

**CS2204**

Program Design and Data  
Structures for Scientific  
Computing

# Announcements

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- **Project #2**
  - We will be creating another class that allows us to represent & manipulate DNA strands
  - We will use a list as our underlying storage container
  - Our goal is to have a faster/smaller representation by utilizing the mutability of lists (as compared to the immutability of strings)
    - You should use mutation operations on your list whenever possible
    - Avoid operations that create new lists



# Matrix Programming

## Indexing

By Richard T. Guy



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Now we'll have a look at some of the ways you can index arrays.

It may not seem important at first, but as we'll see, clever indexing allows you to avoid writing loops,

- reduces the size of your code
- makes it more efficient

# Can *slice* arrays like lists or strings

```
>>> block
```

```
array([[ 10, 20, 30, 40],
       [110, 120, 130, 140],
       [210, 220, 230, 240]])
```

```
>>> block[0:3, 0:2]
```

```
array([[ 10, 20],
       [110, 120],
       [210, 220]])
```

	← 0:2 →			
↑ 0:3 ↓	10	20	30	40
	110	120	130	140
	210	220	230	240

Can *slice* arrays like lists or strings

```
>>> block
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```
array([[ 10, 20, 30, 40],
       [110, 120, 130, 140],
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>>> block[0:3, 0:2]
```

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array([[ 10, 20],
       [110, 120],
       [210, 220]])
```

Are slices **aliases** or copies?

	← 0:2 →			
↑ 0:3 ↓	10	20	30	40
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Can *slice* arrays like lists or strings

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>>> block
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array([[ 10, 20, 30, 40],
       [110, 120, 130, 140],
       [210, 220, 230, 240]])
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```
>>> block[0:3, 0:2]
```

```
array([[ 10, 20],
       [110, 120],
       [210, 220]])
```

Are slices aliases or copies?

- *They are aliases*

← 0:2 →

	10	20	30	40
↑	110	120	130	140
0:3	210	220	230	240
↓				

## Can assign to slices

```
>>> block[1, 1:3] = 0
```

```
>>> block
```

```
array([[ 10,  20,  30,  40],  
       [110,  0,  0, 140],  
       [210, 220, 230, 240]])
```



Slice on both sides to shift data

```
>>> vector = array([10, 20, 30, 40])
```

```
>>> vector[0:3] = vector[1:4]
```

```
>>> vector
```

```
array([20, 30, 40, 40])
```



Not overwritten

Slice on both sides to shift data

```
>>> vector = array([10, 20, 30, 40])
```

```
>>> vector[0:3] = vector[1:4]
```

```
>>> vector
```

```
array([20, 30, 40, 40])
```



Not overwritten

Is this easier to understand than a loop?

Slice on both sides to shift data

```
>>> vector = array([10, 20, 30, 40])
```

```
>>> vector[0:3] = vector[1:4]
```

```
>>> vector
```

```
array([20, 30, 40, 40])
```



Not overwritten

Is this easier to understand than a loop?

- In most cases, yes

Slice on both sides to shift data

```
>>> vector = array([10, 20, 30, 40])
```

```
>>> vector[0:3] = vector[1:4]
```

```
>>> vector
```

```
array([20, 30, 40, 40])
```



Not overwritten

Is this easier to understand than a loop?

- In most cases, yes
- Particularly when shifting up, when a loop would have to be written to count down

# Can use lists or arrays as subscripts

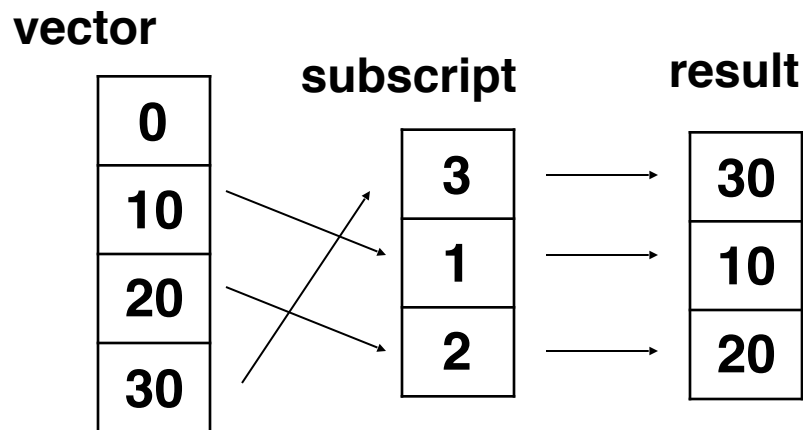
```
>>> vector
```

```
array([0, 10, 20, 30])
```

```
>>> subscript = [3, 1, 2]
```

```
>>> vector[ subscript ]
```

```
array([30, 10, 20])
```



Also works in multiple dimensions

Though operation may not be obvious

```
>>> square = numpy.array([[5, 6], [7, 8]])
```

```
>>> square[ [1] ]
```

```
array([[7, 8]])
```

~~LOOPS~~

# Comparisons

```
>>> vector
```

```
array([0, 10, 20, 30])
```

```
>>> vector < 25
```

```
array([ True,  True,  True, False ],  
      dtype=bool)
```

← Data type is Boolean

The possible state values for a boolean object are True and False with the standard boolean operators, and, or, and not.

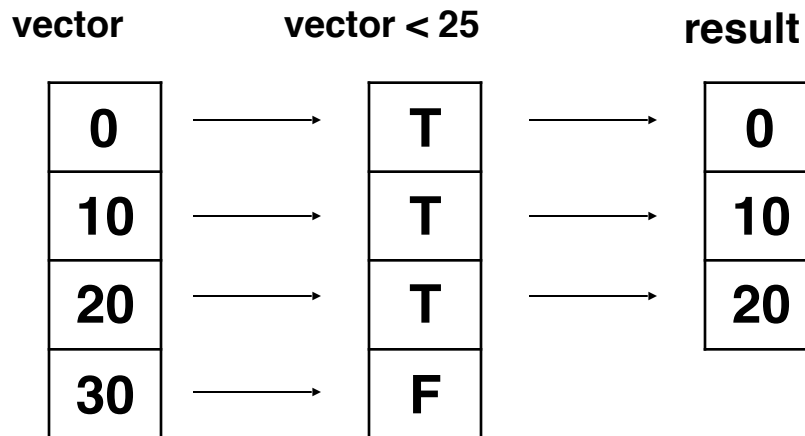
# Use a Boolean subscript as a *mask*

```
>>> vector
```

```
array([0, 10, 20, 30])
```

```
>>> vector[ vector < 25 ]
```

```
array([0, 10, 20])
```





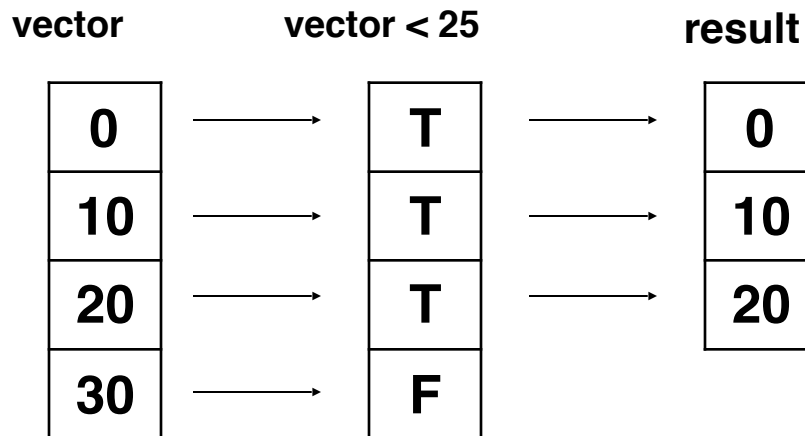
Use a Boolean subscript as a *mask*

```
>>> vector
```

```
array([0, 10, 20, 30])
```

```
>>> vector[ vector < 25 ]
```

```
array([0, 10, 20])
```



Copy or  
alias?

# Use masking for assignment

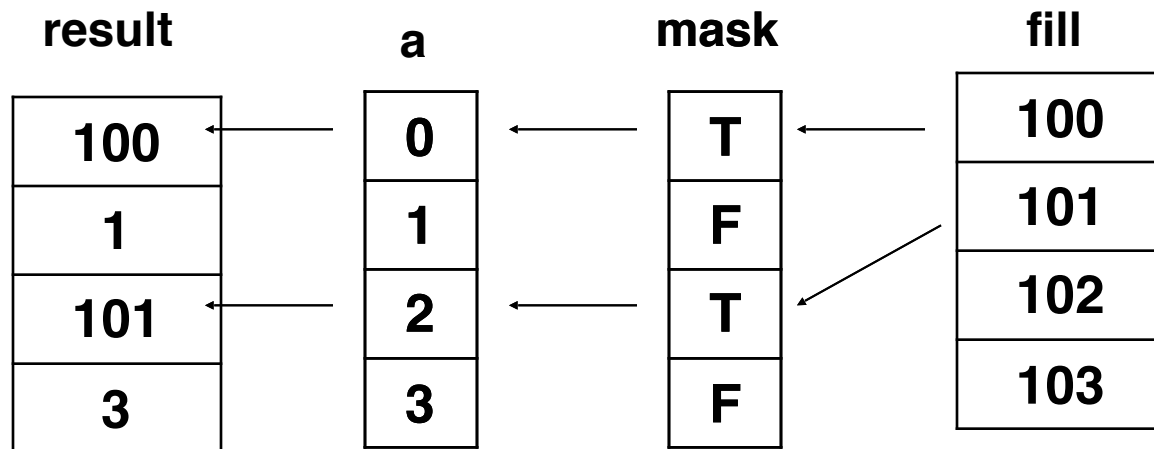
```
>>> a = array([0, 1, 2, 3])
```

```
>>> mask = array([True, False, True, False])
```

```
>>> a[mask] = array([100, 101, 102, 103])
```

```
>>> a
```

```
array([100, 1, 101, 3])
```



Taken  
in  
order

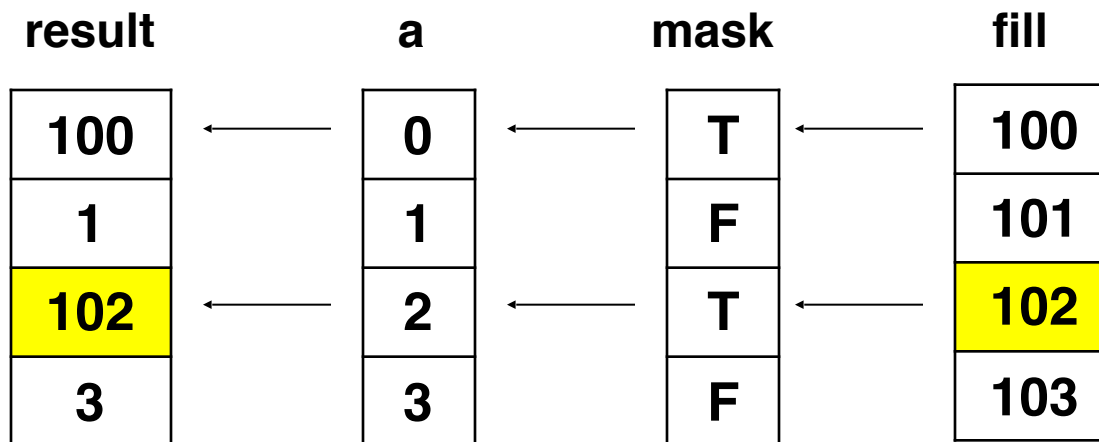
The **putmask** function works slightly differently

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

```
>>> putmask(a, mask,
            array([100, 101, 102, 103]))
```

```
>>> a
```

```
array([100, 1, 102, 3])
```



Taken  
where  
True

Python does not allow objects to re-define the meaning of **and/or/not**

```
>>> vector = array([0, 10, 20, 30])
```

```
>>> vector <= 20
```

```
array([True, True, True, False], dtype=bool)
```

```
>>> (vector <= 20) and (vector >= 20)
```

**ValueError: The truth value of an array with more than one element is ambiguous.**

Use **logical\_and** / **logical\_or** functions

```
>>> logical_and(vector <= 20, vector >= 20)  
array([False, False, True, False], dtype=bool)
```

Or use bitwise operations: **|** for **or**, **&** for **and**

```
>>> (vector <= 20) & (vector >= 20)  
array([False, False, True, False], dtype=bool)
```

## Use **where** instead of **if/else**

```
>>> vector = array([10, 20, 30, 40])
```

```
>>> where(vector < 25, vector, 0)
```

```
array([10, 20, 0, 0])
```

```
>>> where(vector > 25, vector/10, vector)
```

```
array([10, 20, 3, 4])
```

## Review:

- Arrays can be sliced
- Or subscripted with vectors of indices
- Or masked with conditionals

## Review:

- Arrays can be sliced
- Or subscripted with vectors of indices
- Or masked with conditionals

~~LOOPS~~



NumPy arrays make operations on rectangular data easy

But they are not quite mathematical matrices

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

```
>>> a * a
```

Operators act *element-wise*

NumPy arrays make operations on rectangular data easy

But they are not quite mathematical matrices

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

```
>>> a * a
```

```
array([[ 1,  4],  
       [ 9, 16]])
```

Operators act *element-wise*

So this does what you think

```
>>> a + a
```

And NumPy is sensible about *scalar* values

```
>>> a + 1
```

So this does what you think

```
>>> a + a  
array([[ 2,  4],  
       [ 6,  8]])
```

And NumPy is sensible about *scalar* values

```
>>> a + 1
```

So this does what you think

```
>>> a + a  
array([[ 2,  4],  
       [ 6,  8]])
```

And NumPy is sensible about *scalar* values

```
>>> a + 1  
array([[ 2,  3],  
       [ 4,  5]])
```

Lots of useful utilities - Pay attention to dimensions!

**>>> sum(a)**

**>>> sum(a, 0)**

**>>> sum(a, 1)**

	1	→	
0	1	2	3
↓	3	4	7
	4	6	

Lots of useful utilities - Pay attention to dimensions!

```
>>> sum(a)
```

10

```
>>> sum(a, 0)
```

	1	→	
0	1	2	3
↓	3	4	7
	4	6	

```
>>> sum(a, 1)
```

Lots of useful utilities - Pay attention to dimensions!

```
>>> sum(a)
```

10

```
>>> sum(a, 0)
```

array([4, 6])

```
>>> sum(a, 1)
```

	1	→	
	1	2	3
0	3	4	7
↓			
	4	6	



Lots of useful utilities - Pay attention to dimensions!

```
>>> sum(a)
10
```

```
>>> sum(a, 0)
array([4, 6])
```

```
>>> sum(a, 1)
array([3, 7])
```

	1	→	
	1	2	3
0	3	4	7
↓			
	4	6	

```
>>> data[:, 0] # t0 count for all patients  
array([1., 0., 0., 2., 1.])  
  
>>> data[0, :] # all samples for patient 0  
array([1., 3., 3., 5., 12., 10., 9.])
```

Why are these 1D rather than 2D?

## Example: Disease statistics

- We have 5 patients with 7 samples each
- One row per patient
- Columns are hourly responsive T cell counts

```
>>> data[:, 0] # t0 count for all patients
```

```
array([1., 0., 0., 2., 1.])
```

```
>>> data[0, :] # all samples for patient 0
```

```
array([1., 3., 3., 5., 12., 10., 9.])
```

Why are these 1D rather than 2D?

```
>>> mean(data)
```

```
6.8857
```

Gives us the average T cell count for all patients at all times

Intriguing, but not particularly meaningful

```
>>> mean(data, 0) # over time
```

```
array([ 0.8, 2.6, 4.4, 6.4, 10.8, 11., 12.2])
```

The mean of the data along axis 0 gives us the average across all patients for each hour.

This is much more useful: it is the "normal" progress of the disease.

```
>>> mean(data)
```

```
6.8857
```

Gives us the average T cell count for all patients at all times

Intriguing, but not particularly meaningful

```
>>> mean(data, 1) # per patient
```

```
array([ 6.14,  4.28, 16.57,  2.14,  5.29])
```

The mean of the data along axis 1 gives us the average T cell count per patient across all times.

This could be useful if we need to normalize the data.

Select the data for people who started with  
a responsive T cell count of 0

```
>>> data[:, 0]
array([1., 0., 0., 2., 1.])

>>> data[:, 0] == 0.
array([False, True, True, False, False],
      dtype=bool)

>>> data[ data[:, 0] == 0 ]
array([[ 0.,  1.,  2.,  4.,  8.,  7.,  8.],
       [ 0.,  4., 11., 15., 21., 28., 37.]])
```

Find the mean T cell count over time for people who started with a count of 0

>>>

**data[:, 0]**

Column 0

Find the mean T cell count over time for people who started with a count of 0

>>>

**data[:, 0] == 0**



Column 0 is 0



Find the mean T cell count over time for people who started with a count of 0

```
>>> data[ data[:, 0] == 0 ]
```



Rows where column 0 is 0

Find the mean T cell count over time for people who started with a count of 0

```
>>> mean(data[ data[:, 0] == 0 ], 0)
```



Mean along axis 0 of  
rows where column 0 is 0

Find the mean T cell count over time for people who started with a count of 0

```
>>> mean(data[ data[:, 0] == 0 ], 0)  
array([ 0., 2.5, 6.5, 9.5, 14.5, 17.5, 22.5])
```

Find the mean T cell count over time for people who started with a count of 0

```
>>> mean(data[ data[:, 0] == 0 ], 0)  
array([ 0., 2.5, 6.5, 9.5, 14.5, 17.5, 22.5])
```

Key to good array programming: no loops!  
Just as true for MATLAB or R as for NumPy

# What about "real" matrix multiplication?

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

```
>>> dot(a, a)
```

```
>>> v = arange(3) # [0, 1, 2]
```

```
>>> dot(v, v)      # 0*0 + 1*1 + 2*2
```

# What about "real" matrix multiplication?

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

```
>>> dot(a, a)
```

```
array([[ 7, 10],  
       [15, 22]])
```

```
>>> v = arange(3) # [0, 1, 2]
```

```
>>> dot(v, v)     # 0*0 + 1*1 + 2*2
```

# What about "real" matrix multiplication?

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

```
>>> dot(a, a)
```

```
array([[ 7, 10],  
       [15, 22]])
```

```
>>> v = arange(3) # [0, 1, 2]
```

```
>>> dot(v, v)      # 0*0 + 1*1 + 2*2
```

```
5
```

Dot product only works for sensible shapes

```
>>> dot(ones((2, 3)), ones((2, 3)))
```

**ValueError: objects are not aligned**

NumPy does not distinguish row/column vectors

```
>>> v = array([1, 2])
```

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

```
>>> dot(v, a)
```

```
array([ 7, 10])
```

```
>>> dot(a, v)
```

```
array([ 5, 11])
```

$$\begin{bmatrix} 1 & 2 \\ 3 & 4 \end{bmatrix} \begin{bmatrix} 1 \\ 2 \end{bmatrix} = \begin{bmatrix} 7 \\ 10 \end{bmatrix}$$

$$\begin{bmatrix} 1 & 2 \\ 3 & 4 \end{bmatrix} \begin{bmatrix} 1 & 2 \end{bmatrix} = \begin{bmatrix} 5 & 11 \end{bmatrix}$$



Can also use the **matrix** subclass of **array**

```
>>> m = matrix([[1, 2], [3, 4]])
```

```
>>> m
```

```
matrix([[ 1, 2],
         [ 3, 4]])
```

```
>>> m*m # '*' does matrix mult for matrices
```

```
matrix([[ 7, 10],
         [15, 22]])
```

Use **matrix(a)** or **array(m)** to convert

Which should you use?

If your problem is linear algebra, **matrix** will probably be more convenient

- Treats vectors as  $N \times 1$  matrices

Otherwise, use **array**

- Especially if you're representing grids, rather than mathematical matrices

Always look at

[http://www.scipy.org/Numpy\\_Example\\_List\\_With\\_Doc](http://www.scipy.org/Numpy_Example_List_With_Doc)

before writing any functions of your own

**conjugate**

**histogram**

**convolve**

**lstsq**

**correlate**

**npv**

**diagonal**

**roots**

**fft**

**solve**

**gradient**

**svd**

Fast...

...and someone else has debugged them

I will also post an extended example with these lectures notes

# Arrays vs. Lists

---

- Arrays and lists have many similarities, but there are also some important differences
- Similarities between arrays and lists:
  - Both are mutable: both can have elements reassigned in place
  - Arrays and lists are indexed and sliced identically
  - The len command works just as well on arrays as anything else
  - Arrays and lists both have sort and reverse attributes
- Differences between arrays and lists:
  - With arrays, the + and \* signs do not refer to concatenation or repetition, but are element-wise addition/multiplication

# Arrays vs. Lists

---

- The biggest difference between arrays and lists is speed; it's much faster to carry out operations on arrays (and all the terms therein) than on each term in a given list.
- Example: take the following script:

```
tt1 = time.clock()
sarr = 1.*arange(0,10001)/10000
sinarr = sin(sarr)
tt2 = time.clock()
slist = []; sinlist = []
for i in range(10001):
    slist.append(1.*i/10000)
    sinlist.append(sin(slist[i]))
tt3 = time.clock()
```
- $tt2 - tt1$  (the time to process with arrays) is 0.0007 seconds, while  $tt3 - tt2$  (the time to process with lists) is 0.027 seconds.