element of the array object arr pointed to by data. This function deals with "misbehaved" (misaligned and/or swapped) arrays correctly.

```
setitem (int) (PyObject* item, void* data, void* arr)
```

A pointer to a function that sets a Python object item into the array, arr, at the position pointed to by data. This function deals with "misbehaved" arrays. If successful, a zero is returned, otherwise, a negative number is returned.

```
copyswapn (void) (void* dest, void* src, intp n, int swap, void *arr)
copyswap (void) (void* dest, void* src, int swap, void *arr)
```

These members are both pointers to functions to copy data from src to dest and swap if indicated. The value of itemsize is only used for flexible (STRING, UNICODE, and VOID) arrays. The second function copies a single value, while the first loops over n values which are assumed to be

contiguous. These functions can deal with misbehaved src data. If src is NULL then no copy is performed. If swap is 0, then no byteswapping occurs. It is assumed that dest and src do not overlap. If they overlap, then use memmove(...) first followed by copyswap(n) with NULL valued src.

compare (int) (const void\* d1, const void\* d2, void\* arr)

A pointer to a function that compares two elements of the array, arr, pointed to by d1 and d2. This function requires behaved arrays. The return value is 1 if \*d1 > \*d2, 0 if \*d1 == \*d2, and -1 if \*d1 < \*d2. The array object arr is used to retrieve itemsize information for flexible arrays.

argmax (int) (void\* data, intp n, intp\* max\_ind, void\* arr)

A pointer to a function that retrieves the index of the largest of n elements in arr beginning at the element pointed to by data. This function requires that the memory segment be contiguous and behaved. The return value is always 0. The index of the largest element is returned in max\_ind.

dotfunc (void) (void\* ip1, intp is1, void\* ip2, intp is2, void\* op, intp n,
 void\* arr)

A pointer to a function that multiplies two n-length sequences together, adds them, and places the result in element pointed to by op of arr. The start of the two sequenes are pointed to by ip1 and ip2. To get to the next element in each sequence requires a jump of is1 and is2 bytes, respectively. This function requires behaved (though not necessarily contiguous) memory.

scanfunc (int) (FILE\* fd, void\* ip ,void\* sep ,void\* arr)

A pointer to a function that scans (scanf style) one element of the corresponding type from the file descriptor fd into the array memory pointed to by ip. The array is assumed to be behaved. If sep is not NULL, then a separator string is also scanned from the file before returning. The last argument arr is the array to be scanned into. A 0 is returned if the scan is successful. A negative number indicates something went wrong: -1 means the end of file was reached before the separator string scould be scanned, -4 means that the end of file was reached before the element could be scanned, and -3 means that the element could not be interpreted from the format string. Requires a behaved array.

fromstr (int) (char\* str, void\* ip, char\*\* endptr, void\* arr)

A pointer to a function that converts the string pointed to by str to one element of the corresponding type and places it in the memory location pointed to by ip. After the conversion is completed, \*endptr points to the rest of the string. The last argument arr is the array into which ip points (needed for variable-size data-types). Returns 0 on success or -1 on failure. Requires a behaved array.

nonzero (Bool) (void\* data, void\* arr)

A pointer to a function that returns TRUE if the item of arr pointed to by data is nonzero. This function can deal with misbehaved arrays.

fill (void) (void\* data, intp length, void\* arr)

A pointer to a function that fills a contiguous array of given length with data. The first two elements of the array must already be filled-in. From these two values, a delta will be computed and the values from item 3 to the end will be computed by repeatedly adding this computed delta. The data buffer must be well-behaved.

fillwithscalar (void)(void\* buffer, intp length, void\* value, void\* arr)

A pointer to a function that fills a contiguous buffer of the given length with a single scalar value whose address is given. The final argument is the array which is needed to get the itemsize for variable-length arrays.

sort (int) (void \*start, intp length, void \*arr)

An array of function pointers to a particular sorting algorithms. A particular sorting algorithm is obtained using a key (so far PyArray\_QUICKSORT, PyArray\_HEAPSORT, and PyArray\_MERGESORT are defined). These sorts are done in-place assuming contiguous and aligned data.

argsort (int) (void \*start, intp \*result, intp length, void \*arr)

An array of function pointers to sorting algorithms for this data type. The same sorting algorithms as for sort are available. The indices producing the sort are returned in result (which must be initialized with indices 0 to length-1 inclusive).

#### castdict

Either NULL or a dictionary containing low-level casting functions for user-defined data-types. Each function is wrapped in a PyCObject\* and keyed by the data-type number.

scalarkind (PyArray\_SCALARKIND) (PyArrayObject\* arr)

A function to determine how scalars of this type should be interpreted. The argument is NULL or a 0-dimensional array containing the data (if that is needed to determine the kind of scalar). The return value must be of type PyArray\_SCALARKIND.

#### cancastscalarkindto

Either NULL or an array of PyArray\_NSCALARKINDS pointers. These pointers should each be either NULL or a pointer to an array of integers (terminated by PyArray\_NOTYPE) indicating data-types that a scalar of this data-type of the specified kind can be cast to safely (this usually means without losing precision).

#### cancastto

Either NULL or an array of integers (terminated by PyArray\_NOTYPE) indicated data-types that this data-type can be cast to safely (this usually means without losing precision).

The PyArray\_Type typeobject implements many of the features of Python objects including the tp\_as\_number, tp\_as\_sequence, tp\_as\_mapping, and tp\_as\_buffer interfaces. The rich comparison (tp\_richcompare) is also used along with newstyle attribute lookup for methods (tp\_methods) and properties (tp\_getset). The PyArray\_Type can also be sub-typed.



#### TIP

The tp\_as\_number methods use a generic approach to call whatever function has been registered for handling the operation. The function PyNumeric\_SetOps(..) can be used to register functions to handle particular mathematical operations (for all arrays). When the umath module is imported, it sets the numeric operations for all arrays to the corresponding ufuncs.

The tp\_str and tp\_repr methods can also be altered using PyS-tring\_SetStringFunction(...).

# 12.1.3 PyUFunc\_Type

The ufunc object is implemented by creation of the PyUFunc\_Type. It is a very simple type that implements only basic getattribute behavior, printing behavior, and has call behavior which allows these objects to act like functions. The basic idea behind the ufunc is to hold a reference to fast 1-dimensional (vector) loops for each data type that supports the operation. These one-dimensional loops all have

the same signature and are the key to creating a new ufunc. They are called by the generic looping code as appropriate to implement the N-dimensional function. There are also some generic 1-d loops defined for floating and complexfloating arrays that allow you to define a ufunc using a single scalar function (e.g. atanh).

The core of the ufunc is the PyUFuncObject which contains all the information needed to call the underlying C-code loops that perform the actual work. It has the following structure.

```
typedef struct {
    PyObject_HEAD
    int nin;
    int nout;
    int nargs;
    int identity;
    PyUFuncGenericFunction *functions;
    void **data;
    int ntypes;
    int check_return;
    char *name;
    char *types;
    char *doc:
    void *ptr;
    PyObject *obj;
    PyObject *userloops;
} PyUFuncObject;
```

PyObject\_HEAD required for all Python objects.

**nin** The number of input arguments.

**nout** The number of output arguments.

nargs The total number of arguments (nin+nout). This must be less than MAX\_ARGS.

**identity** Either PyUFunc\_One, PyUFunc\_Zero, or PyUFunc\_None to indicate the identity for this operation. It is only used for a reduce-like call on an empty array.

```
functions (void) (char** args, intp* dims, intp* steps, void* extradata)
```

An array of function pointers — one for each data type supported by the ufunc.

This is the vector loop that is called to implement the underlying function

- dims[0] times. The first argument, args, is an array of nargs pointers to behaved memory. Pointers to the data for the input arguments are first, followed by the pointers to the data for the output arguments. How many bytes must be skipped to get to the next element in the sequence is specified by the corresponding entry in the steps array. The last argument allows the loop to receive extra information. This is commonly used so that a single, generic vector loop can be used for multiple functions. In this case, the actual scalar function to call is passed in as extradata. The size of this function pointer array is ntypes.
- data Extra data to be passed to the 1-d vector loops or NULL if no extra-data is needed. This C-array must be the same size (*i.e.* ntypes) as the functions array. NULL is used if extra-data is not needed. Several C-API calls for UFuncs are just 1-d vector loops that make use of this extra data to receive a pointer to the actual function to call.
- **ntypes** The number of supported data types for the ufunc. This number specifies how many different 1-d loops (of the builtin data types) are available.
- **check\_return** Obsolete and unused. However, it is set by the corresponding entry in the main ufunc creation routine: PyUFunc\_FromFuncAndData(...).
- **name** A string name for the ufunc. This is used dynamically to build the \_doc\_ attribute of ufuncs.
- types An array of nargs×ntypes sizeof(char)-bit type\_numbers which contains the type signature for the function for each of the supported (builtin) data types. For each of the ntypes functions, the corresponding set of type numbers in this array shows how the args argument should be interpreted in the 1-d vector loop. These type numbers do not have to be the same type and mixed-type ufuncs are supported.
- **doc** Documentation for the ufunc. Should not contain the function signature as this is generated dynamically when <u>\_doc\_</u> is retrieved.
- **ptr** Any dynamically allocated memory. Currently, this is used for dynamic ufuncs created from a python function to store room for the types, data, and name members.
- **obj** For ufuncs dynamically created from python functions, this member holds a reference to the underlying Python function.

userloops A dictionary of user-defined 1-d vector loops (stored as CObject ptrs) for user-defined types. A loop may be registered by the user for any user-defined type. It is retrieved by type number. User defined type numbers are always larger than PyArray\_USERDEF.

## 12.1.4 PyArrayIter\_Type

This is an iterator object that makes it easy to loop over an N-dimensional array. It is the object returned from the flat attribute of an ndarray. It is also used extensively throughout the implementation internals to loop over an N-dimensional array. The tp\_as\_mapping interface is implemented so that the iterator object can be indexed (using 1-d indexing), and a few methods are implemented through the tp\_methods table. This object implements the next method and can be used anywhere an iterator can be used in Python.

The C-structure corresponding to an object of PyArrayIter\_Type is the PyArrayIter\_Object. The PyArrayIterObject is used to keep track of a pointer into an N-dimensional array. It contains associated information used to quickly march through the array. The pointer can be adjusted in three basic ways: 1) advance to the "next" position in the array in a C-style contiguous fashion, 2) advance to an arbitrary N-dimensional coordinate in the array, and 3) advance to an arbitrary one-dimensional index into the array. The members of the PyArrayIterObject structure are used in these calculations. Iterator objects keep their own dimension and strides information about an array. This can be adjusted as needed for "broadcasting," or to loop over only specific dimensions.

```
typedef struct {
    PyObject_HEAD
    int
          nd_m1;
    intp
          index;
    intp
          size;
          coordinates [MAX_DIMS];
    intp
          dims_m1 [MAX_DIMS];
    intp
    intp
          strides [MAX_DIMS];
          backstrides [MAX_DIMS];
    intp
          factors[MAX_DIMS];
    PyArrayObject *ao;
    char
          *dataptr;
    Bool
          contiguous;
} PyArrayIterObject;
```

 $nd_m1$  N-1 where N is the number of dimensions in the underlying array.

index The current 1-d index into the array.

**size** The total size of the underlying array.

**coordinates** An N-dimensional index into the array.

dims\_m1 The size of the array minus 1 in each dimension.

strides The strides of the array. How many bytes needed to jump to the next element in each dimension.

backstrides How many bytes needed to jump from the end of a dimension back to its beginning. Note that backstrides[k]=strides[k]\*dims\_m1[k], but it is stored here as an optimization.

factors This array is used in computing an N-d index from a 1-d index. It contains needed products of the dimensions.

ao A pointer to the underlying ndarray this iterator was created to represent.

dataptr This member points to an element in the ndarray indicated by the index.

**contiguous** This flag is true if the underlying array is CONTIGUOUS. It is used to simplify calculations when possible.

How to use an array iterator on a C-level is explained more fully in later sections. Typically, you do not need to concern yourself with the internal structure of the iterator object, and merely interact with it through the use of the macros PyArray\_ITER\_NEXT(it), PyArray\_ITER\_GOTO(it, dest), or PyArray\_ITER\_GOTO(it, dest).

index). All of these macros require the argument it to be a PyArrayIterObject\*.

# 12.1.5 PyArrayMultiIter\_Type

This type provides an iterator that encapsulates the concept of broadcasting. It allows N arrays to be broadcast together so that the loop progresses in C-style contiguous fashion over the broadcasted array. The corresponding C-structure is the **PyArrayMultiIterObject** whose memory layout must begin any object, obj, passed in to the **PyArray\_Broadcast(obj)** function. Broadcasting is performed by adjusting array iterators so that each iterator represents the broadcasted shape and size, but has its strides adjusted so that the correct element from the array is used at each iteration.

```
typedef struct {
    PyObject_HEAD
    int numiter;
    intp size;
    intp index;
    int nd;
    int p dimensions[MAX_DIMS];
    PyArrayIterObject *iters[MAX_DIMS];
}
```

**PyObject\_HEAD** Needed at the start of every Python object (holds reference count and type identification).

**numiter** The number of arrays that need to be broadcast to the same shape.

**size** The total broadcasted size.

index The current (1-d) index into the broadcasted result.

**nd** The number of dimensions in the broadcasted result.

dimensions The shape of the broadcated result (only nd slots are used).

**iters** An array of iterator objects that holds the iterators for the arrays to be broadcast together. On return, the iterators are adjusted for broadcasting.

# 12.1.6 PyArrayFlags\_Type

When the flags attribute is retrieved from Python, a special builtin object of this type is constructed. This special type makes it easier to work with the different flags by accessing them as attributes or by accessing them as if the object were a dictionary with the flag names as entries.

# 12.1.7 ScalarArrayTypes

There is a Python type for each of the different builtin data types that can be present in the array Most of these are simple wrappers around the corresponding data type in C. The C-names for these types are **Py**<TYPE>**ArrType\_Type** where <TYPE> can be

Bool, Byte, Short, Int, Long, LongLong, UByte, UShort, UInt, ULong, ULongLong, Float, Double, LongDouble, CFloat, CDouble, CLongDouble, String, Unicode, Void, and Object.

These type names are part of the C-API and can therefore be created in extension C-code. There is also a **PyIntpArrType\_Type** and a **PyUIntpArrType\_Type** that are simple substitutes for one of the integer types that can hold a pointer on the platform. The structure of these scalar objects is not exposed to C-code. The function **PyArray\_ScalarAsCtype(...)** can be used to extract the C-type value from the array scalar and the function **PyArray\_Scalar(...)** can be used to construct an array scalar from a C-value.

## 12.2 Other C-Structures

A few new C-structures were found to be useful in the development of NumPy. These C-structures are used in at least one C-API call and are therefore documented here. The main reason these structures were defined is to make it easy to use the Python ParseTuple C-API to convert from Python objects to a useful C-Object.

## 12.2.1 PyArray\_Dims

Very useful when shape and/or strides information is supposed to be interpreted. The structure is

```
typedef struct {
    intp *ptr;
    int len;
} PyArray_Dims;
```

The members of this structure are

**ptr** A pointer to a list of (intp) integers which usually represent array shape or array strides.

len The length of the list of integers. It is safe to access ptr[0] to ptr[len-1].

# 12.2.2 PyArray\_Chunk

This is equivalent to the buffer object structure in Python up to the ptr member. On 32-bit platforms (*i.e.* if **SIZEOF\_INT**==**SIZEOF\_INTP**) the len member also matches an equivalent member of the buffer object (where it is defined as an int). It is useful to represent a generic single-segment chunk of memory.

```
typedef struct {
    PyObject_HEAD
```

```
PyObject *base;
void *ptr;
intp len;
int flags;
} PyArray_Chunk;
```

The members are

**PyObject\_HEAD** Necessary for all Python objects. Included here so that the PyArray\_Chunk structure matches that of the buffer object (at least to the len member).

**base** The Python object this chunk of memory comes from. Needed so that memory can be accounted for properly.

ptr A pointer to the start of the single-segment chunk of memory.

**len** The length of the segment in bytes.

flags Any data flags (e.g. WRITEABLE) that should be used to interpret the memory.

## 12.2.3 PyArrayInterface

The PyArrayInterface structure is defined so that NumPy and other extension modules can use the rapid array interface protocol. The <code>\_\_array\_struct\_\_</code> method of an object that supports the rapid array interface protocol should return a PyCObject that contains a pointer to a PyArrayInterface structure with the relevant details of the array. After the new array is created, the attribute should be DECREF'd which will free the PyArrayInterface structure. Remember to INCREF the object (whose <code>\_array\_struct\_\_</code> attribute was retrieved) and point the base member of the new PyArrayObject to this same object. In this way the memory for the array will be managed correctly.

```
typedef struct {
    int version;
    int nd;
    char typekind;
    int itemsize;
    int flags;
    intp *shape;
    intp *strides;
```

```
void *data;
} PyArrayInterface;
```

**version** the integer 2 as a sanity check.

nd the number of dimensions in the array.

typekind A character indicating what kind of array is present according to the typestring convention with 't' -> bitfield, 'b' -> boolean, 'i' -> signed integer, 'u' -> unsigned integer, 'f' -> floating point, 'c' -> complex floating point, 'O' -> object, 'S' -> string, 'U' -> unicode, 'V' -> void.

itemsize the number of bytes each item in the array requires.

flags any of the bits CONTIGUOUS (1), FORTRAN (2), ALIGNED (0x100), NOTSWAPPED (0x200), or WRITEABLE (0x400) to indicate something about the data. The ALIGNED, CONTIGUOUS, and FORTRAN flags can actually be determined from the other parameters.

**shape** An array containing the size of the array in each dimension.

**strides** An array containing the number of bytes to jump to get to the next element in each dimension.

data A pointer to the first element of the array.

## 12.2.4 Internally used structures

Internally, the code uses some additional Python objects primarily for memory management. These types are not accessible directly from Python, and are not exposed to the C-API. They are included here only for completeness and assistance in understanding the code.

#### 12.2.4.1 PyUFuncLoopObject

A loose wrapper for a C-structure that contains the information needed for looping. This is useful if you are trying to understand the ufunc looping code. The **PyUFuncLoopObject** is the associated C-structure. It is defined in the **ufuncobject.h** header.

#### 12.2.4.2 PyUFuncReduceObject

A loose wrapper for the C-structure that contains the information needed for reducelike methods of ufuncs. This is useful if you are trying to understand the reduce, accumulate, and reduce-at code. The **PyUFuncReduceObject** is the associated C-structure. It is defined in the **ufuncobject.h** header.

#### 12.2.4.3 PyArrayMapIter\_Type

Advanced indexing is handled with this Python type. It is simply a loose wrapper around the C-structure containing the variables needed for advanced array indexing. The associated C-structure, **PyArrayMapIterObject**, is useful if you are trying to understand the advanced-index mapping code. It is defined in the **arrayobject.h** header. This type is not exposed to Python and could be replaced with a C-structure. As a type it takes advantage of reference-counted memory management.

# Chapter 13

# Complete API

# 13.1 Configuration defines

When NumPy is built, a configuration file is constructed and placed as config.h in the NumPy include directory. This configuration file ensures that specific macros are defined and defines other macros based on whether or not your system has certain features. It is included by the arrayobject.h file.

#### 13.1.1 Guaranteed to be defined

The **SIZEOF**\_<CTYPE> constants are defined so that size of information is available to the pre-processor.

CHAR\_BIT The number of bits of a char. The char is the unit of all size of definitions

**SIZEOF\_SHORT** sizeof(short)

**SIZEOF\_INT** sizeof(int)

**SIZEOF\_LONG** sizeof(long)

**SIZEOF\_LONG\_LONG** sizeof(longlong) where longlong is defined appropriately on the platform (A macro defines **SIZEOF\_LONGLONG** as well.)

SIZEOF\_PY\_LONG\_LONG

**SIZEOF\_FLOAT** sizeof(float)

**SIZEOF\_DOUBLE** sizeof(double)

**SIZEOF\_LONG\_DOUBLE** sizeof(longdouble) (A macro defines **SIZEOF\_LONGDOUBLE** as well.)

SIZEOF\_PY\_INTPTR\_T Size of a pointer on this platform (sizeof(void \*)) (A macro defines SIZEOF\_INTP as well.)

#### 13.1.2 Possible defines

These defines will cause the compilation to ignore compatibility code that is placed in NumPy and use the system code instead. If they are not defined, then the system does not have that capability.

HAVE\_LONGDOUBLE\_FUNCS System has C99 long double math functions.

HAVE\_FLOAT\_FUNCS System has C99 float math functions.

**HAVE\_INVERSE\_HYPERBOLIC** System has inverse hyperbolic functions: asinh, acosh, and atanh.

**HAVE\_INVERSE\_HYPERBOLIC\_FLOAT** System has C99 float extensions to inverse hyperbolic functions: asinhf, acoshf, atanhf

HAVE\_INVERSE\_HYPERBOLIC\_LONGDOUBLE System has C99 long double extensions to inverse hyperbolic functions: asinhl, acoshl, atanhl.

HAVE\_ISNAN System has an isnan function.

**HAVE\_ISINF** System has an isinf function.

**HAVE\_LOG1P** System has log1p function:  $\log(x+1)$ .

**HAVE\_EXPM1** System has expm1 function:  $\exp(x) - 1$ .

# 13.2 Array Data Types

The standard array can have 21 different data types (and has some beginning support for adding your own types). These data types all have an enumerated type, an enumerated typecharacter, and a corresponding array scalar Python type object (placed in a hierarchy). There are also standard C typedefs to make it easier to manipulate elements of the given data type. For the numeric types, there are also bit-width equivalent C typedefs and named typenumbers that make it easier to select the precision desired.



#### WARNING

The names for the types in c code follows c naming conventions more closely. The Python names for these types follow Python conventions. Thus, PyArray\_FLOAT picks up a 32-bit float in C, but "float\_" in python corresponds to a 64-bit double. The bit-width names can be used in both Python and C for clarity.

#### 13.2.1 Enumerated Types

There is a list of enumerated types defined providing the basic 21 data types plus some useful generic names. Whenever the code requires a type number, one of these enumerated types is requested. The types are all called **PyArray\_**<NAME> where <NAME> can be

BOOL, BYTE, UBYTE, SHORT, USHORT, INT, UINT, LONG, ULONG, LONGLONG, ULONGLONG, FLOAT, DOUBLE, LONG-DOUBLE, CFLOAT, CDOUBLE, CLONGDOUBLE, OBJECT, STRING, UNICODE, VOID

NTYPES, NOTYPE, USERDEF

The various character codes indicating certain types are also part of an enumerated list. References to type characters (should they be needed at all) should always use these enumerations. The form of them is **PyArray\_**<NAME>**LTR** where <NAME> can be

BOOL, BYTE, UBYTE, SHORT, USHORT, INT, UINT, LONG, ULONG, LONGLONG, ULONGLONG, FLOAT, DOUBLE, LONG-DOUBLE, CFLOAT, CDOUBLE, CLONGDOUBLE, OBJECT, STRING, VOID

INTP, UINTP

GENBOOL, SIGNED, UNSIGNED, FLOATING, COMPLEX

The latter group of <NAME>s corresponds to letters used in the array interface typestring specification.

#### 13.2.2 **Defines**

13.2.2.1 Max and min values for integers

MAX\_INT<br/><br/>bits>

#### $\mathbf{MAX\_UINT}{<} \mathrm{bits}{>}$

#### MIN\_INT<br/>bits>

These are defined for <br/> <br/> = 8, 16, 32, 64, 128, and 256 and provide the maximum (minimum) value of the corresponding (unsigned) integer type. Note: the actual integer type may not be available on all platforms (i.e. 128-bit and 256-bit integers are rare).

#### $MIN_{-} < type >$

This is defined for <type> = BYTE, SHORT, INT, LONG, LONGLONG, INTP

#### $MAX_{\text{-}} < \text{type} >$

This is defined for all defined for <type> = BYTE, UBYTE, SHORT, USHORT, INT, UINT, LONG, ULONG, LONGLONG, ULONGLONG, INTP, UINTP

#### 13.2.2.2 Number of bits in data types

All SIZEOF\_<CTYPE> constants have corresponding BITSOF\_<CTYPE> constants defined. The BITSOF\_<CTYPE> constants provide the number of bits in the data type. Specifically, the available <CTYPE>s are

BOOL, CHAR, SHORT, INT, LONG, LONGLONG, FLOAT, DOUBLE, LONGDOUBLE

#### 13.2.2.3 Bit-width references to enumerated typenums

All of the numeric data types (integer, floating point, and complex) have constants that are defined to be a specific enumerated type number. Exactly which enumerated type a bit-width type refers to is platform dependent. In particular, the constants available are **PyArray**\_<NAME><BITS> where <NAME> is **INT**, **UINT**, **FLOAT**, **COMPLEX** and <BITS> can be 8, 16, 32, 64, 80, 96, 128, 160, 192, 256, and 512. Obviously not all bit-widths are available on all platforms for all the kinds of numeric types. Commonly 8-, 16-, 32-, 64-bit integers; 32-, 64-bit floats; and 64-, 128-bit complex types are available.

#### 13.2.2.4 Integer that can hold a pointer

The constants **PyArray\_INTP** and **PyArray\_UINTP** refer to an enumerated integer type that is large enough to hold a pointer on the platform. Index arrays

should always be converted to **PyArray\_INTP**, because the dimension of the array is of type intp.

#### 13.2.3 C-type names

There are standard variable types for each of the numeric data types and the bool data type. Some of these are already available in the C-specification. You can create variables in extension code with these types.

If a conflict arises between one or more of these names and other library code, you may define **PY\_ARRAY\_TYPES\_PREFIX** as a prefix to all of these names

#### 13.2.3.1 Boolean

**Bool** unsigned char; The constants FALSE and TRUE are also defined.

#### 13.2.3.2 (Un)Signed Integer

Unsigned versions of the integers can be defined by pre-pending a 'u' to the front of the integer name.

- (u)byte (unsigned) char
- (u)short (unsigned) short
- (u)int (unsigned) int
- (u)long (unsigned) long int
- (u)longlong (unsigned long long int)
- (u)intp (unsigned) Py\_intptr\_t (an integer that is the size of a pointer on the platform).

#### 13.2.3.3 (Complex) Floating point

- (c)float float
- (c)double double
- (c)longdouble long double

complex types are structures with **.real** and **.imag** members (in that order).

#### 13.2.3.4 Bit-width names

There are also typedefs for signed integers, unsigned integers, floating point, and complex floating point types of specific bit-widths. The available type names are

where <br/>bits> is the number of bits in the type and can be **8**, **16**, **32**, **64**, 128, and 256 for integer types; 16, **32**, **64**, 80, 96, 128, and 256 for floating-point types; and 32, **64**, **128**, 160, 192, and 512 for complex-valued types. Which bit-widths are available is platform dependent. The bolded bit-widths are usually available on all platforms.

## 13.2.4 Printf Formatting

For help in printing, the following strings are defined as the correct format specifier in printf and related commands.

# LONGLONG\_FMT, ULONGLONG\_FMT, INTP\_FMT, UINTP\_FMT LONGDOUBLE\_FMT

# 13.3 Array API

# 13.3.1 Array structure and data access

These macros all access the PyArrayObject structure members. The input argument, obj, can be any PyObject\* that is directly interpretable as a PyArrayObject\* (any instance of the PyArray\_Type and its sub-types).

PyArray\_DATA (void\*) (PyObject\* obj)

PyArray\_BYTES (char\*) (PyObject\* obj)

These two macros are similar and obtain the pointer to the data-buffer for the array. The first macro can (and should be) assigned to a particular pointer where the second is for generic processing. If you have not guaranteed a contiguous and/or aligned array then be sure you understand how to access the data in the array to avoid memory and/or alignment problems.

PyArray\_DIMS (intp\*) (PyObject\* arr)

PyArray\_STRIDES (intp\*) (PyObject\* arr)

PyArray\_DIM (intp) (PyObject\* arr, int n)

Return the shape in the n<sup>th</sup> dimension.

PyArray\_STRIDE (intp) (PyObject\* arr, int n)

Return the stride in the n<sup>th</sup> dimension.

PyArray\_BASE (PyObject\*) (PyObject\* arr)

PyArray\_DESCR (PyArray\_Descr\*) (PyObject\* arr)

PyArray\_FLAGS (int) (PyObject\* arr)

PyArray\_ITEMSIZE (int) (PyObject\* arr)

Return the itemsize for the elements of this array.

PyArray\_TYPE (int) (PyObject\* arr)

Return the (builtin) typenumber for the elements of this array.

PyArray\_GETITEM (PyObject \*) (PyObject\* arr, void\* itemptr)

Get a Python object from the ndarray, arr, at the location pointed to by itemptr.

Return NULL on failure.

PyArray\_SETITEM (int) (PyObject\* arr, void\* itemptr, PyObject\* obj)

Convert obj and place it in the ndarray, arr, at the place pointed to by itemptr. Return -1 if an error occurs or 0 on success.

PyArray\_SIZE (intp) (PyObject\* arr)

Returns the total size (in number of elements) of the array.

PyArray\_Size (intp) (PyObject\* obj)

Returns 0 if obj is not a sub-class of bigndarray. Otherwise, returns the total number of elements in the array. Safer version of PyArray\_SIZE(obj).

 ${\bf PyArray\_NBYTES} \ ({\tt intp}) \ ({\tt PyObject*} \ {\tt arr})$ 

Returns the total number of bytes consumed by the array.

#### 13.3.1.1 Data access

These functions and macros provide easy access to elements of the ndarray from C. These work for all arrays. You may need to take care when accessing the data in the array, however, if it is not in machine byte-order, misaligned, or not writeable. In other words, be sure to respect the state of the flags unless you know what you are doing, or have previously guaranteed an array that is writeable, aligned, and in machine byte-order using PyArray\_FromAny. If you wish to handle all types of arrays, the copyswap function for each type is useful for handling misbehaved arrays. Some platforms (e.g. Solaris) do not like misaligned data and will crash if you de-reference a misaligned pointer. Other platforms (e.g. x86 Linux) will just work more slowly with misaligned data.

PyArray\_GetPtr (void\*) (PyArrayObject\* aobj, intp\* ind)

Return a pointer to the data of the ndarray, aobj, at the N-dimensional index given by the c-array, ind, (which must be at least aobj->nd in size). You may want to typecast the returned pointer to the data type of the ndarray.

PyArray\_GETPTR1 (void\*) (PyObject\* obj, <intp> i)

PyArray\_GETPTR2 (void\*) (PyObject\* obj, <intp> i, <intp> j)

PyArray\_GETPTR3 (void\*) (PyObject\* obj, <intp> i, <intp> j, <intp> k)

Quick, inline access to the element at the given coordinates in the ndarray, obj, which must have respectively 1, 2, 3, or 4 dimensions (this is not checked). The corresponding i, j, k, and l coordinates can be any integer but will be interpreted as intp. You may want to typecast the returned pointer to the data type of the ndarray.

#### 13.3.2 Creating arrays

#### 13.3.2.1 From scratch

PyArray\_NewFromDescr (PyObject\*) (PyTypeObject\* subtype, PyArray\_Descr\* descr, intp\* dims, intp\* strides, void\* data, int flags, PyObject\* obj)

This is the main array creation function. Most new arrays are created with this flexible function. The returned object is an object of Python-type subtype, which must be a subtype of **PyArray\_Type**. The array has **nd** dimensions,

described by dims. The data-type descriptor of the new array is descr. If subtype is not &PyArray\_Type (e.g. a Python subclass of the ndarray), then obj is the object to pass to the \_\_array\_finalize\_\_ method of the subclass. If data is NULL, then new memory will be allocated and flags can be non-zero to indicate a Fortran-style contiguous array. If data is not NULL, then it is assumed to point to the memory to be used for the array and the flags argument is used as the new flags for the array (except the state of OWNDATA and UPDATEIFCOPY flags of the new array will be reset). In addition, if data is non-NULL, then strides can also be provided. If strides is NULL, then the array strides are computed as C-style contiguous (default) or Fortran-style contiguous (flags is nonzero for data=NULL or flags & FORTRAN is nonzero non-NULL data). Any provided dims and strides are copied into newly allocated dimension and strides arrays for the new array object.

PyArray\_New (PyObject\*) (PyTypeObject\* subtype, int nd, intp\* dims, int type\_num, intp\* strides, void\* data, int itemsize, int flags, PyObject\* obj)

This is similar to **PyArray\_DescrNew**(...) except you specify the data-type descriptor with type\_num and itemsize, where type\_num corresponds to a builtin (or user-defined) type. If the type always has the same number of bytes, then itemsize is ignored. Otherwise, itemsize specifies the particular size of this array.



#### WARNING

If data is passed to PyArray\_NewFromDescr or PyArray\_New, this memory must not be deallocated until the new array is deleted. If this data came from another Python object, this can be accomplished using Py\_INCREF on that object and setting the base member of the new array to point to that object. If strides are passed in they must be consistent with the dimensions, the itemsize, and the data of the array.

PyArray\_SimpleNew (PyObject\*) (int nd, intp\* dims, int typenum)

Create a new unitialized array of type, typenum, whose size in each of nd dimensions is given by the integer array, dims. This function cannot be used to create a flexible-type array (no itemsize given).

PyArray\_SimpleNewFromData (PyObject\*) (int nd, intp\* dims, int typenum, void\* data)

- Create an array wrapper around data pointed to by the given pointer. The array flags will have a default that the data area is well-behaved and C-style contiguous.
- - Create a new array with the provided data-type descriptor, descr, of the shape determined by nd and dims.
- PyArray\_FILLWBYTE (PyObject\* obj, int val)
  - Fill the array pointed to by obj—which must be a (subclass of) bigndarray—with the contents of val (evaluated as a byte).
- PyArray\_Zeros (PyObject\*) (int nd, intp\* dims, PyArray\_Descr\* dtype, int
  fortran)
  - Construct a new nd-dimensional array with shape given by dims and data type given by dtype. If fortran is non-zero, then a fortran-order array is created. Fill the memory with zeros (or the 0 object if dtype corresponds to PyArray\_OBJECT).
- PyArray\_Empty (PyObject\*) (int nd, intp\* dims, PyArray\_Descr\* dtype, int
  fortran)
  - Construct a new nd-dimensional array with shape given by dims and data type given by dtype. If fortran is non-zero, then a fortran-order array is created.

    The array is uninitialized unless the data type corresponds to PyArray\_OBJECT in which case the array is filled with Py\_None.
- PyArray\_Arange (PyObject\*) (double start, double stop, double step, int typenum)
  - Construct a new 1-dimensional array of data-type, typenum, that ranges from start to stop (exclusive) in increments of step. Equivalent to arange(start, stop, step, typenum).
- PyArray\_ArangeObj (PyObject\*) (PyObject\* start, PyObject\* stop, PyObject\*
  step, PyArray\_Descr\* descr)
  - Construct a new 1-dimensional array of data-type determined by descr, that ranges from start to stop (exclusive) in increments of step. Equivalent to arange(start, stop, step, typenum).

#### 13.3.2.2 From other objects

PyArray\_FromAny (PyObject\*) (PyObject\* op, PyArray\_Descr\* dtype, int min\_depth, int max\_depth, int requirements, PyObject\* context)

This is the main function used to obtain an array from any nested sequence, or object that exposes the array interface, op. The parameters allow specification of the required type, the minimum (min\_depth) and maximum (max\_depth) number of dimensions acceptable, and other requirements for the array. The dtype argument needs to be a PyArray\_Descr structure indicating the desired data-type (including required byteorder). The dtype argument may be NULL, indicating that any data-type (and byteorder) is acceptable. If you want to use NULL for the dtype and ensure the array is not swapped then use PyArray\_CheckFromAny. A value of 0 for either of the depth parameters causes the parameter to be ignored. Any of the following array flags can be added (e.q. using |) to get the requirements argument. If your code can handle general arrays, then requirements may be 0. Also, if op is not already an array (or does not expose the array interface), then a new array will be created (and filled from op using the sequence protocol). The new array will have DEFAULT\_FLAGS. The context argument is passed to the \_\_array\_ method of op and is only used if the array is constructed that way.

**CONTIGUOUS** Make sure the returned array is C-style contiguous

**FORTRAN** Make sure the returned array is Fortran-style contiguous.

**ALIGNED** Make sure the returned array is aligned on proper boundaries for its data type. An aligned array has the data pointer and every strides factor as a multiple of the alignment factor for the data-type-descriptor.

WRITEABLE Make sure the returned array can be written to.

**ENSURECOPY** Make sure a copy is made of op. If this flag is not present, data is not copied if it can be avoided.

**ENSUREARRAY** Make sure the result is a base-class ndarray or bigndarray. By default, if **op** is an instance of a subclass of the bigndarray, an instance of that same subclass is returned. If this flag is set, an ndarray object will be returned instead.

**FORCECAST** Force a cast to the output type even if it cannot be done safely. Without this flag, a data cast will occur only if it can be done safely, otherwise an error is reaised.

**UPDATEIFCOPY** If op is already an array, but does not satisfy the requirements, then a copy is made (which will satisfy the requirements).

If this flag is present and a copy (of an object that is already an array) must be made, then the corresponding UPDATEIFCOPY flag is set in the returned copy and op is made to be read-only. When the returned copy is deleted (presumably after your calculations are complete), its contents will be copied back into op and the op array will be made writeable again. If op is not writeable to begin with, then an error is raised. If op is not already an array, then this flag has no effect.

BEHAVED\_FLAGS ALIGNED | WRITEABLE

CARRAY\_FLAGS CONTIGUOUS | BEHAVED\_FLAGS

CARRAY\_FLAGS\_RO CONTIGUOUS | ALIGNED

FARRAY\_FLAGS FORTRAN | BEHAVED\_FLAGS

FARRAY\_FLAGS\_RO FORTRAN | ALIGNED

**DEFAULT\_FLAGS** CARRAY\_FLAGS

IN\_ARRAY CONTIGUOUS | ALIGNED

IN\_FARRAY FORTRAN | ALIGNED

INOUT\_ARRAY CONTIGUOUS | WRITEABLE | ALIGNED

INOUT\_FARRAY FORTRAN | WRITEABLE | ALIGNED

 $\begin{array}{c} \mathbf{OUT\_ARRAY} \ \ \mathbf{CONTIGUOUS} \ | \ \mathbf{WRITEABLE} \ | \ \mathbf{ALIGNED} \ | \ \mathbf{UPDATEIF-COPY} \\ \end{array}$ 

OUT\_FARRAY FORTRAN | WRITEABLE | ALIGNED | UPDATEIF-COPY

PyArray\_CheckFromAny (PyObject\*) (PyObject\* op, PyArray\_Descr\* dtype,
int min\_depth, int max\_depth, int requirements, PyObject\* context)

Nearly identical to **PyArray\_FromAny**(...) except **requirements** can contain NOTSWAPPED (over-riding the specification in **dtype**), and if requirements contains ENSURECOPY, then DEFAULT\_FLAGS will be automatically OR'd to the requirements as well.

NOTSWAPPED Make sure the returned array has a data-type descriptor that is in machine byte-order, over-riding any specification in the dtype argument. Normally, the byte-order requirement is determined by the dtype argument. If this flag is set and the dtype argument does not indicate a machine byte-order descriptor (or is NULL and the object is already an array with a data-type descriptor that is not in machine byte-order), then a new data-type descriptor is created and used with its byte-order field set to native.

#### BEHAVED\_NS\_FLAGS ALIGNED | WRITEABLE | NOTSWAPPED

PyArray\_FromArray (PyObject\*) (PyArrayObject\* op, PyArray\_Descr\* newtype, int requirements)

Special case of **PyArray\_FromAny** for when op is already an array but it needs to be of a specific newtype (including byte-order) or has certain requirements.

#### PyArray\_FromStructInterface (PyObject\*) (PyObject\* op)

Returns an ndarray object from a Python object that exposes the \_array\_struct\_ method and follows the array interface protocol. If the object does not contain this method then a borrowed reference to Py\_NotImplemented is returned.

#### PyArray\_FromInterface (PyObject\*) (PyObject\* op)

Returns an ndarray object from a Python object that exposes the \_array\_shape\_ and \_array\_typestr\_ methods following the array interface protocol. If the object does not contain one of these method then a borrowed reference to Py\_NotImplemented is returned.

# 

Return an ndarray object from a Python object that exposes the \_array\_ method. The \_array\_ method can take 0, 1, or 2 arguments ([dtype, context]) where context is used to pass information about where the \_array\_ method is being called from (currently only used in ufuncs).

# PyArray\_ContiguousFromAny (PyObject\*) (PyObject\* op, int typenum, int min\_depth, int max\_depth)

This function returns a (C-style) contiguous and behaved function array from any nested sequence or array interface exporting object, op, of (non-flexible) type given by the enumerated typenum, of minimum depth min\_depth, and of maximum depth max\_depth. Equivalent to a call to PyArray\_FromAny with requirements set to DEFAULT\_FLAGS and the type\_num member of the type argument set to typenum.

# **PyArray\_FromObject** (PyObject \*) (PyObject \* op, int typenum, int min\_depth, int max\_depth)

Return an aligned and in native-byteorder array from any nested sequence or array-interface exporting object, op, of a type given by the enumerated typenum. The minimum number of dimensions the array can have is given by min\_depth while the maximum is max\_depth. This is equivalent to a call to PyArray\_FromAny with requirements set to BEHAVED\_FLAGS.

#### PyArray\_EnsureArray (PyObject\* op)

This function **steals a reference** to op and makes sure that op is a base-class ndarray. It special cases array scalars, but otherwise calls **PyArray\_FromAny**(op, NULL, 0, 0, ENSUREARRAY).

PyArray\_FromString (PyObject\*) (char\* string, intp slen, PyArray\_Descr\* dtype, intp num, char\* sep)

Construct a one-dimensional ndarray of a single type from a binary or (ASCII) text string of length slen. The data-type of the array to-be-created is given by dtype. If num is -1, then copy the entire string and return an appropriately sized array, otherwise, num is the number of items to copy from the string. If sep is NULL (or ""), then interpret the string as bytes of binary data, otherwise convert the sub-strings separated by sep to items of data-type dtype. Some data-types may not be readable in text mode and an error will be raised if that occurs. All errors return NULL.

Construct a one-dimensional ndarray of a single type from a binary or text file. The open file pointer is fp, the data-type of the array to be created is given by dtype. This must match the data in the file. If num is -1, then read until the end of the file and return an appropriately sized array, otherwise, num is the number of items to read. If sep is NULL (or ""), then read from the file in binary mode, otherwise read from the file in text mode with sep providing the item separator. Some array types cannot be read in text mode in which case an error is raised.

 $\label{eq:pyArray_FromBuffer} \textbf{PyObject*} \ (\textbf{PyObject*} \ \textbf{buf}, \textbf{PyArray\_Descr*} \ \textbf{dtype}, \ \textbf{intp} \\ \textbf{count}, \ \textbf{intp} \ \textbf{offset})$ 

Construct a one-dimensional ndarray of a single type from an object, buf, that exports the (single-segment) buffer protocol (or has an attribute \_buffer\_ that returns an object that exports the buffer protocol). A writeable buffer will be tried first followed by a read-only buffer. The WRITEABLE flag of the returned array will reflect which one was successful. The data is assumed to start at offset bytes from the start of the memory location for the object.

The type of the data in the buffer will be interpreted depending on the datatype descriptor, dtype. If count is negative then it will be determined from the size of the buffer and the requested itemsize, otherwise, count represents how many elements should be converted from the buffer.

### PyArray\_CopyInto (int) (PyArrayObject\* dest, PyArrayObject\* src)

Copy from the source array, src, into the destination array, dest, performing a data-type conversion if necessary. If an error occurs return -1 (otherwise 0). The total number of elements in dest must be an integer multiple of the total number of elements in src.

## PyArray\_GETCONTIGUOUS (PyArrayObject\*) (PyObject\* op)

If op is already (C-style) contiguous and well-behaved then just return a reference, otherwise return a (contiguous and well-behaved) copy of the array. The parameter op must be a (sub-class of an) ndarray and no checking for that is done.

## PyArray\_FROM\_O (PyObject\*) (PyObject\* obj)

Convert obj to an ndarray. The argument can be any nested sequence or object that exports the array interface. This is a macro form of PyArray\_FromAny using NULL, 0, 0, 0 for the other arguments. Your code must be able to handle any data-type descriptor and any combination of data-flags to use this macro.

## PyArray\_FROM\_OF (PyObject\*) (PyObject\* obj, int requirements)

Similar to PyArray\_FROM\_O except it can take an argument of requirements indicating properties the resulting array must have. Available requirements that can be enforced are CONTIGUOUS, FORTRAN, ALIGNED, WRITE-ABLE, NOTSWAPPED, ENSURECOPY, UPDATEIFCOPY, FORCE-CAST, and ENSUREARRAY. Standard combinations of flags can also be used:

## PyArray\_FROM\_OT (PyObject\*) (PyObject\* obj, int typenum)

Similar to **PyArray\_FROM\_O** except it can take an argument of typenum specifying the type-number the returned array.

PyArray\_FROM\_OTF (PyObject\*) (PyObject\* obj, int typenum, int requirements) Combination of **PyArray\_FROM\_OF** and **PyArray\_FROM\_OT** allowing both a typenum and a flags argument to be provided..

# PyArray\_FROMANY (PyObject\*) (PyObject\* obj, int typenum, int min, int max, int requirements)

Similar to **PyArray\_FromAny** except the data-type is specified using a typenumber. PyArray\_DescrFromType(typenum) is passed directly to PyArray\_FromAny. This macro also adds DEFAULT\_FLAGS to requirements if ENSURECOPY is passed in as requirements.

# 13.3.3 Dealing with types

## 13.3.3.1 General check of Python Type

### PyArray\_Check (op)

Evaluates true if op is a Python object that is a sub-type of **PyArray\_Type**.

## PyArray\_CheckExact (op)

Evaluates true if op is a Python object with type **PyArray\_Type**.

## PyArray\_IsZeroDim (op)

Evaluates true if op is an instance of (a subclass of) **PyArray\_Type** and has 0 dimensions.

## PyArray\_IsScalar (op, cls)

Evaluates true if op is an instance of Py<cls>ArrType\_Type.

## PyArray\_CheckScalar (op)

Evaluates true if op is either an array scalar (an instance of a sub-type of **Py-GenericArr\_Type**), or an instance of (a sub-class of) **PyArray\_Type** whose dimensionality is 0.

## PyArray\_IsPythonScalar (op)

Evaluates true if op is a builtin Python "scalar" object (int, float, complex, str, unicode, long, bool).

## PyArray\_IsAnyScalar (op)

Evaluates true if op is either a Python scalar or an array scalar (an instance of a sub-type of **PyGenericArr\_Type**).

#### 13.3.3.2 Data-type checking

For the typenum macros, the argument is an integer representing an enumerated array data type. For the array type checking macros the argument must be a PyObject\* that can be directly interpreted as a PyArrayObject\*.

PyTypeNum\_ISUNSIGNED (num)

PyDescr\_ISUNSIGNED (descr)

 $\mathbf{PyArray\_ISUNSIGNED} \ (\mathbf{obj})$ 

Type represents an unsigned integer.

PyTypeNum\_ISSIGNED (num)

PyDescr\_ISSIGNED (descr)

PyArray\_ISSIGNED (obj)

Type represents a signed integer.

PyTypeNum\_ISINTEGER (num)

PyDescr\_ISINTEGER (descr)

PyArray\_ISINTEGER (obj)

Type represents any integer.

PyTypeNum\_ISFLOAT (num)

PyDescr\_ISFLOAT (descr)

PyArray\_ISFLOAT (obj)

Type represents any floating point number.

PyTypeNum\_ISCOMPLEX (num)

PyDescr\_ISCOMPLEX (descr)

PyArray\_ISCOMPLEX (obj)

Type represents any complex floating point number.

PyTypeNum\_ISNUMBER (num)

PyDescr\_ISNUMBER (descr)

```
PyArray_ISNUMBER (obj)
```

Type represents any integer, floating point, or complex floating point number.

PyTypeNum\_ISSTRING (num)

PyDescr\_ISSTRING (descr)

PyArray\_ISSTRING (obj)

Type represents a string data type.

PyTypeNum\_ISPYTHON (num)

PyDescr\_ISPYTHON (descr)

PyArray\_ISPYTHON (obj)

Type represents an enumerated type corresponding to one of the standard Python scalar (bool, int, float, or complex).

PyTypeNum\_ISFLEXIBLE (num)

PyDescr\_ISFLEXIBLE (descr)

PyArray\_ISFLEXIBLE (obj)

Type represents one of the flexible array types (STRING, UNICODE, or VOID).

PyTypeNum\_ISUSERDEF (num)

PyDescr\_ISUSERDEF (descr)

PyArray\_ISUSERDEF (obj)

Type represents a user-defined type.

PyTypeNum\_ISEXTENDED (num)

PyDescr\_ISEXTENDED (descr)

PyArray\_ISEXTENDED (obj)

Type is either flexible or user-defined.

PyTypeNum\_ISOBJECT (num)

PyDescr\_ISOBJECT (descr)

PyArray\_ISOBJECT (obj)

Type represents Object data type.

PyTypeNum\_ISBOOL (num)

PyDescr\_ISBOOL (descr)

PyArray\_ISBOOL (obj)

Type represents Boolean data type.

## PyArray\_ISNOTSWAPPED (m)

Evaluates true if the data area of the ndarray  ${\tt m}$  is in machine byte-order according to its descriptor.

 $\textbf{PyArray\_EquivTypes} \hspace{0.1cm} \texttt{(Bool)} \hspace{0.1cm} \texttt{(PyArray\_Descr* type1, PyArray\_Descr* type2)}$ 

Return TRUE if type1 and type2 actually represent equivalent types for this platform (the fortran member of each type is ignored). For example, on 32-bit platforms, PyArray\_LONG and PyArray\_INT are equivalent. Otherwise return FALSE.

PyArray\_EquivArrTypes (Bool) (PyArrayObject\* a1, PyArrayObject\* a2)

Return TRUE if a1 and a2 are arrays with equivalent types for this platform.

PyArray\_EquivTypenums (Bool) (int typenum1, int typenum2)

Special case of **PyArray\_EquivTypes(...)** that does not accept flexible data types but may be easier to call.

PyArray\_EquivByteorders (int) (<byteorder> b1, <byteorder> b2)

True if byteorder characters (PyArray\_LITTLE, PyArray\_BIG, PyArray\_NATIVE, PyArray\_IGNORE) are either equal or equivalent as to their specification of a native byte order. Thus, on a little-endian machine PyArray\_LITTLE and PyArray\_NATIVE are equivalent where they are not equivalent on a bigendian machine.

#### 13.3.3.3 Converting data types

PyArray\_Cast (PyObject\*) (PyArrayObject\* arr, int typenum)

Mainly for backwards compatibility to the Numeric C-API and for simple casts to non-flexible types. Return a new array object with the elements of arr cast to the data-type typenum which must be one of the enumerated types and not a flexible type.

## PyArray\_CastToType (Py0bject\*) (PyArrayObject\*arr, PyArray\_Descr\*type)

Return a new array of the type specified, casting the elements of arr as appropriate. The fortran member of the type structure will be respected in producing the strides of the output array.

## PyArray\_CastTo (int) (PyArrayObject\* out, PyArrayObject\* in)

Cast the elements of the array in into the array out. The output array should be writeable, have an integer-multiple of the number of elements in the input array (more than one copy can be placed in out), and have a data type that is one of the builtin types. Returns 0 on success and -1 if an error occurs.

# PyArray\_GetCastFunc (PyArray\_VectorUnaryFunc\*) (PyArray\_Descr\* from, int totype)

Return the low-level casting function to cast from the given descriptor to the builtin type number. If no casting function exists return NULL and set an error. Using this function instead of direct access to from->f->cast will allow support of any user-defined casting functions added to a descriptors casting dictionary.

## PyArray\_CanCastSafely (int) (int fromtype, int totype)

Returns non-zero if an array of data type fromtype can be cast to an array of data type totype without losing information. An exception is that 64-bit integers are allowed to be cast to 64-bit floating point values even though this can lose precision on large integers so as not to proliferate the use of long doubles without explict requests. Flexible array types are not checked according to their lengths with this function.

## PyArray\_CanCastTo (int) (PyArray\_Descr\* fromtype, PyArray\_Descr\* totype)

Returns non-zero if an array of data type fromtype (which can include flexible types) can be cast safely to an array of data type totype (which can include flexible types). This is basically a wrapper around PyArray\_CanCastSafely with additional support for size checking if fromtype and totype are PyArray\_STRING or PyArray\_UNICODE.

#### PyArray\_ObjectType (int) (PyObject\* op, int mintype)

This function is useful for determining a common type that two or more arrays can be converted to. It only works for non-flexible array types as no itemsize information is passed. The mintype argument represents the minimum type

acceptable, and op represents the object that will be converted to an array. The return value is the enumerated typenumber that represents the data-type that op should have.

# PyArray\_ArrayType (void) (PyObject\*op, PyArray\_Descr\* mintype, PyArray\_Descr\* outtype)

This function works similarly to **PyArray\_ObjectType**(...) except it handles flexible arrays. The mintype argument can have an itemsize member and the outtype argument will have an itemsize member at least as big but perhaps bigger depending on the object op.

# PyArray\_ConvertToCommonType (PyArrayObject\*\*) (PyObject\* op, int\* n)

Convert a sequence of Python objects contained in op to an array of ndarrays each having the same data type. The type is selected based on the typenumber (larger type number is chosen over a smaller one) ignoring objects that are only scalars. The length of the sequence is returned in n, and an n-length array of PyArrayObject pointers is the return value (or NULL if an error occurs). The returned array must be freed by the caller of this routine (using PyDataMem\_FREE) and all the array objects in it DECREF'd or a memory-leak will occur. The example template-code below shows a typically usage.

## PyArray\_Zero (char\*) (PyArrayObject\* arr)

A pointer to newly created memory of size arr->itemsize that holds the representation of 0 for that type. The returned pointer, ret, **must be freed** using **PyDataMem\_FREE**(ret) when it is not needed anymore.

```
PyArray_One (char*) (PyArrayObject* arr)
```

A pointer to newly created memory of size arr->itemsize that holds the representation of 1 for that type. The returned pointer, ret, **must be freed** using **PyDataMem\_FREE**(ret) when it is not needed anymore.

### PyArray\_ValidType (int) (int typenum)

Returns TRUE if typenum represents a valid typenumber (builtin or user-defined or character code). Otherwise, this function returns FALSE.

### 13.3.3.4 New data types

## PyArray\_InitArrFuncs (void) (PyArray\_ArrFuncs\* f)

Initialize all function pointers and members to NULL.

## PyArray\_RegisterDataType (int) (PyArray\_Descr\* dtype)

Register a data-type as a new user-defined data type for arrays. The type must have most of its entries filled in. This is not always checked and errors can produce segfaults. In particular, the typeobj member of the dtype structure must be filled with a Python type that has a fixed-size element-size that corresponds to the elsize member of dtype. Also the "f" member must have the required functions: nonzero, copyswap, copyswapn, getitem, setitem, and cast (some of the cast functions may be NULL if no support is desired). To avoid confusion, you should choose a unique character typecode but this is not enforced and not relied on internally.

A user-defined type number is returned that uniquely identifies the type. A pointer to the new structure can then be obtained from **PyArray\_DescrFromType** using the returned type number. A -1 is returned if an error occurs. If this dtype has already been registered (checked only by the address of the pointer), then return the previously-assigned type-number.

# PyArray\_RegisterCastFunc (int) (PyArray\_Descr\* descr, int totype, PyArray\_VectorUnaryF castfunc)

Register a low-level casting function, castfunc, to convert from the data-type, descr, to the given data-type number, totype. Any old casting function is over-written. A 0 is returned on success or a -1 on failure.

# PyArray\_RegisterCanCast (int) (PyArray\_Descr\* descr, int totype, PyArray\_SCALARKIND scalar)

Register the data-type number, totype, as castable from data-type object, descr, of the given scalar kind. Use scalar = PyArray\_NOSCALAR to register that an array of data-type descr can be cast safely to a data-type whose type\_number is totype.

### 13.3.3.5 Special functions for PyArray\_OBJECT

## PyArray\_INCREF (int) (PyArrayObject\* op)

Used for arrays of Python objects, op, to increment the reference count of every object in the array. A -1 is returned if an error occurs, otherwise 0 is returned.

### PyArray\_XDECREF (int) (PyArrayObject\* op)

Used for arrays of Python objects, op, to decrement the reference count of every object in the array. Normal return value is 0. A -1 is returned if an error occurs.

## PyArray\_FillObjectArray (void) (PyArrayObject\* arr, PyObject\* obj)

Fill a newly created **PyArray\_OBJECT** array, **arr**, with a single value, **obj**. No checking is performed but **arr** must be of data-type **PyArray\_OBJECT** and be single-segment and uninitialized (no previous objects in position). Use **PyArray\_DECREF**(arr) if you need to decrement all the items in the object array prior to calling this function.

## 13.3.4 Array flags

## 13.3.4.1 Basic Array Flags

A bigndarray can have a data segment that is not a simple contiguous chunk of well-behaved memory you can manipulate. It may not be aligned with word boundaries (very important on some platforms). It might have its data in a different byte-order than the machine recognizes. It might not be writeable. It might be in Fortan-contiguous order. The array flags are used to indicate what can be said about data associated with an array.

**CONTIGUOUS** The data area is in C-style contiguous order (last index varies the fastest).

**FORTRAN** The data area is in Fortran-style contiguous order (first index varies the fastest).

**OWNDATA** The data area is owned by this array.

**ALIGNED** The data area is aligned appropriately (for all strides).

WRITEABLE The data area can be written to.

Notice that the above flags are are defined so that a new array has these flags as TRUE.

**UPDATEIFCOPY** The data area represents a (well-behaved) copy whose information should be transferred back to the original when this array is deleted.

### 13.3.4.2 Combinations of array flags

BEHAVED\_FLAGS ALIGNED | WRITEABLE

CARRAY\_FLAGS CONTIGUOUS | BEHAVED\_FLAGS

CARRAY\_FLAGS\_RO CONTIGUOUS | ALIGNED

FARRAY\_FLAGS FORTRAN | BEHAVED\_FLAGS

FARRAY\_FLAGS\_RO FORTRAN | ALIGNED

DEFAULT\_FLAGS CARRAY\_FLAGS

UPDATE\_ALL\_FLAGS CONTIGUOUS | FORTRAN | ALIGNED

#### 13.3.4.3 Flag-like constants

These constants are used in PyArray\_FromAny (and its macro forms) to specify desired properties of the new array.

**FORCECAST** Cast to the desired type, even if it can't be done without losing information.

**ENSURECOPY** Make sure the resulting array is a copy of the original.

**ENSUREARRAY** Make sure the resulting object is an actual ndarray (or bign-darray), and not a sub-class.

**NOTSWAPPED** Only used in PyArray\_CheckFromAny to over-ride the byte-order of the data-type object passed in.

BEHAVED\_NS\_FLAGS ALIGNED | WRITEABLE | NOTSWAPPED

#### 13.3.4.4 Flag checking

For all of these macros m must be an instance of a (subclass of) **PyArray\_Type**, but no checking is done.

#### PyArray\_CHKFLAGS (m, flags)

The first parameter, m, must be a (big)ndarray or subclass. The parameter, flags, should be an integer consisting of bitwise combinations of the possible flags an array can have: CONTIGUOUS, FORTRAN, OWNDATA, ALIGNED, WRITEABLE, UPDATEIFCOPY.

## PyArray\_ISCONTIGUOUS (m)

Evaluates true if  ${\tt m}$  is C-style contiguous.

## PyArray\_ISFORTRAN (m)

Evaluates true if m is Fortran-style contiguous.

## PyArray\_ISWRITEABLE (m)

Evaluates true if the data area of m can be written to

## PyArray\_ISALIGNED (m)

Evaluates true if the data area of m is properly aligned on the machine.

## PyArray\_ISBEHAVED (m)

Evalutes true if the data area of m is **ALIGNED**, and **WRITEABLE** and in machine byte-order according to its descriptor.

## PyArray\_ISBEHAVED\_RO (m)

Evaluates true if the data area of m is ALIGNED and in machine byte-order.

## PyArray\_ISCARRAY (m)

Evaluates true if the data area of m is **CONTIGUOUS**, and PyArray\_ISBEHAVED(m) is true.

## PyArray\_ISFARRAY (m)

Evaluates true if the data area of  ${\tt m}$  is in **FORTRAN** and PyArray\_ISBEHAVED(m) is true.

#### PyArray\_ISCARRAY\_RO (m)

Evaluates true if the data area of m is **CONTIGUOUS** and PyArray ISBEHAVED(m) is true.

## PyArray\_ISFARRAY\_RO (m)

Evaluates true if the data area of m is **FORTRAN** and PyArray\_ISBEHAVED(m) is true.

## PyArray\_ISONESEGMENT (m)

Evaluates true if the data area of m consists of a single (C-style or Fortran style) contiguous segment

### PyArray\_UpdateFlags (void) (PyArrayObject\* arr, int flagmask)

The CONTIGUOUS, ALIGNED, and FORTRAN array flags can be "calculated" from the array object itself. This routine updates one or more of these flags of arr as specified in flagmask by performing the required calculation.

## 13.3.5 Array method alternative API

#### 13.3.5.1 Conversion

PyArray\_GetField (PyObject\*) (PyArrayObject\* self, PyArray\_Descr\* dtype,
 int offset)

Equivalent to self.getfield(dtype, offset). Return a new array of the given type using the data in the current array at a specified offset in bytes. The offset plus the itemsize of the new array type must be less than self->descr->elsize or an error is raised. The same shape and strides as the original array are used. Therefore, this function has the effect of returning a field from a record array. But, it can also be used to select specific bytes or groups of bytes from any array type.

PyArray\_SetField (int) (PyArrayObject\* self, PyArray\_Descr\* dtype, int offset, PyObject\* val)

Equivalent to self.setfield(value, dtype, offset). Set the field starting at offset in bytes and of the given dtype to val. The offset plus dtype->elsize must be less than self->descr->elsize or an error is raised. Otherwise, the val argument is converted to an array and copied into the field pointed to. If necessary, the elements of val are repeated to fill the destination array, But, the number of elements in the destination must be an integer multiple of the number of elements in val.

## PyArray\_Byteswap (PyObject\*) (PyArrayObject\* self, Bool implace)

Equivalent to self.byteswap(inplace). Return an array whose data area is byteswapped. If inplace is non-zero, then do the byteswap inplace and return a reference to self. Otherwise, create a byteswapped copy and leave self unchanged.

## PyArray\_NewCopy (PyObject\*) (PyArrayObject\* old, int fortran)

Equivalent to self.copy(fortran). Make a copy of the old array. The returned array is always ALIGNED and WRITEABLE with data interpreted the same

as the old array. If fortran is 0, then a C-style contiguous array is returned. If fortran is 1, then a Fortran-style contiguous array is returned. If fortran < 0, then the array returned is Fortran-style contiguous if the old is, otherwise, it is C-style contiguous.

### PyArray\_ToList (PyObject\*) (PyArrayObject\* self)

Equivalent to self.tolist(). Return a nested Python list from self.

PyArray\_ToFile (PyObject\*) (PyArrayObject\* self, FILE\* fp, char\* sep, char\*
format)

Write the contents of self to the file pointer fp in C-style contiguous fashion. Write the data as binary bytes if sep is the string "" or NULL. Otherwise, write the contents of self as text using the sep string as the item separator. Each item will be printed to the file. If the format string is not NULL or "", then it is a python print statement format string showing how the items is to be written.

## PyArray\_Dump (int) (PyObject\* self, PyObject\* file, int protocol)

Pickle the object in self to the given file (either a string or a Python file object). If file is a Python string it is considered the name of a file which is opened in binary mode. The given protocol is used (if protocol is negative, then protocol 2 is used). This is a simple wrapper around cPickle.dump(self, file, protocol)

## PyArray\_Dumps (PyObject\*) (PyObject\* self, int protocol)

Pickle the object in self to a string and return the string as a Python string. Use the Pickle protocol provided (or 2 if protocol is negative).

## PyArray\_FillWithScalar (int) (PyArrayObject\* arr, PyObject\* obj)

Fill the array, arr, with the given scalar object, obj. The object is first converted to the data type of arr, and then copied into every location. A -1 is returned if an error occurs, otherwise 0 is returned.

#### PyArray\_View (PyObject\*) (PyArrayObject\* self, PyArray\_Descr\* dtype)

Equivalent to self.view(dtype). Return a new view of the array self as possibly a different data type, dtype. If dtype is NULL, then the returned array will have the same data type as self. The new data type must be consistent with the size of self. Either the itemsizes must be identical, or self must be

single-segment and the total number of bytes must be the same. In the latter case the dimensions of the returned array will be altered in the last (or first for FORTRAN arrays) dimension. The data area of the returned array and self is exactly the same.

## 13.3.5.2 Shape Manipulation

**PyArray\_Newshape** (PyObject\*) (PyArrayObject\* self, PyArray\_Dims\* newshape)

Result will be a new array pointing to the same data as in self, but having a shape given by newshape. Call will succeed as long as self is contiguous or newshape is different from the shape of self only by inserting or deleting ones. The method self.reshape(...) also works for Fortran arrays, and always produces a copy for discontiguous arrays.

PyArray\_Reshape (PyObject\*) (PyArrayObject\* self, PyObject\* shape)

Equivalent to self.reshape(shape) where shape is a sequence. Converts shape to a PyArray\_Dims structure and calls PyArray\_Newshape internally.

PyArray\_Squeeze (PyObject\*) (PyArrayObject\* self)

Equivalent to self.squeeze(). Return a new view of self with all of the length-1 dimensions removed from the shape. Note, matrix objects are always 2dimensional, and thus this call has no effect if self is a matrix sub-class.

PyArray\_SwapAxes (PyObject\*) (PyArrayObject\* self, int a1, int a2)

Equivalent to self.swapaxes(a1, a2). The returned array is a new view of the data in self with the given axes, a1 and a2, swapped.

PyArray\_Resize (PyObject\*) (PyArrayObject\* self, PyArray\_Dims\* newshape,
 int refcheck)

Equivalent to self.resize(newshape, refcheck=refcheck). This function only works on single-segment arrays. It changes the shape of self inplace and will reallocate the memory for self if newshape has a different total number of elements then the old shape. If reallocation is necessary, then self must own its data, have self->base==NULL, have self->weakrefs==NULL, and (unless refcheck is 0) not be referenced by any other array. The new array is returned.

PyArray\_Transpose (PyObject\*) (PyArrayObject\*self, PyArray\_Dims\* permute)

Equivalent to self.transpose(permute). Permute the axes of the ndarray object self according to the data structure permute return the result. If permute is NULL, then the resulting array has its axes reversed. For example if self has shape  $10 \times 20 \times 30$ , and permute.ptr is (0,2,1) the shape of the result is  $10 \times 30 \times 20$ . If permute is NULL, the shape of the result is  $30 \times 20 \times 10$ .

### PyArray\_Flatten (PyObject\*) (PyArrayObject\* self, int fortran)

Equivalent to self.flatten(fortran). Return a 1-d copy of the array. If fortran is positive the elements are scanned out in fortran order (first-dimension varies the fastest). Otherwise, the elements of self are scanned in C-order (last dimension varies the fastest). If fortran is less than 0, then the result of PyArray\_ISFORTRAN(self) is used to determine which order to flatten.

PyArray\_Ravel (PyObject\*) (PyArrayObject\* self, int fortran)

Equivalent to self.ravel(fortran). Same basic functionality as **PyArray\_Flatten**(self, fortran) except if fortran is 0 and self is C-style contiguous, the shape is altered but no copy is performed.

#### 13.3.5.3 Item selection and manipulation

 $\mathbf{PyArrpay\_Take} \ \ (\mathtt{PyObject*}) \ (\mathtt{PyArrayObject*} \\ \mathrm{self}, \ \mathtt{PyObject*} \\ \mathrm{indices}, \ \mathtt{int} \ \mathrm{axis})$ 

Equivalent to self.take(indices, axis) except axis=None in Python is obtained by setting axis=MAX\_DIMS in C. Extract the items from self along the given axis indicated by the integer-valued indices.

PyArray\_Put (PyObject\*) (PyArrayObject\* self, PyObject\* values, PyObject\*
indices)

Equivalent to self.put(values, indices). Put values into self at the corresponding (flattened) indices. If values is too small it will be repeated as necessary.

PyArray\_PutMask (PyObject\*) (PyArrayObject\* self, PyObject\* values, PyObject\*
mask)

Equivalent to self.putmask(values, mask). Place the values in self wherever corresponding positions (using a flattened context) in mask are true. The mask and self arrays must have the same total number of elements. If values is too small, it will be repeated as necessary.

PyArray\_Repeat (PyObject\*) (PyArrayObject\* self, PyObject\* op, int axis)

Equivalent to self.repeat(op, axis). Copy the elements of self, op times along the given axis. Either op is a scalar integer or a sequence of length self->dimensions[axis] indicating how many times to repeat each item along the axis.

### PyArray\_Choose (PyObject\*) (PyArrayObject\* self, PyObject\* op)

Equivalent to self.choose(op). Create a new array by selecting elements from the sequence of arrays in op based on the integer values in self. The arrays must all have the same shape and the entries in self should be between 0 and len(op).

## PyArray\_Sort (PyObject\*) (PyArrayObject\* self, int axis)

Equivalent to self.sort(axis). Return an array with the items of self sorted along axis.

## PyArray\_ArgSort (PyObject\*) (PyArrayObject\* self, int axis)

Equivalent to self.argsort(axis). Return an array of indices such that selection of these indices along the given axis would return a sorted version of self.

## PyArray\_LexSort (PyObject\*) (PyObject\* sort\_keys, int axis)

Given a sequence of arrays (sort\_keys) of the same shape, return an array of indices (similar to PyArray\_ArgSort(...)) that would sort the arrays lexicographically. A lexicographic sort specifies that when two keys are found to be equal, the order is based on comparison of subsequent keys. A merge sort (which leaves equal entries unmoved) is required to be defined for the types. The sort is accomplished by sorting the indices first using the first sort\_key and then using the second sort\_key and so forth. This is equivalent to the lexsort(sort\_keys, axis) Python command. Be sure to understand the order the sort\_keys must be in (reversed from what you might think).

#### PyArray\_SearchSorted (PyObject\*) (PyArrayObject\* self, PyObject\* values)

Equivalent to self.searchsorted(values). Assuming self is a 1-d array in ascending order representing bin boundaries then the output is an array the same shape as values of bin numbers, giving the bin into which each item in values would be placed. No checking is done on whether or not self is in ascending order.

# PyArray\_Diagonal (PyObject\*) (PyArrayObject\* self, int offset, int axis1, int axis2)

Equivalent to self.diagonal(offset, axis1, axis2). Return the offset diagonals of the 2-d arrays defined by axis1 and axis2.

## PyArray\_Nonzero (PyObject\*) (PyArrayObject\* self)

Equivalent to self.nonzero(). Returns a tuple of index arrays that select elements of self that are nonzero. If (nd=PyArray\_NDIM(self))==1, then a single index array is returned. The index arrays have data type PyArray\_INTP. If a tuple is returned (nd\neq1), then its length is nd.

PyArray\_Compress (PyObject\*) (PyArrayObject\* self, PyObject\* condition,
 int axis)

Equivalent to self.compress (condition axis). Return the elements along axis corresponding to elements of condition that are true.

#### 13.3.5.4 Calculation

PyArray\_ArgMax (PyObject\*) (PyArrayObject\* self, int axis)

Equivalent to self.argmax(axis). Return the index of the largest element of self along axis.

PyArray\_ArgMin (PyObject\*) (PyArrayObject\* self, int axis)

Equivalent to self.argmin(axis). Return the index of the smallest element of self along axis.

PyArray\_Max (PyObject\*) (PyArrayObject\* self, int axis)

Equivalent to self.max(axis). Return the largest element of self along the given axis.

PyArray\_Min (PyObject\*) (PyArrayObject\* self, int axis)

Equivalent to self.min(axis). Return the smallest element of self along the given axis.

PyArray\_Ptp (PyObject\*) (PyArrayObject\* self, int axis)

Equivalent to self.ptp(axis). Return the difference between the largest element of self along axis and the smallest element of self along axis.

PyArray\_Mean (PyObject\*) (PyArrayObject\* self, int axis, int rtype)

- Equivalent to self.mean(axis, rtype). Returns the mean of the elements along the given axis, using the enumerated type rtype as the data type to sum in. Default sum behavior is obtained using PyArray\_NOTYPE for rtype.
- - Equivalent to self.trace(offset, axis1, axis2, rtype). Return the sum (using rtype as the data type of summation) over the offset diagonal elements of the 2-d arrays defined by axis1 and axis2 variables. A positive offset chooses diagonals above the main diagonal. A negative offset selects diagonals below the main diagonal.
- - Equivalent to self.clip(min, max). Clip an array, self, so that values larger than max are fixed to max and values less than min are fixed to min.
- PyArray\_Conjugate (PyObject\*) (PyArrayObject\* self)
  - Equivalent to self.conjugate() and self.conj() Return the complex conjugate of self. If self is not of complex data type, then return self with an reference.
- PyArray\_Round (PyObject\*) (PyArrayObject\* self, int decimals)
  - Equivalent to self.round(decimals). Returns the array with elements rounded to the nearest decimal place. The decimal place is defined as the 10<sup>-decimals</sup> digit so that negative decimals cause rounding to the nearest 10's, 100's, etc.
- PyArray\_Std (PyObject\*) (PyArrayObject\* self, int axis, int rtype)
  - Equivalent to self.std(axis, rtype). Return the standard deviation using data along axis converted to data type rtype.
- PyArray\_Sum (PyObject\*) (PyArrayObject\* self, int axis, int rtype)
  - Equivalent to self.sum(axis, rtype). Return 1-d vector sums of elements in self along axis. Perform the sum after converting data to data type rtype.
- PyArray\_CumSum (PyObject\*) (PyArrayObject\* self, int axis, int rtype)
  - Equivalent to self.cumsum(axis, rtype). Return cumulative 1-d sums of elements in self along axis. Perform the sum after converting data to data type rtype.

PyArray\_Prod (PyObject\*) (PyArrayObject\* self, int axis, int rtype)

Equivalent to self.prod(axis, rtype). Return 1-d products of elements in self along axis. Perform the product after converting data to data type rtype.

PyArray\_CumProd (PyObject\*) (PyArrayObject\* self, int axis, int rtype)

Equivalent to self.cumprod(axis, rtype). Return 1-d cumulative products of elements in self along axis. Perform the product after converting data to data type rtype.

PyArray\_All (PyObject\*) (PyArrayObject\* self, int axis)

Equivalent to self.all(axis). Return an array with True elements for every 1-d sub-array of self defined by axis in which all the elements are True.

PyArray\_Any (PyObject\*) (PyArrayObject\* self, int axis)

Equivalent to self.any(axis). Return an array with True elements for every 1-d sub-array of self defined by axis in which any of the elements are True.

#### 13.3.6 Functions

## 13.3.6.1 Array Functions

PyArray\_AsCArray (int) (PyObject\*\* op, void\* ptr, intp\* dims, int nd, int
typenum, int itemsize)

Sometimes it is useful to access a multidimensional array as a C-style multidimensional array so that algorithms can be implemented using C's a[i][j][k] syntax. This routine returns a pointer, ptr, that simulates this kind of C-style array, for 1-, 2-, and 3-d ndarrays.

- op The address to any Python object. This Python object will be replaced with an equivalent well-behaved, C-style contiguous, ndarray of the given data type specifice by the last two arguments. Be sure that stealing a reference in this way to the input object is justified.
- ptr The address to a (ctype\* for 1-d, ctype\*\* for 2-d or ctype\*\*\* for 3-d) variable where ctype is the equivalent C-type for the data type. On return, ptr will be addressable as a 1-d, 2-d, or 3-d array.
- **dims** An output array that contains the shape of the array object. This array gives boundaries on any looping that will take place.
- **nd** The dimensionality of the array (1, 2, or 3).

**type\_num** The expected data type of the array.

**itemsize** This argument is only needed when type\_num represents a Flexible array. Otherwise it should be 0.

The simulation of a C-style array is not complete for 2-d and 3-d arrays. For example, the simulated arrays of pointers cannot be passed to subroutines expecting specific, statically-defined 2-d and 3-d arrays.

## PyArray\_Free (int) (PyObject\* op, void\* ptr)

Must be called with the same objects and memory locations returned from **PyArray\_AsCArray**(
This function cleans up memory that otherwise would get leaked.

## PyArray\_Concatenate (PyObject\*) (PyObject\* obj, int axis)

Join the sequence of objects in obj together along axis into a single array. If the dimensions or types are not compatible an error is raised.

## PyArray\_InnerProduct (PyObject\*) (PyObject\* obj1, PyObject\* obj2)

Compute a product-sum over the last dimensions of obj1 and obj2. Neither array is conjugated.

# $\mathbf{PyArray\_MatrixProduct} \ \ (\mathtt{Py0bject*}) \ (\mathtt{Py0bject*} \ \mathrm{obj1}, \ \mathtt{Py0bject*} \ \mathrm{obj})$

Compute a product-sum over the last dimension of obj1 and the second-to-last dimension of obj2. For 2-d arrays this is a matrix-product. Neither array is conjugated.

## PyArray\_CopyAndTranspose (PyObject\*) (PyObject\* op)

A specialized copy and transpose function that works only for 2-d arrays. The returned array is a transposed copy of op.

#### PyArray\_Correlate (PyObject\*) (PyObject\* op1, PyObject\* op2, int mode)

Compute the 1-d correlation of the 1-d arrays op1 and op2. The correlation is computed at each output point by multiplying op1 by a shifted version of op2 and summing the result. As a result of the shift, needed values outside of the defined range of op1 and op2 are interpreted as zero. The mode determines how many shifts to return: 0 - return only shifts that did not need to assume zero-values; 1 - return an object that is the same size as op1, 2 - return all possible shifts (any overlap at all is accepted).

PyArray\_Where (PyObject\*) (PyObject\* condition, PyObject\* x, PyObject\*
y)

If both x and y are NULL, then return **PyArray\_Nonzero**(condition). Otherwise, both x and y must be given and the object returned is shaped like condition and has elements of x and y where condition is respectively true or false.

#### 13.3.6.2 Other functions

PyArray\_CheckStrides (Bool) (int elsize, int nd, intp numbytes, intp\* dims, intp\* newstrides)

Determine if newstrides is a strides array consistent with the memory of an nddimensional array with shape dims and element-size, elsize. The newstrides array is checked to see if jumping by the provided number of bytes in each direction will ever mean jumping more than numbytes which is the assumed size of the available memory segment. If numbytes is 0, then an equivalent numbytes is computed assuming nd, dims, and elsize refer to a singlesegment array. Return TRUE if newstrides is acceptable, otherwise return FALSE.

PyArray\_MultiplyList (intp) (intp\* seq, int n)

PyArray\_MultiplyIntList (int) (int\* seq, int n)

Both of these routines multiply an n-length array, seq, of integers and return the result. No overflow checking is performed.

PyArray\_CompareLists (int) (intp\* l1, intp\* l2, int n)

Given two n-length arrays of integers, 11, and 12, return 1 if the lists are identical. Otherwise, return 0.

# 13.3.7 Array Iterators

An array iterator is a simple way to access the elements of an N-dimensional array quickly and efficiently. Section 15.1.1 provides more description and examples of this useful approach to looping over an array.

PyArray\_IterNew (PyObject\*) (PyObject\* arr)

Return an array iterator object from the array, arr. This is equivalent to arr.flat.

The array iterator object makes it easy to loop over an N-dimensional discontiguous array in C-style contiguous fashion.

### PyArray\_IterAllButAxis (PyObject\*) (PyObject\* arr, int axis)

Return an array iterator that will iterate over all axes but the one provided. The returned iterator cannot be used with PyArray\_ITER\_GOTO1D. This iterator could be used to write something similar to what ufuncs do wherein the loop over the largest axis is done by a separate sub-routine.

### PyArrayIter\_Check (int) (PyObject\* op)

Evaluates true if op is an array iterator (or instance of a subclass of the array iterator type).

## PyArray\_ITER\_RESET (void) (PyArrayIterObject\* iterator)

Reset an iterator to the beginning of the array.

## PvArray\_ITER\_NEXT (void) (PvArrayIterObject\* iterator)

Incremement the index and the dataptr members of the iterator to point to the next element of the array. If the array is not (C-style) contiguous, also increment the N-dimensional coordinates array.

# PyArray\_ITER\_GOTO (void) (PyArrayIterObject\* iterator, intp\* destination)

Set the iterator index, dataptr, and coordinates members to the location in the array indicated by the N-dimensional array, destination, which must have size at least iterator->nd\_m1+1.

## PyArray\_ITER\_GOTO1D (PyArrayIterObject\* iterator, index)

Set the iterator index and dataptr to the location in the array indicated by the 1-dimensional flat integer index. The index can be any integer but will be cast to intp.

# 13.3.8 Broadcasting (multi-iterators)

## PyArray\_MultiIterNew (PyObject\*) (int num, ...)

A simplified interface to broadcasting. This function takes the number of arrays to broadcast and then num extra (PyObject\*) arguments. These arguments are converted to arrays and iterators are created. PyArray\_Broadcast is then called on the resulting multi-iterator object. The resulting, broadcasted multi-iterator object is then returned. A broadcasted operation can then be performed using a single loop and using PyArray\_MultiIter\_NEXT(..)

### PyArray\_MultiIter\_RESET (void) (PyObject\* multi)

Reset all the iterators to the beginning in a multi-iterator object, multi.

## PyArray\_MultiIter\_NEXT (void) (PyObject\* multi)

Advance each iterator in a multi-iterator object, multi, to its next (broadcasted) element.

### PyArray\_MultiIter\_GOTO (void) (PyObject\* multi, intp\* destination)

Advance each iterator in a multi-iterator object,  $\mathtt{multi}$ , to the given N-dimensional destination where N is the number of dimensions in the broadcasted array.

## PyArray\_MultiIter\_GOTO1D (void) (PyObject\* multi, intp index)

Advance each iterator in a multi-iterator object, multi, to the corresponding location of the 1-d index into the broadcasted array.

## PyArray\_Broadcast (int) (PyArrayMultiIterObject\* mit)

This function encapsulates the broadcasting rules. The mit container should already contain iterators for all the arrays that need to be broadcast. On return, these iterators will be adjusted so that iteration over each simultaneously will accomplish the broadcasting. A negative number is returned if an error occurs.

### PvArray\_RemoveLargest (int) (PvArrayMultiIterObject\* mit)

This function takes a multi-iterator object that has been previously "broad-casted," finds the longest dimension in the broadcasted result and adapts all the iterators so as not to iterate over that dimension (by effectively making them of length-1 in that dimension). The longest dimension is returned unless mit->nd is 0, then -1 is returned. This function is useful for constructing ufunc-like routines that broadcast their inputs correctly and then call a strided 1-d version of the routine as the inner-loop. This 1-d version is usually optimized for speed.

# 13.3.9 Array Scalars

## PyArray\_Return (PyObject\*) (PyArrayObject\* arr)

This function simply checks to see if arr is a 0-dimensional array and, if so, returns the appropriate array scalar. It should be used whenever 0-dimensional arrays could be returned to Python.

PyArray\_Scalar (PyObject\*) (void\* data, PyArray\_Descr\* dtype, PyObject\*
 itemsize)

Return an array scalar object of the given enumerated typenum and itemsize by *copying* from memory pointed to by data. If swap is nonzero then this function will byteswap the data if appropriate to the data type because array scalars are always in correct machine-byte order.

## PyArray\_ToScalar (PyObject\*) (void\* data, PyArrayObject\* arr)

Return an array scalar object of the type and itemsize indicated by the array object arr copied from the memory pointed to by data and swapping if the NOTSWAPPED flag in arr is 0.

PyArray\_FromScalar (PyObject\*) (PyObject\*scalar, PyArray\_Descr\*outcode)

Return a 0-dimensional array of type determined by outcode from scalar which should be an array-scalar object. If outcode is NULL, then the type is determined from scalar.

## PyArray\_ScalarAsCtype (void) (PyObject\* scalar, void\* ctypeptr)

Return in ctypeptr a pointer to the actual value in an array scalar. There is no error checking, scalar must be an array-scalar object, and ctypeptr must have enough space to hold the correct type. For flexible types, a pointer to the data is copied into the memory of ctypeptr, for all other types, the actual data is copied into the address pointed to by ctypeptr.

PyArray\_CastScalarToCtype (void) (PyObject\*scalar, void\*ctypeptr, PyArray\_Descr\*outcode)

Return the data (cast to the data type indicated by outcode) from the arrayscalar, scalar, into the memory pointed to by ctypeptr (which must be large enough to handle the incoming memory).

## PyArray\_TypeObjectFromType (PyObject\*) (int type)

Returns a scalar type-object from a type-number. Equivalent to PyArray\_DescrFromType(type)->typeobj except for reference counting and error-checking. Returns a new reference to the typeobject on success or NULL on failure.

**PyArray\_ScalarKind** (PyArray\_SCALARKIND) (int typenum, PyArrayObject \*\*arr)

Return the kind of scalar represented by the typenumber and the array in \*arr (if arr is not NULL). The array is assumed to be rank-0 and only used if the typenum represents a signed integer. If arr is not NULL and the first element is negative then PyArray\_INTNEG\_SCALAR is returned, otherwise PyArray\_INTPOS\_SCALAR is returned. The possible return values are PyArray\_<a href="kind">kind</a> SCALAR where <a href="kind">kind</a> can be INTPOS, INTNEG, FLOAT, COMPLEX, BOOL, or OBJECT. PyArray\_NOSCALAR is also an enumerated value PyArray\_SCALARKIND variables can take on.

PyArray\_CanCoerceScalar (int) (char thistype, char neededtype, PyArray\_SCALARKIND scalar)

Implements the rules for scalar coercion. Scalars are only silently coerced from thistype to neededtype if this function returns nonzero. If scalar is PyArray\_NOSCALAR, then this function is equivalent to PyArray\_CanCastSafely. The rule is that scalars of the same KIND can be coerced into arrays of the same KIND. This rule means that high-precision scalars will never cause low-precision arrays of the same KIND to be upcast.

# 13.3.10 Data-type descriptors

When a data-type descriptor object is returned it is a new reference. Functions that take PyArray\_Descr\* objects and return arrays steal references to their inputs unless otherwise noted.

PyArrayDescr\_Check (int) (PyObject\* obj)

Evaluates as true if obj is a dtypedescr object (PyArray\_Descr\* object).

PyArray\_DescrNew (PyArray\_Descr\*) (PyArray\_Descr\* obj)

Return a new dtypedescr object copied from obj (the fields reference is just updated so that the new object points to the same fields dictionary).

# PyArray\_DescrNewFromType (PyArray\_Descr\*) (int typenum)

Create a new dtypedescr object from the built-in (or user-registered) data-type indicated by typenum. All builtin types should not have any of their fields changed. This creates a new copy of the PyArray\_Descr structure so that you can fill it in as appropriate. This function is especially needed for flexible data-types which need to have a new elsize member in order to be meaningful in array construction.

# PyArray\_DescrNewByteorder (PyArray\_Descr\*) (PyArray\_Descr\* obj, char newendian)

Create a new dtypedescr object with the byteorder set according to newendian. All referenced data-type descriptors (in subdescr and fields) are also changed (recursively). If a byteorder of PyArray\_IGNORE is encountered it is left alone. If newendian is PyArray\_SWAP, then all byte-orders are swapped. Other valid newendian values are PyArray\_NATIVE, PyArray\_LITTLE, and PyArray\_BIG which all cause the returned data-typed descriptor (and all it's referenced data-type descriptors) to have the corresponding byte-order.

# PyArray\_DescrFromObject (PyArray\_Descr\*) (PyObject\*op, PyArray\_Descr\* mintype)

Determine an appropriate data-type descriptor from the object op (which should be a "nested" sequence object) and the minimum data-type descriptor mintype (which can be NULL). Similar in behavior to array(op).dtypedescr. Don't confuse this function with with PyArray\_DescrConverter. This function looks essentially looks at all the objects in the (nested) sequence and determines the data-type from what it finds.

# PyArray\_DescrFromScalar (PyArray\_Descr\*) (PyObject\* scalar)

Return a data-type descriptor from an array-scalar object. No checking is done to be sure that scalar is an array scalar. If it is not, then the typecode will correspond to a PyArray\_OBJECT.

# PyArray\_DescrFromType (PyArray\_Descr\*) (int typenum)

Returns a dtypedescr object corresponding to typenum. The typenum can be one of the enumerated types, a character code for one of the enumerated types, or a user-defined type.

# PyArray\_DescrConverter (int) (PyObject\* obj, PyArray\_Descr\*\* dtype)

Convert any compatible Python object, obj, to a dtypedescr object in dtype. A large number of Python objects can be converted to dtypedescr objects. See Chapter 7 for a complete description. This version of the converter converts None objects to a PyArray\_LONG data-type descriptor. This function can be used in "O&" PyArg\_ParseTuple processing.

## PyArray\_DescrConverter2 (int) (int) (PyObject\* obj, PyArray\_Descr\*\* dtype)

Convert any compatible Python object, obj, to a dtypedescr object in dtype. This version of the converter converts None objects so that the returned datatype is NULL. This function can also be used in "O&" PyArg\_ParseTuple processing.

## 13.3.11 Conversion Utilities

### 13.3.11.1 For use with PyArg\_ParseTuple

All of these functions can be used in **PyArg\_ParseTuple**(...) with the "O&" format specifier to automatically convert any Python Object to the required C-Object. All of these functions return PY\_SUCCESS if successful and PY\_FAIL if not. The first argument to all of these function is a Python object. The second argument is the *address* of the c-type to convert the Python object to.



## WARNING

Be sure to understand what steps you should take to manage the memory when using these conversion functions. These functions can require freeing memory, incrementing or decrementing reference counts of specific objects based on your use.

## PyArray\_Converter (int) (PyObject\* obj, PyObject\*\* address)

Convert any Python object to a PyArrayObject. If PyArray\_Check(obj) is TRUE then its reference count is incremented and a reference placed in address. If obj is not an array, then convert it to an array using PyArray\_FromAny. No matter what is returned, you must DECREF the object returned by this routine in address when you are done with it.

## PyArray\_IntpConverter (int) (PyObject\* obj, PyArray\_Dims\* seq)

Convert any Python sequence, obj, smaller than MAX\_DIMS to an array of intp's. The Python sequence could also be a single number. The seq variable is a pointer to a structure with members ptr and len. On successful return, seq->ptr contains a pointer to memory that must be freed to avoid a memory leak. The restriction on memory size allows this converter to be conveniently used for sequences intended to be interpreted as array shapes.

# PyArray\_BufferConverter (int) (PyObject\* obj, PyArray\_Chunk\* buf)

Convert any Python object, obj, with a (single-segment) buffer interface to a variable with members that detail the object's use of its chunk of memory. The

buf variable is a pointer to a structure with base, ptr, len, and flags members. The PyArray\_Chunk structure is binary compatibile with the Python's buffer object through its len member on 32-bit platforms and its ptr member on 64-bit platforms. On return, the base member is set to obj (or its base if obj is already a buffer object pointing to another object). If you need to hold on to the memory be sure to INCREF the base member. The chunk of memory is pointed to by buf->ptr member and has length buf->len. The flags member of buf is BEHAVED\_RO with the WRITEABLE flag set if obj has a writeable buffer interface.

## PyArray\_AxisConverter (int) (PyObject\* obj, int\* axis)

Convert a Python object, obj, representing an axis argument to the proper value for passing to the functions that take an integer axis. Specifically, if obj is None, set axis to MAX\_DIMS which is interpreted by all these C-API functions correctly.

## PyArray\_BoolConverter (int) (PyObject\* obj, Bool\* value)

Convert any Python object, obj, to TRUE or FALSE, and place the result in value.

# PyArray\_ByteorderConverter (int) (PyObject\* obj, char\* endian)

Convert Python strings into the corresponding byte-order character: '>', '<', 's', '=', or '|'.

# PyArray\_SortkindConverter (int) (PyObject\* obj, PyArray\_SORTKIND\* sort)

Convert Python strings into one of PyArray\_QUICKSORT (startswith 'q' or 'Q'), PyArray\_HEAPSORT (startswith 'h' or 'H'), or PyArray\_MERGESORT (startswith 'm' or 'M').

#### 13.3.11.2 Other conversions

#### PyArray\_PyIntAsInt (int) (PyObject\* op)

Convert all kinds of Python objects (including arrays and array scalars) to a standard integer. On error, -1 is returned and an exception set. You may find useful the macro:

#define error\_converting(x) (((x) == -1) && PyErr\_Occurred()

### PyArray\_PyIntAsIntp (intp) (PyObject\* op)

Convert all kinds of Python objects (including arrays and array scalars) to a (platform-pointer-sized) integer. On error, -1 is returned and an exception set.

## PyArray\_IntpFromSequence (int) (PyObject\* seq, intp\* vals, int maxvals)

Convert any Python sequence (or single Python number) passed in as seq to (up to) maxvals pointer-sized integers and place them in the vals array. The sequence can be smaller then maxvals as the number of converted objects is returned.

### PyArray\_TypestrConvert (int) (int itemsize, int gentype)

Convert typestring characters (with itemsize) to basic enumerated data types. The typestring character corresponding to signed and unsigned integers, floating point numbers, and complex-floating point numbers are recognized and converted. Other values of gentype are returned. This function can be used to convert, for example, the string 'f4' to PyArray-FLOAT32.

#### 13.3.12 Miscellaneous

## 13.3.12.1 Importing the API

In order to make use of the C-API from another extension module, the import\_array() command must be used. If the extension module is self-contained in a single .c file, then that is all that needs to be done. If however, the extension module involve multiple files where the C-API is needed then some additional steps must be taken.

#### import\_array (int) (void)

This function must be called in the initialization section of a module that will make use of the C-API. It imports the module where the function-pointer table is stored and points the correct variable to it.

### PY\_ARRAY\_UNIQUE\_SYMBOL

## NO\_IMPORT\_ARRAY

Using these #defines you can use the C-API in multiple files for a single extension module. In each file you must define PY\_ARRAY\_UNIQUE\_SYMBOL to some name that will hold the C-API (e.g. myextension\_ARRAY\_API). This must be done **before** including the numpy/arrayobject.h file. In the module

intialization routine you call import\_array(). In addition, in the files that do not have the module initialization sub\_routine define NO\_IMPORT\_ARRAY prior to including numpy/arrayobject.h.

Suppose I have two files coolmodule.c and coolhelper.c which need to be compiled and linked into a single extension module. Suppose coolmodule.c contains the required initcool module initialization function (with the import\_array() function called). Then, coolmodule.c would have at the top:

```
#define PY_ARRAY_UNIQUE_SYMBOL cool_ARRAY_API
#include numpy/arrayobject.h
```

On the other hand, coolhelper.c would contain at the top:

```
#define PY_ARRAY_UNIQUE_SYMBOL cool_ARRAY_API
#define NO_IMPORT_ARRAY
#include numpy/arrayobject.h
```

#### PY ARRAY TYPES PREFIX

This constant can be defined to a user-defined name to attach as a prefix to the data-type names defined by numpy/arrayobject.h in case a clash is found with another library. The altered names are: longlong, ulonglong, Bool, longdouble, byte, ubyte, ushort, uint, ulong, cfloat, cdouble, clongdouble, Int8, UInt8, Int16, UInt16, Int32, UInt32, Int64, UInt64, Int128, UInt128, Int256, UInt256, Float16, Complex32, Float32, Complex64, Float64, Complex128, Float80, Complex160, Float96, Complex192, Float128, Complex256, intp, uintp.

#### PyArray\_GetNDArrayCVersion (unsigned int) (void)

This just returns the value NDARRAY\_VERSION. Because it is in the C-API, however, comparing the output of this function from the value defined in the current header gives a way to test if the C-API has changed thus requiring a re-compilation of extension modules that use the C-API.

#### 13.3.12.2 Internal Flexibility

#### PyArray\_SetNumericOps (int) (PyObject\* dict)

NumPy stores an internal table of Python callable objects that are used to implement arithmetic operations for arrays as well as certain array calculation methods. This function allows the user to replace any or all of these Python

objects with their own versions. The keys of the dictionary, dict, are the named functions to replace and the paired value is the Python callable object to use. Care should be taken that the function used to replace an internal array operation does not itself call back to that internal array operation (unless you have designed the function to handle that), or an unchecked infinite recursion can result (possibly causing program crash). The key names that represent operations that can be replaced are:

add, subtract, multiply, divide, remainder, power, sqrt, negative, absolute, invert, left\_shift, right\_shift, bitwise\_and, bitwise\_xor, bitwise\_or, less, less\_equal, equal, not\_equal, greater, greater\_equal, floor\_divide, true\_divide, logical\_or, logical\_and, floor, ceil, maximum, minimum.

These functions are included here because they are used at least once in the array object's methods. The function returns -1 (without setting a Python Error) if one of the objects being assigned is not callable.

# PyArray\_GetNumericOps (PyObject\*) (void)

Return a Python dictionary containing the callable Python objects stored in the the internal arithmetic operation table. The keys of this dictionary are given in the explanation for PyArray\_SetNumericOps.

## PyArray\_SetStringFunction (void) (PyObject\* op, int repr)

This function allows you to alter the tp\_str and tp\_repr methods of the array object to any Python function. Thus you can alter what happens for all arrays when str(arr) or repr(arr) is called from Python. The function to be called is passed in as op. If repr is non-zero, then this function will be called in response to repr(arr), otherwise the function will be called in response to str(arr). No check on whether or not op is callable is performed. The callable passed in to op should expect an array argument and should return a string to be printed.

#### 13.3.12.3 Memory management

PyDataMem\_NEW (char\*) (size\_t nbytes)

PyDataMem\_FREE (char\* ptr)

PyDataMem\_RENEW (char\*) (void \* ptr, size\_t newbytes)

Macros to allocate, free, and reallocate memory. These macros are used internally to create arrays.

```
PyDimMem_NEW (intp*) (nd)

PyDimMem_FREE (intp* ptr)

PyDimMem_RENEW (intp*) (intp* ptr, intp newnd)

Macros to allocate, free, and reallocate dimension and strides memory.
_pya_malloc (nbytes)
_pya_free (ptr)
_pya_realloc (ptr, nbytes)
```

These macros use different memory allocators, depending on the constant PyArray\_USE\_PYMEM Currently, the system malloc or the Python Object allocator can be used.

#### 13.3.12.4 Threading support

These macros are only useful if ALLOW\_THREADS is defined during compilation of the extension module. Otherwise, these macros are equivalent to whitespace. Python uses a single Global Interpreter Lock (GIL) for each Python process so that only a single thread may excecute at a time (even on multi-cpu machines). When calling out to a compiled function that may take time to compute, the GIL should be released so that other Python threads can run while the time-consuming calculations are performed. This can be accomplished using two groups of macros. Typically, if one macro in a group is used in a code block, all of them must be used in the same code block.

- **Group 1** This group is used to call code that may take some time but does not use any Python C-API calls. Thus, the GIL should be released during its calculation.
  - **BEGIN\_THREADS\_DEF** Place in the variable declaration area. This macro sets up the variable needed for storing the Python state.
  - **BEGIN\_THREADS** Place right before code that does not need the Python interpreter (no Python C-API calls). This macro saves the Python state and releases the GIL.
  - **END\_THREADS** Place right after code that does not need the Python interpreter. This macro acquires the GIL and restores the Python state from the saved variable.

Group 2 This group is used to re-acquire the Python GIL after it has been released. For example, suppose the GIL has been released (using the previous calls), and then some path in the code (perhaps in a different subroutine) requires use of the Python C-API, then these macros are useful to acquire the GIL. These macros accomplish essentially a reverse of the previous three (acquire the LOCK saving what state it had) and then re-release it with the saved state.

**ALLOW\_C\_API\_DEF** Place in the variable declaration area to set up the necessary variable.

**ALLOW\_C\_API** Place before code that needs to call the Python C-API (when it is known that the GIL has already been released).

**DISABLE\_C\_API** Place after code that needs to call the Python C-API (to re-release the GIL).



#### WARNING

Never use semicolons after the threading support macros.

#### 13.3.12.5 Priority

PyArray\_PRIOIRTY Default priority for arrays.

PyArray\_BIG\_PRIORITY Default priority for the big array.

PyArray\_SUBTYPE\_PRIORITY Default subtype priority.

PyArray\_GetPriority (double) (PyObject\* obj, double def)

Return the <u>\_array\_priority\_</u> attribute (converted to a double) of obj or def if no attribute of that name exists. Fast returns that avoid the attribute lookup are provided for objects of type **PyArray\_Type**.

#### 13.3.12.6 Default buffers

PyArray\_BUFSIZE Default size of the user-settable internal buffers.

PyArray\_MIN\_BUFSIZE Smallest size of user-settable internal buffers.

PyArray\_MAX\_BUFSIZE Largest size allowed for the user-settable buffers.

#### 13.3.12.7 Other constants

PyArray\_NUM\_FLOATTYPE The number of floating-point types

MAX\_DIMS The maximum number of dimensions allowed in arrays.

**NDARRAY\_VERSION** The current version of the ndarray object (check to see if this variable is defined to guarantee the numpy/arrayobject.h header is being used).

**FALSE** Defined as 0 for use with Bool.

**TRUE** Defined as 1 for use with Bool.

**PY\_FAIL** The return value of failed converter functions which are called using the "O&" syntax in PyArg\_ParseTuple-like functions.

**PY\_SUCCEED** The return value of successful converter functions which are called using the "O&" syntax in PyArg\_ParseTuple-like functions.

#### 13.3.12.8 Miscellaneous Macros

# $\mathbf{MAX}$ (a,b)

Returns the maximum of R and b. If (a) and (b) are expressions they are evaluated twice.

## MIN (a,b)

Returs the minimum of a and b. If (a) and (b) are expressions they are evaluated twice.

#### tMAX (a, b, typ)

Returns the maximum of a and b and these are only evaluated once. However, the c-type of both a and b must be provided in typ.

## tMIN (a, b, typ)

Returns the minimum of a and b (only evaluating them once). The c-type of both a and b must be provided by typ.

## **REFCOUNT** (PyObject\* op)

Returns the reference count of any Python object.

## PyArray\_XDECREF\_ERR (PyObject \*obj)

DECREF's an array object which may have the UPDATEIFCOPY flag set without causing the contents to be copied back into the original array. Resets the WRITEABLE flag on the base object. This is useful for recovering from an error condition when UPDATEIFCOPY is used.

### 13.3.12.9 Enumerated Types

**PyArray\_SORTKIND** A special variable-type which can take on the values **PyArray\_**<KIND> where <KIND> is

## QUICKSORT, HEAPSORT, MERGESORT

PyArray\_NSORTS is defined to be the number of sorts.

**PyArray\_SCALARKIND** A special variable type indicating the number of "kinds" of scalars distinguished in determining scalar-coercion rules. This variable can take on the values PyArray\_<KIND> where <KIND> can be

NOSCALAR, BOOL\_SCALAR, INTPOS\_SCALAR, INTNEG\_SCALAR, FLOAT\_SCALAR, COMPLEX\_SCALAR, OBJECT\_SCALAR

**PyArray\_NSCALARKINDS** is defined to be the number of scalar kinds (not including **PyArray\_NOSCALAR**).

**PyArray\_ORDER** A variable type indicating the order that an array should be interpreted in. The value of a variable of this type can be PyArray\_<ORDER> where <ORDER> is

ANYORDER, CORDER, FORTRANORDER

## 13.4 UFunc API

#### 13.4.1 Constants

UFUNC\_ERR\_<HANDLER>

<HANDLER> can be IGNORE, WARN, RAISE, or CALL

UFUNC\_<THING>\_<ERR>

<THING> can be MASK, SHIFT, or FPE, and <ERR> can be DIVIDE-BYZERO, OVERFLOW, UNDERFLOW, and INVALID.

PyUFunc\_<VALUE> <VALUE> can be One (1), Zero (0), or None (-1)

## 13.4.2 Macros

#### LOOP\_BEGIN\_THREADS

Used in universal function code to only release the Python GIL if loop->obj is not true (this is not an OBJECT array loop).

#### LOOP\_END\_THREADS

Used in universal function code to re-acquire the Python GIL if it was released (because loop->obj was not true).

## UFUNC\_CHECK\_ERROR (loop)

A macro used internally to check for errors and goto fail if found. This macro requires a fail label in the current code block. The loop variable must have at least members (obj, errormask, and errorobj). If loop->obj is nonzero, then PyErr\_Occurred() is called (meaning the GIL must be held). If loop->obj is zero, then if loop->errormask is nonzero, PyUFunc\_checkfperr is called with arguments loop->errormask and loop->errobj. If the result of this check of the IEEE floating point registers is true then the code redirects to the fail label.

## UFUNC\_CHECK\_STATUS (ret)

A macro that expands to platform-dependent code. The ret variable can can be any integer. The UFUNC\_FPE\_<ERR> bits are set in ret according to the status of the corresponding IEEE error flags of the floating point processor.

## 13.4.3 Functions

Create a new broadcasting universal function from required variables. Each ufunc builds around the notion of an element-by-element operation. The number of inputs to this operation is  ${\tt nin}$ . The number of outputs is  ${\tt nout}$ . The  ${\tt ntypes}$  variable is how many different data-type "signatures" the ufunc has implemented. The  ${\tt func}$  argument must point to an array of length  ${\tt ntypes}$  containing PyUFuncGenericFunction items. These items are pointers to functions that acutally implement the underlying (element-by-element) function N times. The types array must be of length  $({\tt nin+nout})*{\tt ntypes}$ , and it contains the (built-in) data types that the corresponding function in the  ${\tt func}$ 

array can deal with. The data argument should be NULL or a pointer to an array of size ntypes. The data array may contain arbitrary extra-data to be passed to the corresponding 1-d loop function in the func array. The name of the ufunc is passed in as name. A documenation string can be passed in as doc. The documentation string should not contain the name of the function or the calling signature as that will be dynamically determined from the object and available when accessing the \_doc\_ attribute of the ufunc. The check\_return variable is unused and present for backwards compatibility. A corresponding check\_return integer does get set in the ufunc object created.

### 

This function allows the user to register a 1-d loop with an already-created ufunc to be used whenever the ufunc is called with any of its input arguments as the user-defined data-type. This is needed in order to make ufuncs work with built-in data-types. The data-type must have been previously registered with the numpy system. The loop is passed in as function which can take arbitrary data passed in as data. The data-types the loop requires are passed in as arg\_types which must be a pointer to persistent memory at least as large as ufunc->nin + ufunc->nout.

## PyUFunc\_GenericFunction (int) (PyUFuncObject\*self, PyObject\*args, PyArrayObject\*\* mps)

A generic ufunc call. The ufunc is passed in as self, the arguments to the ufunc as args. The mps argument is an array of PyArrayObject pointers containing the converted input arguments as well as the ufunc outputs on return. The user is responsible for managing this array and receives a new reference for each array in mps. The total number of arrays in mps is given by self->nin + self->nout.

#### PyUFunc\_checkfperr (int) (int errmask, PyObject\* errobj)

A simple interface to the IEEE error-flag checking support. The errmask argument is a mask of UFUNC\_MASK\_<ERR> bitmasks indicating which errors to check for (and how to check for them). The errobj must be a Python tuple with two elements: a string containing the name which will be used in any communication of error and either a callable Python object (call-back function) or None. The callable object will only be used if UFUNC\_ERR\_CALL is set as the desired error checking method. This routine manages the GIL and

is safe to call even after releasing the GIL. If an error in the IEEE-compatibile hardware is determined a -1 is returned, otherwise a 0 is returned.

#### PyUFunc\_clearfperr (void) ()

Clear the IEEE error flags.

PyUFunc\_GetPyValues (void) (char\* name, int\* bufsize, int\* errmask, PyObject\*\*
errobj)

Get the python values used for ufunc processing from the local, global, and builtin namespace (in that order) unless the defaults have been set in which case the name lookup is bypassed. The name is placed as a string in the first element of \*errobj. The second element is the looked-up function to call on error callback. The value of the looked-up buffer-size to use is passed into bufsize, and the value of the error mask is placed into errmask.

#### 13.4.4 Generic functions

At the core of every ufunc is a collection of type-specific functions that defines the basic functionality for each of the supported types. These functions must evaluate the underlying function  $N \geq 1$  times. Extra-data may be passed in that may be used during the calculation. This feature allows some general functions to be used as these basic looping functions. The general function has all the code needed to point variables to the right place and set up a function call. The general function assumes that the actual function to call is passed in as the extra data and calls it with the correct values. All of these functions are suitable for placing directly in the array of functions stored in the functions member of the PyUFuncObject structure.

PyUFunc\_f\_f\_As\_d\_d (void) (char\*\* args, intp\* dimensions, intp\* steps, void\*
func)

PyUFunc\_d\_d (void) (char\*\* args, intp\* dimensions, intp\* steps, void\* func)

PyUFunc\_f\_f (void) (char\*\* args, intp\* dimensions, intp\* steps, void\* func)

PyUFunc\_g\_g (void) (char\*\* args, intp\* dimensions, intp\* steps, void\* func)

PyUFunc\_F\_F\_As\_D\_D (void) (char\*\* args, intp\* dimensions, intp\* steps,
 void\* func)

PyUFunc\_F\_F (void) (char\*\* args, intp\* dimensions, intp\* steps, void\* func)

PyUFunc\_D\_D (void) (char\*\* args, intp\* dimensions, intp\* steps, void\* func)

PyUFunc\_G\_G (void) (char\*\* args, intp\* dimensions, intp\* steps, void\* func)

Type specific, core 1-d functions for ufuncs where each calculation is obtained by calling a function taking one input argument and returning one output. This function is passed in func. The letters correspond to dtypechar's of the supported data types (f - float, d - double, g - long double, F - cfloat, D - cdouble, G - clongdouble). The argument func must support the same signature. The \_As\_X\_X variants assume ndarray's of one data type but cast the values to use an underlying function that takes a different data type. Thus, PyUFunc\_f\_f\_As\_d\_d uses ndarrays of data type PyArray\_FLOAT but calls out to a c-function that takes double and returns double.

PyUFunc\_ff\_As\_dd\_d (void) (char\*\* args, intp\* dimensions, intp\* steps,
 void\* func)

PyUFunc\_ff\_f (void) (char\*\* args, intp\* dimensions, intp\* steps, void\* func)

PyUFunc\_dd\_d (void) (char\*\* args, intp\* dimensions, intp\* steps, void\* func)

PyUFunc\_gg\_g (void) (char\*\* args, intp\* dimensions, intp\* steps, void\* func)

PyUFunc\_FF\_F\_As\_DD\_D (void) (char\*\* args, intp\* dimensions, intp\* steps,
 void\* func)

PyUFunc\_DD\_D (void) (char\*\* args, intp\* dimensions, intp\* steps, void\*
func)

PyUFunc\_FF\_F (void) (char\*\* args, intp\* dimensions, intp\* steps, void\* func)

PyUFunc\_GG\_G (void) (char\*\* args, intp\* dimensions, intp\* steps, void\*
func)

Type specific, core 1-d functions for ufuncs where each calculation is obtained by calling a function taking two input arguments and returning one output. The underlying function to call is passed in as func. The letters correspond to dtypechar's of the specific data type supported by the general-purpose function. The argument func must support the corresponding signature. The \_As\_XX\_X variants assume ndarrays of one data type but cast the values at each iteration of the loop to use the underlying function that takes a different data type.

PyUFunc\_O\_O (void) (char\*\* args, intp\* dimensions, intp\* steps, void\* func)

PyUFunc\_OO\_O (void) (char\*\* args, intp\* dimensions, intp\* steps, void\*
func)

One-input, one-output, and two-input, one-output core 1-d functions for the **PyArray\_OBJECT** data type. These functions handle reference count issues and return early on error. The actual function to call is **func** and it must accept calls with the signature (PyObject\*) (PyObject\*) for **PyUFunc\_O\_O** or (PyObject\*) (PyObject\*, PyObject\*) for **PyUFunc\_OO\_O**.

PyUFunc\_O\_O\_method (void) (char\*\* args, intp\* dimensions, intp\* steps,
 void\* func)

This general purpose 1-d core function assumes that func is a string representing a method of the input object. For each iteration of the loop, the Python obejct is extracted from the array and its func method is called returning the result to the output array.

PyUFunc\_OO\_O\_method (void) (char\*\* args, intp\* dimensions, intp\* steps,
 void\* func)

This general purpose 1-d core function assumes that func is a string representing a method of the input object that takes one argument. The first argument in args is the method whose function is called, the second argument in args is the argument passed to the function. The output of the function is stored in the third entry of args.

PyUFunc\_On\_Om (void) (char\*\* args, intp\* dimensions, intp\* steps, void\*
func)

This is the 1-d core function used by the dynamic ufuncs created by umath.frompyfunc(function, nin, nout). In this case func is a pointer to a PyUFunc\_PyFuncData structure which has definition {int nin; int nout; PyObject\* callable}. At each iteration of the loop, the nin input objects are exctracted from their object arrays and placed into an argument tuple, the Python callable is called with the input arguments, and the nout outputs are placed into their object arrays.

## 13.5 Importing the API

PY\_UFUNC\_UNIQUE\_SYMBOL

NO\_IMPORT\_UFUNC

import\_ufunc (int) (void)

These are the constants and functions for accessing the ufunc C-API from extension modules in precisely the same way as the array C-API can be accessed. The import\_ufunc() function must always be called (in the initialization subroutine of the extension module). If your extension module is in one file then that is all that is required. The other two constants are useful if your extension module makes use of multiple files. In that case, define PY\_UFUNC\_UNIQUE\_SYMBOL to something unique to your code and then in source files that do not contain the module initialization function but still need access to the UFUNC API, define PY\_UFUNC\_UNIQUE\_SYMBOL to the same name used previously and also define NO\_IMPORT\_UFUNC.

The C-API is actually an array of function pointers. This array is created (and pointed to by a global variable) by import\_ufunc. The global variable is either statically defined or allowed to be seen by other files depending on the state of Py\_UFUNC\_UNIQUE\_SYMBOL and NO\_IMPORT\_UFUNC.

## Chapter 14

## How to extend NumPy

## 14.1 Writing an extension module

While the ndarray object is designed to allow rapid computation in Python, it is also designed to be general-purpose and satisfy a wide-variety of computational needs. As a result, if absolute speed is essential, there is no replacement for a well-crafted, compiled loop specific to your application and hardware. This is one of the reasons that numpy includes f2py so that an easy-to-use mechanisms for linking (simple) C/C++ and (arbitrary) Fortran code directly into Python are available. You are encouraged to use and improve this mechanism. The purpose of this section is not to document this tool but to document the more basic steps to writing an extension module that this tool depends on.

When an extension module is written, compiled, and installed to somewhere in the Python path (sys.path), the code can then be imported into Python as if it were a standard python file. It will contain objects and methods that have been defined and compiled in C code. The basic steps for doing this in Python are well-documented and you can find more information in the documentation for Python itself available online at www.python.org http://www.python.org.

In addition to the Python C-API, there is a full and rich C-API for NumPy allowing sophisticated manipulations on a C-level. However, for most applications, only a few API calls will typically be used. If all you need to do is extract a pointer to memory along with some shape information to pass to another calculation routine, then you will use very different calls, then if you are trying to create a new array-like type or add a new data type for ndarrays. This chapter documents the API calls and macros that are most commonly used.

## 14.2 Required subroutine

There is exactly one function that must be defined in your C-code in order for Python to use it as an extension module. The function must be called init<name> where <name> is the name of the module from Python. This function must be declared so that it is visible to code outside of the routine. Besides adding the methods and constants you desire, this subroutine must also contain calls to import\_array() and/or import\_ufunc() depending on which C-API is needed. Forgetting to place these commands will show itself as an ugly segmentation fault (crash) as soon as any C-API subroutine is actually called. It is actually possible to have multiple init<name> functions in a single file in which case multiple modules will be defined by that file. However, there are some tricks to get that to work correctly and it is not covered here.

A minimal init<name> method looks like

```
PyMODINIT_FUNC
init<name>(void)
{
    (void)Py_InitModule(''<name>'', mymethods);
    (void) import_array();
}
```

The mymethods must be an array (usually statically declared) of PyMethodDef structures which contain method names, actual C-functions, a variable indicating whether the method uses keyword arguments or not, and docstrings. These are explained in the next section. If you want to add constants to the module, then you store the returned value from Py\_InitModule which is a module object. The most general way to add itmes to the module is to get the module dictionary using Py-Module\_GetDict(module). With the module dictionary, you can add whatever you like to the module manually. An easier way to add objects to the module is to use one of three additional Python C-API calls that do not require a separate extraction of the module dictionary. These are documented in the Python documentation, but repeated here for convenience:

```
PyModule_AddObject (int) (PyObject* module, char* name, PyObject* value)

PyModule_AddIntConstant (int) (PyObject* module, char* name, long value)

PyModule_AddStringConstant (int) (PyObject* module, char* name, char* value)
```

All three of these functions require the module object (the return value of Py\_InitModule). The name is a string that labels the value in the module. Depending on which function is called, the value argument is either a general object (Py-Module\_AddObject steals a reference to it), an integer constant, or a string constant.

## 14.3 Defining functions

The second argument passed in to the Py\_InitModule function is a structure that makes it easy to to define functions in the module. In the example given above, the mymethods structure would have been defined earlier in the file (usually right before the init<name> subroutine) to

Each entry in the mymethods array is a PyMethodDef structure containing 1) the Python name, 2) the C-function that implements the function, 3) flags indicating whether or not keywords are accepted for this function, and 4) The docstring for the function. Any number of functions may be defined for a single module by adding more entries to this table. The last entry must be all NULL as shown to act as a sentinel. Python looks for this entry to know that all of the functions for the module have been defined.

The last thing that must be done to finish the extension module is to actually write the code that performs the desired functions. There are two kinds of functions: those that don't accept keyword arguments, and those that do.

## 14.3.1 Functions without keyword arguments

Functions that don't accept keyword arguments should be written as

```
static PyObject*
nokeyword_cfunc (PyObject *dummy, PyObject *args)
```

```
{
    /* convert Python arguments */
    /* do function */
    /* return something */
}
```

The dummy argument is not used in this context and can be safely ignored. The args argument contains all of the arguments passed in to the function as a tuple. You can do anything you want at this point, but usually the easiest way to manage the input arguments is to call PyArg\_ParseTuple(args, format\_string, addresses\_to\_C\_variables...) or PyArg\_UnpackTuple(tuple, "name", min, max, ...). A good description of how to use the first function is contained in the Python C-API reference manual under section 5.5 (Parsing arguments and building values). You should pay particular attention to the "O&" format which uses converter functions to go between the Python object and the C object. All of the other format functions can be (mostly) thought of as special cases of this general rule. There are several converter functions defined in the NumPy C-API that may be of use. In particular, the PyArray\_DescrConverter function is very useful to support arbitrary data-type specification. This function transforms any valid data-type Python object into a PyArray\_Descr \* object. Remember to pass in the address of the C-variables that should be filled in.

There are lots of examples of how to use PyArg\_ParseTuple throughout the NumPy source code. The standard usage is like

It is important to keep in mind that you get a borrowed reference to the object when using the "O" format string. However, the converter functions usually require some form of memory handling. In this example, if the conversion is successful, dtype will hold a new reference to a PyArray\_Descr \* object, while input will hold a borrowed reference.

After the input arguments are processed, the code that actually does the work is written (likely calling other functions as needed). The final step of the C-function is to return something. If an error is encountered then NULL should be returned (making sure an error has actually been set). If nothing should be returned then increment Py\_None and return it. If a single object should be returned then it is

returned (ensuring that you own a reference to it first). If multiple objects should be returned then you need to return a tuple. The Py\_BuildValue(format\_string, c\_variables...) function makes it easy to build tuples of Python objects from C variables. Pay special attention to the difference between 'N' and 'O' in the format string or you can easily create memory leaks. The 'O' format string increments the reference count of the PyObject\* C\_variable it corresponds to, while the 'N' format string steals a reference to the corresponding PyObject\* C\_variable.

#### 14.3.2 Functions with keyword arguments

These functions are very similar to functions without keyword arguments. The only difference is that the function signature is

```
static PyObject*
keyword_cfunc (PyObject *dummy, PyObject *args, PyObject *kwds)
{
...
}
```

The kwds argument holds a Python dictionary whose keys are the names of the keyword arguments and whose values are the corresponding keyword-argument values. This dictionary can be processed however you see fit. The easiest way to handle it, however, is to replace the PyArg\_ParseTuple(args, format\_string, addresses...) function with a call to PyArg\_ParseTupleAndKeywords(args, kwds, format\_string, char \*kwlist[], addresses...). The kwlist parameter to this function is a NULL-terminated array of strings providing the expected keyword arguments. There should be one string for each entry in the format\_string. Using this function will raise a TypeError if invalid keyword arguments are passed in.

For more help on this function please see section 1.8 (Keyword Paramters for Extension Functions) of the Extending and Embedding tutorial in the Python documentation.

## 14.3.3 Reference counting

The biggest difficulty when writing extension modules is reference counting. It is an important reason for the popularity of f2py, weave, pyrex, ctypes, etc.... If you mishandle reference counts you can get problems from memory-leaks to segmentation faults. The only strategy I know of to handle reference counts correctly is blood, sweat, and tears. First, you force it into your head that every Python variable has a reference count. Then, you understand exactly what each function does to

the reference count of your objects, so that you can properly use DECREF and INCREF when you need them. Reference counting can really test the amount of patience and diligence you have towards your programming craft. Despite the grim depiction, most cases of reference counting are quite straightforward with the most common difficulty being not using DECREF on objects before exiting early from a routine due to some error. In second place, is the common error of not owning the reference on an object that is passed to a function or macro that is going to steal the reference (e.g. PyTuple\_SET\_ITEM, and most functions that take PyArray\_Descrobjects).

Typically you get a new reference to a variable when it is created or is the return value of some function (there are some prominent exceptions, however — such as getting an item out of a tuple or a dictionary). When you own the reference, you are responsible to make sure that Py\_DECREF(var) is called when the variable is no longer necessary (and no other function has "stolen" its reference). Also, if you are passing a Python object to a function that will "steal" the reference, then you need to make sure you own it (or use Py\_INCREF to get your own reference). You will also encounter the notion of borrowing a reference. A function that borrows a reference does not alter the reference count of the object and does not expect to "hold on" to the reference. It's just going to use the object temporarily. When you use PyArg\_ParseTuple or PyArg\_UnpackTuple you receive a borrowed reference to the objects in the tuple and should not alter their reference count inside your function. With practice, you can learn to get reference counting right, but it can be frustrating at first.

One common source of reference-count errors is the **Py\_BuildValue** function. Pay careful attention to the difference between the 'N' format character and the 'O' format character. If you create a new object in your subroutine (such as an output array), and you are passing it back in a tuple of return values, then you should most-likely use the 'N' format character in **Py\_BuildValue**. The 'O' character will increase the reference count by one. This will leave the caller with two reference counts for a brand-new array. When the variable is deleted and the reference count decremented by one, there will still be that extra reference count, and the array will not be deallocated. You will have a reference-counting induced memory leak. Using the 'N' character will avoid this situation as it will return to the caller an object (inside the tuple) with a single reference count.

## 14.4 Dealing with array objects

Most extension modules for NumPy will need to access the memory for an ndarray object (or one of it's sub-classes). The easiest way to do this doesn't require you to know much about the internals of NumPy. The method is to

- 1. Ensure you are dealing with a well-behaved array (aligned, in machine byteorder and single-segment) of the correct type and number of dimensions.
  - (a) By converting it from some Python object using PyArray\_FromAny or a macro built on it.
  - (b) By constructing a new ndarray of your desired shape and type using PyArray\_NewFromDescr or a simpler macro or function based on it.
- 2. Get the shape of the array and a pointer to its actual data.
- 3. Pass the data and shape information on to a subroutine or other section of code that actually performs the computation.
- 4. If you are writing the algorithm, then I recommend that you use the stride information contained in the array to access the elements of the array (the PyArray\_GETPTR macros make this painless). Then, you can relax your requirements so as not to force a single-segment array and the data-copying that might result.

Each of these sub-topics is covered in the following sub-sections.

## 14.4.1 Converting an arbitrary sequence object

The main routine for obtaining an array from any Python object that can be converted to an array is PyArray\_FromAny. This function is very flexible with many input arguments. Several macros make it easier to use the basic function. **PyArray\_FROM\_OTF** is arguably the most useful of these macros for the most common uses. It allows you to convert an arbitrary Python object to an array of a specific builtin data-type (e.g. float), while specifying a particular set of requirements (e.g. contiguous, aligned, and writeable). The syntax is

PyArray\_FROM\_OTF (PyObject\*) (PyObject\* obj, int typenum, int requirements)

Return an indarray from any Python object, obj, that can be converted to an array. The number of dimensions in the returned array is determined by the

object. The desired data-type of the returned array is provided in typenum which should be one of the enumerated types. The requirements for the returned array can be any combination of standard array flags. Each of these arguments is explained in more detail below. You receive a new reference to the array on success. On failure, NULL is returned and an exception is set.

obj The object can be any Python object convertable to an ndarray. If the object is already (a subclass of) the ndarray that satisfies the requirements then a new reference is returned. Otherwise, a new array is constructed. The contents of obj are copied to the new array unless the array interface is used so that data does not have to be copied. Objects that can be converted to an array include: 1) any nested sequence object, 2) any object exposing the array interface, 3) any object with an \_array\_method (which should return an ndarray), and 4) any scalar object (becomes an zero-dimensional array). Sub-classes of the ndarray that otherwise fit the requirements will be passed through. If you want to ensure a base-class array, then use ENSUREARRAY in the requirements flag. A copy is made only if necessary. If you want to guarantee a copy, then pass in ENSURECOPY to the requirements flag.

**typenum** One of the enumerated types or PyArray\_NOTYPE if the datatype should be determined from the object itself. The C-based names can be used:

PyArray\_BOOL, PyArray\_BYTE, PyArray\_UBYTE, PyArray\_SHORT, PyArray\_USHORT, PyArray\_INT, PyArray\_UINT, PyArray\_LONG, PyArray\_ULONG, PyArray\_LONGLONG, PyArray\_DOUBLE, PyArray\_LONGDOUBLE, PyArray\_CFLOAT, PyArray\_CDOUBLE, PyArray\_CLONGDOUBLE, PyArray\_OBJECT.

Alternatively, the bit-width names can be used as supported on the platform. For example:

PyArray\_INT8, PyArray\_INT16, PyArray\_INT32, PyArray\_INT64, PyArray\_UINT8, PyArray\_UINT16, PyArray\_UINT32, PyArray\_UINT64, PyArray\_FLOAT32, PyArray\_FLOAT64, PyArray\_COMPLEX64, PyArray\_COMPLEX128.

The object will be converted to the desired type only if it can be done without losing precision. Otherwise NULL will be returned and an error raised. Use FORCECAST in the requirements flag to override this behavior.

requirements The memory model for an ndarray admits arbitrary strides in each dimension to advance to the next element of the array. Often, however, you need to interface with code that expects a C-contiguous or a Fortran-contiguous memory layout. In addition, an ndarray can be misaligned (the address of an element is not at an integral multiple of the size of the element) which can cause your program to crash (or at least work more slowly) if you try and dereference a pointer into the array data. Both of these problems can be solved by converting the Python object into an array that is more "well-behaved" for your specific usage.

The requirements flag allows specification of what kind of array is acceptable. If the object passed in does not satisfy this requirements then a copy is made so that thre returned object will satisfy the requirements. these ndarray can use a very generic pointer to memory. This flag allows specification of the desired properties of the returned array object. All of the flags are explained in the detailed API chapter. The flags most commonly needed are IN\_ARRAY, OUT\_ARRAY, and INOUT\_ARRAY:

- IN\_ARRAY Equivalent to CONTIGUOUS | ALIGNED. This combination of flags is useful for arrays that must be in C-contiguous order and aligned. These kinds of arrays are usually input arrays for some algorithm.
- **OUT\_ARRAY** Equivalent to CONTIGUOUS | ALIGNED | WRITE-ABLE. This combination of flags is useful to specify an array that is in C-contiguous order, is aligned, and can be written to as well. Such an array is usually returned as output (although normally such output arrays are created from scratch).
- INOUT\_ARRAY Equivalent to CONTIGUOUS | ALIGNED | WRITE-ABLE | UPDATEIFCOPY. This combination of flags is useful to specify an array that will be used for both input and output. If a copy is needed, then when the temporary is deleted (by your use of Py\_DECREF at the end of the interface routine), the temporary array will be copied back into the original array passed in. Use of the UPDATEIFCOPY flag requires that the input object is already an array (because other objects cannot be automatically updated in this fashion). If an error occurs use PyArray\_DECREF\_ERR(obj) on an array with the UPDATEIFCOPY flag set. This will delete the array without causing the contents to be copied back into the original array.

Other useful flags that can be OR'd as additional requirements are:

**FORCECAST** Cast to the desired type, even if it can't be done without losing information.

ENSURECOPY Make sure the resulting array is a copy of the original. ENSUREARRAY Make sure the resulting object is an actual ndarray and not a sub-class.



#### NOTE

Whether or not an array is byte-swapped is determined by the data-type of the array. Native byte-order arrays are always requested by PyArray\_FROM\_OTF and so there is no need for a NOTSWAPPED flag in the requirements argument. There is also no way to get a byte-swapped array from this routine.

#### 14.4.2 Creating a brand-new ndarray

Quite often new arrays must be created from within extension-module code. Perhaps an output array is needed and you don't want the caller to have to supply it. Perhaps only a temporary array is needed to hold an intermediate calculation. Whatever the need there are simple ways to get an ndarray object of whatever datatype is needed. The most general function for doing this is PyArray\_NewFromDescr. All array creation functions go through this heavily re-used code. Because of its flexibility, it can be somewhat confusing to use. As a result, simpler forms exist that are easier to use.

#### PyArray\_SimpleNew (PyObject\*)(int nd, intp\* dims, int typenum)

This function allocates new memory and places it in an ndarray with nd dimensions whose shape is determined by the array of at least nd items pointed to by dims. The memory for the array is uninitialized (unless typenum is PyArray\_OBJECT in which case each element in the array is set to NULL). The typenum argument allows specification of any of the builtin data-types such as PyArray\_FLOAT or PyArray\_LONG. The memory for the array can be set to zero if desired using PyArray\_FILLWBYTE(return\_object, 0).

PyArray\_SimpleNewFromData (PyObject\*) (int nd, intp\* dims, int typenum, void\* data)

Sometimes, you want to wrap memory allocated elsewhere into an ndarray object for downstream use. This routine makes it straightforward to do that. The first three arguments are the same as in **PyArray\_SimpleNew**, the final argument is a pointer to a block of contiguous memory that the ndarray should use as it's data-buffer which will be interpreted in C-style contiguous fashion. A new reference to an ndarray is returned, but the ndarray will not own its data. When this ndarray is deallocated, the pointer will not be freed.

You should ensure that the provided memory is not freed while the returned array is in existence. The easiest way to handle this is if data comes from another reference-counted Python object. The reference count on this object should be increased after the pointer is passed in, and the base member of the returned ndarray should point to the Python object that owns the data. Then, when the ndarray is deallocated, the base-member will be DECREF'd appropriately. If you want the memory to be freed as soon as the ndarray is deallocated then simply set the OWNDATA flag on the returned ndarray.

# 14.4.3 Getting at ndarray memory and accessing elements of the ndarray

If obj is an ndarray (PyArrayObject\*), then the data-area of the ndarray is pointed to by the void\* pointer PyArray\_DATA(obj) or the char\* pointer PyArray\_BYTES(obj). Remember that (in general) this data-area may not be aligned according to the data-type, it may represent byte-swapped data, and/or it may not be writeable. Suppose the data area is aligned and in native byte-order, then how to get at a specific element of the array is determined by the array of intp variables: PyArray\_STRIDES(obj). In particular, this c-array of integers shows how many bytes must be advanced to get to the next element in each dimension. For arrays less than 4-dimensions there are PyArray\_GETPTR<k>(obj, ...) macros that make using the array strides easier where <k> is the integer 1, 2, 3, or 4. The arguments .... represent <k> non-negative integer indices into the array. For example, suppose E is a 3-dimensional ndarray. A (void\*) pointer to the element E[i,j,k] is obtained as PyArray\_GETPTR3(E, i, j, k).

As explained previously, C-style contiguous arrays and Fortran-style contiguous arrays have particular striding patterns. Two array flags (CONTIGUOUS and FORTRAN) are kept up-to-date to indicate whether or not the striding pattern of a particular array matches the C-style contiguous or Fortran-style contiguous or neither. Whether or not the striding pattern matches a standard C or Fortran one can be tested Using PyArray\_ISCONTIGUOUS(obj) and PyArray\_ISFORTRAN(obj) respectively. Most third-party libraries expect contiguous arrays. But, often it is not difficult to support general-purpose striding. I encourage you to use the striding

information in your own code whenever possible and reserve single-segment requirements for wrapping third-party code. Using the striding information provided with the ndarray rather than requiring a contiguous striding reduces copying that otherwise must be made.

## 14.5 Example

The following example shows how you might write a wrapper that accepts two input arguments (that will be converted to an array) and an output argument (that must be an array). The function returns None and updates the output array.

```
static PyObject *
example_wrapper(PyObject *dummy, PyObject *args)
   PyObject *arg1=NULL, *arg2=NULL, *out=NULL;
   PyObject *arr1=NULL, *arr2=NULL, *oarr=NULL;
    if (!PyArg_ParseTuple(args, ''000&'', &arg1, *arg2,
        &PyArrayType, *out)) return NULL;
    arr1 = PyArray_From_OTF(arg1, PyArray_DOUBLE, IN_ARRAY);
    if (arr1 == NULL) return NULL;
    arr2 = PyArray_From_OTF(arg2, PyArray_DOUBLE, IN_ARRAY);
    if (arr2 == NULL) goto fail;
    oarr = PyArray_From_OTF(out, PyArray_DOUBLE, INOUT_ARRAY);
    if (oarr == NULL) goto fail;
    /* code that makes use of arguments */
    /* You will probably need at least
       nd = PyArray_NDIM(<..>) -- number of dimensions
       dims = PyArray_DIMS(<..>) -- intp array of length nd
                                     showing length in each dim.
       dptr = (double *)PyArray_DATA(<..>) -- pointer to data.
       If an error occurs goto fail.
   Py_DECREF(arr1);
   Py_DECREF(arr2);
   Py_DECREF(oarr);
   Py_INCREF(Py_None);
   return Py_None;
```

```
fail:
    Py_XDECREF(arr1);
    Py_XDECREF(arr2);
    PyArray_XDECREF_ERR(oarr);
    return NULL;
}
```

## Chapter 15

## Beyond the Basics

## 15.1 Iterating over elements in the array

#### 15.1.1 Basic Iteration

One common algorithmic requirement is to be able to walk over all elements in a multidimensional array. The array iterator object makes this easy to do in a generic way that works for arrays of any dimension. Naturally, if you know the number of dimensions you will be using, then you can always write nested for loops to accomplish the iteration. If, however, you want to write code that works with any number of dimensions, then you can make use of the array iterator. An array iterator object is returned when accessing the .flat attribute of an array.

Basic usage is to call **PyArray\_IterNew** (array) where array is an ndarray object (or one of its sub-classes). The returned object is an array-iterator object (the same object returned by the .flat attribute of the ndarray). This object is usually cast to PyArrayIterObject\* so that its members can be accessed. The only members that are needed are iter->size which contains the total size of the array, iter->index, which contains the current 1-d index into the array, and iter->dataptr which is a pointer to the data for the current element of the array. Sometimes it is useful to access iter->ao which is a pointer to the underlying ndarray object.

After processing data at the current element of the array, the next element of the array can be obtained using **PyArray\_ITER\_NEXT**(iter). The iteration always proceeds in a C-style contiguous fashion (last index varying the fastest). The **PyArray\_ITER\_GOTO**(iter, destination) can be used to jump to a particular point in the array, where destination is an array of intp data-type with space to handle at least the number of dimensions in the underlying array. Occasionally it

is useful to use **PyArray\_ITER\_GOTO1D**(iter, index) which will jump to the 1-d index given by the value of index. The most common usage, however, is given in the following example.

```
PyObject *obj; /* assumed to be some ndarray object */
PyArrayIterObject *iter;
...
iter = (PyArrayIterObject *)PyArray_IterNew(obj);
if (iter == NULL) goto fail; /* Assume fail has clean-up code */
while (iter->index < iter->size) {
    /* do something with the data at it->dataptr */
    PyArray_ITER_NEXT(it);
}
```

You can also use **PyArrayIter\_Check(obj)** to ensure you have an iterator object and **PyArray\_ITER\_RESET(iter)** to reset an iterator object back to the beginning of the array.

It should be emphasized at this point that you may not need the array iterator if your array is already contiguous (it will work but will be a bit slower than the best code you could write). The major purpose of array iterators is to encapsulate iteration over N-dimensional arrays with arbitrary strides. If you already know your array is contiguous (Fortran or C), then simply adding the element-size to a running pointer variable will step you through the array very efficiently. In other words, code like this will probably be faster for you in the contiguous case.

```
size = PyArray_SIZE(obj);
dptr = PyArray_DATA(obj);
itemsize = PyArray_ITEMSIZE(obj);
while(size--) {
   /* do something with the data at dptr */
   dptr += itemsize;
}
```

### 15.1.2 Iterating over all but one axis

A common algorithm is to loop over all elements of an array and perform some function with each element by issuing a function call. As function calls can be time consuming, one way to speed up this kind of algorithm is to write the function so it takes a vector of data and then write the iteration so the function call is performed

for an entire dimension of data at a time. This increases the amount of work done per function call, thereby reducing the function-call over-head to a small(er) fraction of the total time. Even if the interior of the loop is performed without a function call it can be advantageous to perform the inner loop over the dimension with the highest number of elements to take advantage of speed enhancements available on micro-processors that use pipelining to enhance fundmental operations.

The **PyArray\_IterAllButAxis**(array, dim) constructs an iterator object that is modified so that it will not iterate over the dimension indicated by dim. The only restriction on this iterator object, is that the **PyArray\_Iter\_GOTO1D**(it, ind) macro cannot be used (thus flat indexing won't work either if you pass this object back to Python — so you shouldn't do this). Note that the returned object from this routine is still usually cast to PyArrayIterObject \*. All that's been done is to modify the strides and dimensions of the returned iterator to simulate iterating over array[...,0,...] where 0 is placed on the dim<sup>th</sup> dimension.

#### 15.1.3 Iterating over multiple arrays

Very often, it is desireable to iterate over several arrays at the same time. The universal functions are an example of this kind of behavior. If all you want to do is iterate over arrays with the same shape, then simply creating several iterator objects is the standard procedure. For example, the following code iterates over two arrays assumed to be the same shape and size (actually they just have to have the same total number of elements):

```
/* It is already assumed that obj1 and obj2
    are ndarrays of the same shape and size.
*/
iter1 = (PyArrayIterObject *)PyArray_IterNew(obj1);
if (iter1 == NULL) goto fail;
iter2 = (PyArrayIterObject *)PyArray_IterNew(obj2);
if (iter2 == NULL) goto fail; /* assume iter1 is DECREF'd at fail */
while (iter2->index < iter2->size) {
    /* process with iter1->dataptr and iter2->dataptr */
    PyArray_ITER_NEXT(iter1);
    PyArray_ITER_NEXT(iter2);
}
```

#### 15.1.4 Broadcasting over multiple arrays

When multiple arrays are involved in an operation, you may want to use the same broadcasting rules that the math operations (i.e. the ufuncs) use. This can be done easily using the PyArrayMultiIterObject. This is the object returned from the Python command numpy.broadcast and it is almost as easy to use from C. The function PyArray\_MultiIterNew (n, ...) is used (with n input objects in place of ...). The input objects can be arrays or anything that can be converted into an array. A pointer to a PyArrayMultiIterObject is returned. Broadcasting has already been accomplished which adjusts the iterators so that all that needs to be done to advance to the next element in each array is for PyArray\_ITER\_NEXT to be called for each of the inputs. This incrementing is automatically performed by PyArray\_MultiIter\_NEXT(obj) macro (which can handle a multiterator obj as either a PyArrayMultiObject\* or a PyObject\*). The data from input number i is available using PyArray\_MultiIter\_DATA(obj, i) and the total (broadcasted) size as PyArray\_MultiIter\_SIZE(obj). An example of using this feature follows.

```
mobj = PyArray_MultiIterNew(2, obj1, obj2);
size = PyArray_MultiIter_SIZE(obj);
while(size--) {
    ptr1 = PyArray_MultiIter_DATA(mobj, 0);
    ptr2 = PyArray_MultiIter_DATA(mobj, 1);
    /* code using contents of ptr1 and ptr2 */
    PyArray_MultiIter_NEXT(mobj);
}
```

The function PyArray\_RemoveLargest(multi) can be used to take a multiiterator object and adjust all the iterators so that iteration does not take place over the largest dimension (it makes that dimension of size 1). The code being looped over that makes use of the pointers will very-likely also need the strides data for each of the iterators. This information is stored in multi->iters[i]->strides.

## 15.2 Creating a new universal function

The umath module is a computer-generated C-module that creates many ufuncs. It provides a great many examples of how to create a universal function. Creating your own ufunc that will make use of the ufunc machinery is not difficult either. Suppose you have a function that you want to operate element-by-element over its inputs. By creating a new ufunc you will obtain a function that handles

broadcasting

- N-dimensional looping
- automatic type-conversions with limited memory usage
- optional output arrays

It is not difficult to create your own ufunc. All that is required is a 1-d loop for each data-type you want to support. Each 1-d loop must have a specific signature that is common to all universal functions. Only ufuncs for the fixed-size data-types can be used. The function call used to create a new ufunc is given below.

func A pointer to an array of 1-d functions to use. This array must be at least ntypes long. Each entry in the array must be a PyUFuncGenericFunction function. This function has the following signature. An example of a valid 1d loop function is also given.

```
void loop1d (char** args, intp* dimensions, intp* steps, void* data)
```

- args An array of pointers to the actual data for the input and output arrays. The input arguments are given first followed by the output arguments.
- **dimensions** A pointer to the size of the dimension over which this function is looping.
- **steps** A pointer to the number of bytes to jump to get to the next element in this dimension for each of the input and output arguments.
- data Arbitrary data (extra arguments, function names, etc.) that can be stored with the ufunc and will be passed in when it is called.

```
i1 += is1; i2 += is2; op += os;
}
```

- data An array of data. There should be ntypes entries (or NULL) one for every loop function defined for this ufunc. This data will be passed in to the 1-d loop. One common use of this data variable is to pass in an actual function to call to compute the result when a generic 1-d loop (e.g. PyUFunc\_d\_d) is being used.
- types An array of type-number signatures (type char). This array should be of size (nin+nout)\*ntypes and contain the data-types for the corresponding 1-d loop. The inputs should be first followed by the outputs. For example, suppose I have a ufunc that supports 1 integer and 1 double 1-d loop (length-2 func and data arrays) that takes 2 inputs and returns 1 output that is always a complex double, then the types array would be

```
char my_sigs[] = \
{PyArray_INT, PyArray_INT, PyArray_CDOUBLE,
PyArray_DOUBLE, PyArray_DOUBLE, PyArray_CDOUBLE};
```

The bit-width names can also be used (e.g. PyArray\_INT32, PyArray\_COMPLEX128) if desired.

**ntypes** The number of data-types supported. This is equal to the number of 1-d loops provided.

nin The number of input arguments.

nout The number of output arguments.

- identity Either PyUFunc\_One, PyUFunc\_Zero, PyUFunc\_None. This specifies what should be returned when an empty array is passed to the reduce method of the ufunc.
- **name** A NULL-terminated string providing the name of this ufunc (should be the Python name it will be called).
- doc A documentation string for this ufunc (will be used in generating the response to <ufunc\_name>.\_\_doc\_\_). Do not include the function signature or the name as this is generated automatically.
- **check\_return** Not presently used, but this integer value does get set in the structure-member of similar name.

The returned usunc object is a callable Python object. It should be placed in a (module) dictionary under the same name as was used in the name argument to the usunc-creation routine. The following example is adapted from the umath module:

```
static PyUFuncGenericFunction atan2_functions[]=\
    {PyUFunc_ff_f, PyUFunc_dd_d,
     PyUFunc_gg_g, PyUFunc_00_0_method};
static void* atan2_data[]=\
    {(void *)atan2f,(void *) atan2,
     (void *)atan21,(void *)"arctan2"};
static char atan2_signatures[]=\
    {PyArray_FLOAT, PyArray_FLOAT, PyArray_FLOAT,
     PyArray_DOUBLE, PyArray_DOUBLE,
     PyArray_DOUBLE, PyArray_LONGDOUBLE,
     PyArray_LONGDOUBLE, PyArray_LONGDOUBLE
     PyArray_OBJECT, PyArray_OBJECT,
     PyArray_OBJECT};
/* in the module initialization code */
PyObject *f, *dict, *module;
dict = PyModule_GetDict(module);
f = PyUFunc_FromFuncAndData(atan2_functions,
    atan2_data, atan2_signatures, 4, 2, 1,
    PyUFunc_None, "arctan2",
    "a safe and correct arctan(x1/x2)", 0);
PyDict_SetItemString(dict, "arctan2", f);
Py_DECREF(f);
. . .
```

## 15.3 User-defined data-types

NumPy comes with 21 builtin data-types. While this covers a large majority of possible use cases, it is conceivable that a user may have a need for an additional data-type. There is some support for adding an additional data-type into the NumPy system. This additional data-type will behave much like a regular data-type except ufuncs must have 1-d loops registered to handle it separately. Also checking for whether or not other data-types can be cast "safely" to and from this new type or

not will always return "can cast."

#### 15.3.1 Adding the new data-type

To begin to make use of the new data-type, you need to first define a new Python type to hold the scalars of your new data-type. It should be acceptable to inherit from one of the array scalars if your new type has a binary compatible layout. This will allow your new data type to have the methods and attributes of array scalars. New data-types must have a fixed memory size (if you want to define a data-type that needs a flexible representation, like a variable-precision number, then use a pointer to the object as the data-type). The memory layout of the object structure for the new Python type must be PyObject\_HEAD followed by the fixed-size memory needed for the data-type. For example, a suitable structure for the new Python type is:

```
typedef struct {
    PyObject_HEAD;
    some_data_type obval;
    /* the name can be whatever you want */
} PySomeDataTypeObject;
```

After you have defined a new Python type object, you must then define a new PyArray\_Descr structure whose typeobject member will contain a pointer to the data-type you've just defined. In addition, the required functions in the ".f" member must be defined: nonzero, copyswap, copyswapn, setitem, getitem, and cast. The more functions in the ".f" member you define, however, the more useful the new data-type will be. It is very important to intialize unused functions to NULL. This can be achieved using **PyArray\_InitializeNewArrFuncs**(f).

Once a new PyArray\_Descr structure is created and filled with the needed information and useful functions you call **PyArray\_RegisterDataType**(new\_descr). The return value from this call is an integer providing you with a unique type\_number that specifies your data-type. This type number should be stored and made available by your module so that other modules can use it to recognize your data-type.

## 15.3.2 Registering a casting function

You may want to allow builtin (and other user-defined) data-types to be cast automatically to your data-type. In order to make this possible, you must register a casting function with the data-type you want to be able to cast from. This requires writing low-level casting functions for each conversion you want to support and

then registering these functions with the data-type descriptor. A low-level casting function has the signature.

```
castfunc (void) (void* from, void* to, intp n, void* fromarr, void* toarr)
```

Cast n elements from one type to another. The data to cast from is in a contiguous, correctly-swapped and aligned chunk of memory pointed to by from. The buffer to cast to is also contiguous, correctly-swapped and aligned. The fromarr and toarr arguments should only be used for flexible-element-sized arrays (string, unicode, void).

An example castfunc is

This could then be registered to convert doubles to floats using the code

#### 15.3.3 Registering coercion rules

By default, all user-defined data-types are not presumed to be safely castable to any builtin data-types. In addition builtin data-types are not presumed to be safely castable to user-defined data-types. This situation limits the ability of user-defined data-types to participate in the coercion system used by ufuncs and other times when automatic coercion takes place in NumPy. This can be changed by registering data-types as safely castable from a particlar data-type object. The function PyArray\_RegisterCanCast (from\_descr, totype\_number, scalarkind) should be used to specify that the data-type object from\_descr can be cast to the data-type with type number totype\_number. If you are not trying to alter scalar coercion rules, then use PyArray\_NOSCALAR for the scalarkind argument.

If you want to allow your new data-type to also be able to share in the scalar coercion rules, then you need to specify the scalarkind function in the data-type

object's ".f" member to return the kind of scalar the new data-type should be seen as (the value of the scalar is available to that function). Then, you can register data-types that can be cast to separately for each scalar kind that may be returned from your user-defined data-type. If you don't register scalar coercion handling, then all of your user-defined data-types will be seen as **PyArray\_NOSCALAR**.

#### 15.3.4 Registering a ufunc loop

You may also want to register low-level usunc loops for your data-type so that an ndarray of your data-type can have math applied to it seamlessly. Only one usunc loop for a given user-defined data-type may be registered for each usunc. Registering a new loop silently replaces any previously registered loops for that (user-defined) data-type.

Before you can register a 1-d loop for a ufunc, the ufunc must be previously created. Then you call **PyUFunc\_RegisterLoopForType**(...) with the information needed for the loop. The return value of this function is 0 if the process was successful and -1 with an error condition set if it was not successful.

**ufunc** The ufunc do attach this loop to.

**usertype** The user-defined type this loop should be indexed under (only one loop per ufunc per user type is allowed). This number must be a user-defined type or an error occurs.

function The ufunc inner loop. This function must have the signature as explained in Section 15.2.

arg\_types (optional) If given, this should contain a persistent array of integers of at least size ufunc.nin + ufunc.nout containing the data-types expected by the loop function. If this is NULL, then it will be assumed that all data-types are of type usertype.

data (optional) Specify any optional data needed by the function which will be passed when the function is called.

## 15.4 Subtyping the ndarray in C

One of the lesser-used features that has been lurking in Python since 2.2 is the ability to sub-class types in C. This facility is one of the important reasons for

basing NumPy off of Numeric. A sub-type in C allows much more flexibility with regards to memory management. Sub-typing in C is not difficult even if you have only a rudimentary understanding of how to create new types for Python. While it is easiest to sub-type from a single parent type, sub-typing from multiple parent types is also possible. Multiple inheritence in C is generally less useful than it is in Python because a restriction on Python sub-types is that they have a binary compatible memory layout. Perhaps for this reason, it is somewhat easier to sub-type from a single parent type.

All C-structures corresponding to Python objects must begin with PyObject\_HEAD (or PyObject\_VAR\_HEAD). In the same way, any sub-type must have a C-structure that begins with exactly the same memory layout as the parent type (or all the parent types). The reason for this is that Python may attempt to access a member of the sub-type structure as if it had the parent structure (i.e. it will cast a given pointer to a pointer to the parent structure and then dereference one of it's members). If the memory layouts are not compatible, then this attempt will cause unpredictable behavior (eventually leading to a memory violation and program crash).

One of the elements in PyObject\_HEAD is a pointer to a type-object structure. A new Python type is created by creating a new type-object structure and populating it with functions and pointers to describe the desired behavior of the type. Typically, a new C-structure is also created to contain the instance-specific information needed for each object of the type as well. For example, &PyArray\_Type is a pointer to the type-object table for the ndarray while PyArrayObject \* is a pointer to a particular instance of an ndarray (one of the members of the ndarray structure is, in turn, a pointer to the type-object table &PyArray\_Type). Finally PyType\_Ready(<pointer\_to\_type\_object>) must be called for every new Python type.

## 15.4.1 Creating sub-types

To create a sub-type, a similar proceedure must be followed except only behaviors that are different require new entries in the type-object structure. All other entires can be NULL and will be filled in by **PyType\_Ready** with appropriate functions from the parent type(s). In particular, to create a sub-type in C follow these steps:

1. If needed create a new C-structure to handle each instance of your type. A typical C-structure would be

```
typedef _new_struct {
    PyArrayObject base;
```

```
/* new things here */
} NewArrayObject;
```

Notice that the full PyArrayObject is used as the first entry in order to ensure that the binary layout of instances of the new type is identical to the PyArrayObject.

- 2. Fill in a new Python type-object structure with pointers to new functions that will over-ride the default behavior while leaving any function that should remain the same unfilled (or NULL). The tp\_name element should be different.
- 3. Fill in the tp\_base member of the new type-object structure with a pointer to the (main) parent type object. For multiple-inheritance, also fill in the tp\_bases member with a tuple containing all of the parent objects in the order they should be used to define inheritance. Remember, all parent-types must have the same C-structure for multiple inheritance to work properly.
- 4. Call **PyType\_Ready**(<pointer\_to\_new\_type>). If this function returns a negative number, a failure occurred and the type is not initialized. Otherwise, the type is ready to be used. It is generally important to place a reference to the new type into the module dictionary so it can be accessed from Python.

More information on creating sub-types in C can be learned by reading PEP 253 (available at http://www.python.org/dev/peps/pep-0253).

### 15.4.2 Specific features of ndarray sub-typing

#### 15.4.2.1 The \_array\_finalize\_ attribute

Several array-creation functions of the ndarray allow specification of a particular sub-type to be created. This allows sub-types to be handled seamlessly in many routines. When a sub-type is created in such a fashion, however, neither the \_\_new\_\_ method nor the \_\_init\_\_ methods gets called. Instead, the sub-type is allocated and the appropriate instance-structure members are filled in. Finally, the \_\_array\_finalize\_\_ attribute is looked-up in the object dictionary. If it is present and not None, then it can be either a CObject containing a pointer to a PyArray\_FinalizeFunc or it can be a method taking a single argument (which could be None).

If the **\_\_array\_finalize**\_\_ attribute is a CObject, then the pointer must be a pointer to a function with the signature:

```
(int) (PyArrayObject *, PyObject *)
```

The first argument is the newly created sub-type. The second argument (if not NULL) is the "parent" array (if the array was created using slicing or some other operation where a clearly-distinguishable parent is present). This routine can do anything it wants to. It should return a -1 on error and 0 otherwise.

If the <u>\_array\_finalize\_</u> attribute is not None nor a CObject, then it must be a method that takes the parent array as the argument (which could be None if there is no parent), and returns nothing. Errors in this method will be caught, however.

#### 15.4.2.2 The \_array\_priority\_ attribute

This attribute allows simple but flexible determination of which sub-type should be considered "primary" when an operation involving two or more sub-types arises. In operations where different sub-types are being used, the sub-type with the largest \_array\_priority\_ attribute will determine the sub-type of the output(s). If two sub-types have the same \_array\_priority\_ then the sub-type of the first argument determines the output. The default \_array\_priority\_ attribute returns a value of 0.0 for the base ndarray type and 1.0 for a sub-type.

#### 15.4.2.3 The \_array\_wrap\_ attribute

This attribute of the sub-type

## Chapter 16

## Third-party tools

## 16.1 Calling other compiled libraries from Python

While Python is a great language and a pleasure to code in, its dynamic nature results in overhead that can cause some code (*i.e.* raw computations inside of for loops) to be up 10-100 times slower than equivalent code written in a static compiled language. For scientific computation that extra slow-down can often not be spared for certain portions of your code. Therefore one of the most common needs is to call out from Python code to a fast, machine-code routine compiled using C/C++ or Fortran. The fact that this is relatively easy to do is a big reason why Python is such an excellent high-level language for scientific and engineering programming.

Their are two basic approaches to calling compiled code: writing an extension module that is then imported to Python using the import command, or calling the subroutine directly from Python using the ctypes module (included in the standard distribution with Python 2.5). The first method is the most common (but with the inclusion of ctypes into Python 2.5 this status may change). The ctypes methods exposes a raw interface to the compiled code and is therefore not tolerant of programmer mistakes which can easily cause Python to crash. Robust use of the ctypes module typically involves an additional layer of Python code in order to check the data types and array bounds of objects passed to the underlying subroutine. This additional layer of checking might make the interface slightly slower than the C-written interface. However, it should be neglible if the C-routine being called is doing any significant amount of work. If you are a great Python programmer with weak C-skills, ctypes may be an easy way for you to write a useful interface to a library of compiled code

#### 16.1.1 Hand-generated wrappers

Except for c-types, all of the other methods for calling C and Fortran code make use

#### 16.1.2 Using f2py

## 16.2 Other tools installed separately

- 16.2.1 Using weave
- 16.2.2 Using PyRex

### 16.2.3 Using ctypes

Ctypes is a python extension module (downloaded separately for Python <2.5 and included with Python 2.5) that allows you to call an arbitrary function in a shared library directly from Python. This approach allows you to interface with C-code directly from Python. The drawback, however, is that coding mistakes can lead to segmentation violations (crashes) very easily because there is little type or bounds checking done on the parameters. This is especially true when array data is passed in as a pointer to a raw memory location. The responsibility is then on you that the subroutine will not access memory outside the actual array. But, if you don't mind living a little dangerously, ctypes can be an effective tool for quickly taking advantage of a large shared library.

#### 16.2.4 Other tools

- 16.2.4.1 SWIG
- 16.2.4.2 Boost
- 16.2.4.3 SIP
- 16.2.4.4 PyFort
- 16.2.4.5 Instant

## Chapter 17

## Code Explanations

This Chapter attempts to explain the logic behind some of the new pieces of code. The purpose behind these explanations is to enable somebody to be able to understand the ideas behind the implementation a little better so that they can be improved on, borrowed from, or optimized.

## 17.1 Code generation

Besides the use of .src files whose repeat blocks are processed prior to compilation, python-generated code is used to construct the umath module and

The code-generation mechanism has been explained in the previous section.

- 17.2 Array Scalars
- 17.3 N-d Array Iteration
- 17.4 Advanced Indexing
- 17.5 Universal Functions

## Index

absolute, 145 add, 144 arccos, 147 arccosh, 148 arcsin, 147 arcsinh, 148 arctan, 147 arctan2, 147 arctanh, 148

conj, 145 conjugate, 145 cos, 147 cosh, 147

divide, 144

 $\exp, 145$ 

floor\_divide, 144

hypot, 147

 $\log, 145$  $\log 10, 146$ 

multiply, 144

negative, 145

power, 145

remainder, 145

 $\sin, 147$ 

sinh, 147 sqrt, 146 subtract, 144

tan, 147 tanh, 148 true\_divide, 144