Numpy Note

1. basics

1.1 type

- numpy has its own datatypes.
 - float: np.float16, np.float32, np.float64
 - int: np.int8, np.int16, np.int32, np.int64
 - unsigned_int: np.uint8, np.uint16, np.uint32,np.uint64
 - complex: np.complex64, np.complex128
 - bool: np.bool
 - byte: np.byte, signed char np.ubyte: unsigned char
 - more conventioned name from C: np.short, np.long, np.double etc.
- all above datatype can be used as function to convert type np.float32(1.0)
- like pandas, for given np.array.astype('str') convert types
- np.ndarray.view(dtype = None): create a new view of the same data.
 - a.view(some_dtype) or a.view(dtype=some_dtype) constructs a view of the array's memory with a different data-type. This can cause a reinterpretation of the bytes of memory.
 - a.view(ndarray_subclass) or a.view(type=ndarray_subclass) just returns an instance of ndarray_subclass that looks at the same array (same shape, dtype, etc.) This does not cause a reinterpretation of the memory.

1.2 arrays:

- numpy's basic object is np.ndarray
- Common ways to build array:
 - from built-in python array_like object, use np.array()
 - numpy function to create arrays:
 - np.zeros(size, dtype): zero array
 - np.arange([start,]stop,[step,], dtype): syntax like range() but create corresponding array

- np.linspace(start, stop, num=50, endpoint= True, retsetp=False, dtype, axis=0): numpy version of matlab linspace
 - start=, stop=: beginning and end point
 - num: int, number of points
 - endpoint: bool, if include stop
 - retsetp: bool, if true, return a tuple (array, step)
 - axis: axis to expand result. relevant only if start, stop are ndarray themselves.
- np.indices(dimensions, dtype, sparse): return indices along all axies. Specifically, if the dimensions = (d0,d1,...,d_n-1), the result is a np. ndarray with shape (n,d0,d1,....d_n-1). The first dimension determines the axis for which indices are provided.

Example: (See also stackoverflow)

• other numpy functions' result, like functions in the random module below.

1.3 access array

Accessor method takes an array_like object to access part of data. It can be any array_like object other than tuple, which is reserved for special use.

There is no single index for multidimensional array as in matlab.

- basic accessor is bracket np.ndarray[].
 - accept single integer, can be positive/negative, a [-1]
 - accept multidimensional index in the same bracket, a[2,3] if length of number < array dims, return subarray
 - For 2d array, a[2,3] == a[2][3] but the former is more efficient, just like in pandas
 - range slice works as expected, a [:-1] means exclude the last element. Note that it provide new view of data but not new copy

- array accept index arrays like matlab, ndarray [index_array], this returns copy of data, this is one of the advanced indexing
 - for 1d array: return a new array of mapping: index -> element the index array can be multidimensional
 - for nd array: used less. Suppose array shape is (d0,d1,...d_n-1). The syntax should be ndarray[index_array_1,index_array_2,...], to select certain dimension fully, use:
 - if each index array has the same shape x (shape match) and there are n index arrays. The result is an array of shape x with mapping ndarray[index_1,index_2,..index_n] -> element
 - if the index array does not have the same shape, numpy attempts broadcasting the match the shape
 - if there are l < n index arrays (partial indexing): the omitting axis's elements are all selected. y[np.array([0,2,4])] selects the 1,3,5 row of the 2d array y.
- array accepts boolean array, leads to new copy of data. this is the other one of the advanced indexing
 - if boolean array has the same shape as the input array. It returns 1d array with elements of true. This is equivalent to y[np.nonzero(bool_array)]
 - if boolean array has less dimension than the array, The result dim are all selected. The returned array has shape [# of true element, omitted dims] Note that any selected dimension should have the same shape between input array and index bool array.

For example, using a 2-D boolean array of shape (2,3) with four True elements to select rows from a 3-D array of shape (2,3,5) results in a 2-D result of shape (4,5). But a 2-D boolean array of shape (2,4) is invalid

• np.newaxis can be used to expand array. For example

```
1 y = np.arange(35).reshape(5,7)
2 y[:,np.newaxis,:].shape # 5,1,7
```

• ... may be used to indicate selecting in full any remaining unspecified dimensions.

```
1 y = np.arange(30).reshape(1,1,2,3,5)
2 y[0,...,1,1].shape # 1,2
```

• all access methods if used in assignment changes original data.

```
1 >>> x = np.arange(0, 50, 10)
2 >>> x
3 array([ 0, 10, 20, 30, 40])
4 >>> x[np.array([1, 1, 3, 1])] += 1
5 >>> x
6 array([ 0, 11, 20, 31, 40])
```

Where people expect that the 1st location will be incremented by 3. In fact, it will only be incremented by 1. The reason is because a new array is extracted from the original (as a temporary) containing the values at 1, 1, 3, 1, then the value 1 is added to the temporary, and then the temporary is assigned back to the original array. Thus the value of the array at x[1]+1 is assigned to x[1] three times, rather than being incremented 3 times.

tuple when supplied is not interpreted as an index array but a list of indices, i.e. indicate single element access described in the first section. a[(1,1)] == a[1,1]

1.4 array methods:

Shape manipulation

- ndarray. shape: attribute, shapes of the array
- np.reshape(array,newshape, order = {'C','F','A'}): reshape, return new view whenever possible.
 - newshape is a tuple
 - order specifies how ndarray is arranged in memory, row-major('C') or column major ('F')
 - also available as ndarray.reshape()
- ndarray.flat: 1d iterator over array, a numpy.flattier instance
- ndarray.flatten(order =): collapse data into 1d array

Transpose

- ndarray. T: transpose of array
- np.transpose(array, axes=): permute dimensions. by default, reverse dimensions.
 - axes =: list of int, order of axes, starting from 0
- np.swapaxes(array, axis1,axis2): swap axes,

• np.moveaxis(a, start, dest): move axis at start to dest, keep other axes' relative orders intact.

Change Dimension:

- ndarray.ndims: attribute, number of dims
- np.expand_dims(array, axis): add dimension at location
- np.squeeze(array, axis): remove dims with size 1
 - axis =: if not specified, remove all.
- np.broadcast_to(array, shape): broadcast array to new shape
- np.broadcast_arrays(array1,array2,...): broad cast to make all arrays match. return list of arrays

Stack and merge

- numpy.concatenate(list of arrays,axis= 0): concatenate array1,array2,... along axis
- numpy.stack(list of arrays, axis = 0): join arrays along a new axis

Tiling arrays

- np.tile(array,reps): if reps = [x0,x1,...,x_n-1]repeat array in ith dimension for x_i times.
 - if array.ndims < n: prepend new axis to array to match n. So a shape (3,) array is promoted to (1, 3) for 2-D replication, or shape (1, 1, 3) for 3-D replication.
 - if array.ndims > n: prepend reps with 1 to match array.ndims.for an A of shape (2, 3, 4, 5), a reps of (2, 2) is treated as (1, 1, 2, 2).
- np.repeat(array, repeats, axis=None): repeat array along a single axis.
 - if axis = None, use array.flatten() as input, return 1d array
 - repeats =: int, number of repetition

Add and remove elements

- np.unique(array,...): find unique elements. merge df.value_counts(), df.nunique and other functions
 - return_index =: bool, return indices of input array that give unique value

- return_inverse =: bool, return indices of unique array to reconstruct original array. unique_array[inverse_index] = original_array.
- return_counts: counts of frequency for each unique value
- axis: axis to operate on, if =None, flatten array first. if specific axis, along that axis, flatten all subarrays.
- return tuple of (unique_array, indice, indice_inverse, counts)
- np.trim_zeros(1darray,trim = {'f','b','fb'}): trim front and back zeros of given 1 d array.
- np.delete(array, obj, axis =): return a new array where along new axis subarrays indexed by list of integers obj are removed.
- np.insert(array, obj, axis =): opposite of np.delete

2. random module

Random module is used to generate random samples from various distributions

2.1 random sampling from uniform/normal

- to generate random sample use np.random module
- random.rand(d0,d1,...,dn): random sampling from uniform distribution of [0,1] with shape [d0,d1,...,dn]
- random.randn(d0,d1,..,dn): random sampling from standard normal
- random.randint(low =, high =, size =): random sampling from uniform integer distribution. Right end is excluded. [low,high)
 - low: int, if high = None, this is the maximal integer. draw from [1,low], otherwise from [low, high]
 - high: int, the maximum
 - size: list/tuple, shape of np.array
- random.random_integers(low= ,high= ,size=): (deprecated) similar to randint, but both ends are inclusive. [low,high]

- random.random_sample(size): uniform from half-open interval [0,1), similar to rand but accepts tuple. Equivalent to random.random(size), random.ranf(size), random.sample(size).
- random.choice(a= , size= , replace =, p =): select from 1d array a, with given probability p, allow replacement if replace = True
 - a: 1d array-like object
 - p: 1d array-like object, if not given, assume uniform

2.2 permutation

• random.shuffle(x): shuffle array-like x along first axis. The modification is inplace

• random.permutation(x):similar to random.shuffle(), but make a copy and return result

2.3 Seed control

There are two ways to work with set seed.

- Set the global seed. Use random.seed(int).python np.random.seed(0)np.rand()np.random.seed(0)np.rand()# should be the same result
- 2. Create a container random.RandomState(seed=int) to fix seed for this particular container. Use member methods to generate random sample. For example RandomState(1000).randint(0,1,[10]), all functions covered in section 2.2 are available in the object's member methods. python container_A = random.RandomState(10)container_A.randn(10)container_B = random.RandomState(100)container_B.randn(100). Useful for managing multiple seeds.

2.4 Other distributions

There are other distributions like logistic(), beta(), binominal() and poisson(). The syntaxs are similar. Consists of (distribution_para, size)

I list some here:

- standard_normal(size)
- standard_t(size)
- f(dfnum =, dfden =, size): dfnum is for numerator. dfden is for denominator
- lognormal(mean, sigma, size):
- uniform([low,high,size]): half-open interval [low,high), float

3 Advanced mechanism

3.1 memory layout

• An instance of class ndarray consists of a contiguous one-dimensional segment of computer memory (owned by the array, or by some other object), combined with an indexing scheme that maps N integers into the location of an item in the block.

3.2 view and copy

• simple selection/slicing returns view of data. indexing based on index array /boolean array returns copy of data. See example below

```
1 Z = np.zeros(9)
2 Z_view = Z[0:3] # simple selection
3 Z_view[...] = 1
4 print(Z) # [1. 1. 1. 0. 0. 0. 0. 0.]
5
6 Z = np.zeros(9)
7 Z_copy = Z[[0,1,2]] # advanced index
8 Z_copy[...] = 1
9 print(Z) # [0. 0. 0. 0. 0. 0. 0. 0.]
```

• to check if the result is a view or copy use . base to return the orginal data and is to check

```
1 Z = np.random.uniform(0,1,(5,5))
2 Z1 = Z[:3,:]
3 Z2 = Z[[0,1,2], :]
```

```
4 Z1.base is Z # True
5 Z2.base is Z # False
6 Z2.base is None # True
```

- See function's document for whether a view or copy is returned
 - np.reshape() return new view whenever possible
 - np.flatten() returns new copy
- In arithmetic calculations between arrays, temporary copies are implicitly made. If arrays are big, think about other possible ways. For example, use np.add/minus/...(,out=) the keyword out= specifies the output numpy array, if the intermediate result need not to be saved, we can speed up slightly via coding the intermediate result with out on the same ndarray.