GIRIRAJ KRISHNA SHARMA 20051973

```
[2]: import numpy as np
     print(np.__version__)
    1.24.2
[3]: # Question : Create a 1D array of numbers from 0 to 9
     # Output : #> array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
     # Solution
     X = np.arange(10)
     Х
[3]: array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
[4]: # Question : Create a 3×3 numpy array of all True's
     # Solution
     np.full((3,3), True, dtype=bool)
     np.full((9), True, dtype=bool).reshape(3,3)
    np.ones((3,3), dtype=bool)
    np.ones((9), dtype=bool).reshape(3,3)
[4]: array([[ True, True, True],
            [ True, True, True],
            [ True, True, True]])
[5]: # Question : Extract all odd numbers from array
     # input: arr = np.array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
     # output: array([1, 3, 5, 7, 9])
     #Solution
```

```
arr = np.arange(10)
     arr[arr%2 == 1]
[5]: array([1, 3, 5, 7, 9])
[6]: # Question: Replace all odd numbers in arr with -1
     # input: arr = np.array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
     # output: array([ 0, -1, 2, -1, 4, -1, 6, -1, 8, -1])
     # Solution
     arr = np.arange(10)
     arr[arr\%2 == 1] = -1
     arr
[6]: array([0, -1, 2, -1, 4, -1, 6, -1, 8, -1])
[7]: # Question: Replace all odd numbers in arr with -1 without changing arr
     # input: arr = np.array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
     # output: out
     # array([ 0, -1, 2, -1, 4, -1, 6, -1, 8, -1])
     # arr
     # array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
     # Solution
     arr = np.arange(10)
     out = arr.copy()
     out[out\%2 == 1] = -1
     print('Modified Array')
     out
     print('\nOriginal Array')
     arr
    Modified Array
    Original Array
[7]: array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
```

```
[8]: # Question: Convert a 1D array to a 2D array with 2 rows
      # input: np.arange(10)
      # output array([[0, 1, 2, 3, 4],
                      [5, 6, 7, 8, 9]])
      # Solution
      arr = np.arange(10)
      arr.reshape(2,5)
      # Another solution
      arr = np.arange(10)
      arr.reshape(2, -1) # Setting to -1 automatically decides the number of cols
 [8]: array([[0, 1, 2, 3, 4],
             [5, 6, 7, 8, 9]])
 [9]: # Question: Stack arrays a and b vertically
      # input: a = np.arange(10).reshape(2,-1)
              b = np.repeat(1, 10).reshape(2, -1)
      # output: array([[0, 1, 2, 3, 4],
                      [5, 6, 7, 8, 9],
      #
                       [1, 1, 1, 1, 1],
      #
                      [1, 1, 1, 1, 1]])
      # Solution
      a = np.arange(10).reshape(2,-1)
      b = np.repeat(1, 10).reshape(2,-1)
      np.vstack([a,b])
 [9]: array([[0, 1, 2, 3, 4],
             [5, 6, 7, 8, 9],
             [1, 1, 1, 1, 1],
             [1, 1, 1, 1, 1]])
[10]: # Question: Stack the arrays a and b horizontally.
      # Input: a = np.arange(10).reshape(2,-1)
               b = np.repeat(1, 10).reshape(2, -1)
      # Output: array([[0, 1, 2, 3, 4, 1, 1, 1, 1],
                      [5, 6, 7, 8, 9, 1, 1, 1, 1, 1]])
      # Solution:
```

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a = np.arange(10).reshape(2,-1)
      b = np.repeat(1, 10).reshape(2, -1)
      np.hstack([a,b])
[10]: array([[0, 1, 2, 3, 4, 1, 1, 1, 1, 1],
             [5, 6, 7, 8, 9, 1, 1, 1, 1, 1]])
[11]: # Question: Create the following pattern without hardcoding. Use only numpy.
       ⇔functions and the below input array a.
      # Input: a = np.array([1,2,3])
      # Output: array([1, 1, 1, 2, 2, 2, 3, 3, 3, 1, 2, 3, 1, 2, 3, 1, 2, 3])
      # Solution
      a = np.array([1,2,3])
      np.r_[np.repeat(a, 3), np.tile(a, 3)]
      #other solution
      np.hstack((np.repeat(a, 3), np.tile(a, 3)))
[11]: array([1, 1, 1, 2, 2, 2, 3, 3, 3, 1, 2, 3, 1, 2, 3, 1, 2, 3])
[12]: # Question: Get the common items between a and b
      # Input: a = np.array([1,2,3,2,3,4,3,4,5,6])
               b = np.array([7,2,10,2,7,4,9,4,9,8])
      # Output: array([2, 4])
      # Solution:
      a = np.array([1,2,3,2,3,4,3,4,5,6])
      b = np.array([7,2,10,2,7,4,9,4,9,8])
      np.intersect1d(a,b)
[12]: array([2, 4])
[13]: # Question: From array a remove all items present in array b
      # Input: a = np.array([1,2,3,4,5])
              b = np.array([5,6,7,8,9])
      # Output: array([1,2,3,4])
```

```
# Solution
      a = np.array([1,2,3,4,5])
      b = np.array([5,6,7,8,9])
      np.setdiff1d(a,b)
[13]: array([1, 2, 3, 4])
[14]: # Question: Get the positions where elements of a and b match
      # Input: a = np.array([1,2,3,2,3,4,3,4,5,6])
              b = np.array([7,2,10,2,7,4,9,4,9,8])
      # Output: (array([1, 3, 5, 7]),)
      # Solution
      a = np.array([1,2,3,2,3,4,3,4,5,6])
      b = np.array([7,2,10,2,7,4,9,4,9,8])
      np.where(a == b)
[14]: (array([1, 3, 5, 7], dtype=int64),)
[15]: # Question: Get all items between 5 and 10 from a.
      # Input: a = np.array([2, 6, 1, 9, 10, 3, 27])
      # Output: (array([6, 9, 10]),)
      # Solution
      a = np.array([2, 6, 1, 9, 10, 3, 27])
      a[(a >= 5) & (a <= 10)]
[15]: array([6, 9, 10])
[16]: # Question: Convert the function maxx that works on two scalars, to work on two
       ⇔arrays.
      # Input:
      def \max(x, y):
          if x >= y:
              return x
          else:
```

```
return y
# maxx(1, 5)
#> 5
# Output:
\# a = np.array([5, 7, 9, 8, 6, 4, 5])
\# b = np.array([6, 3, 4, 8, 9, 7, 1])
# pair_max(a, b)
# array([ 6., 7., 9., 8., 9., 7., 5.])
# Solution
def pair_max(x, y):
    # here I am using map to make tuple from a and b, other solution is using \Box
\hookrightarrow zip(a,b)
    maximum = [maxx(a,b) for a,b in map(lambda a,b:(a,b),x,y)]
    # using zip
   # maximum = [maxx(a,b) for a,b in zip(x,y)]
    return np.array(maximum)
a = np.array([5, 7, 9, 8, 6, 4, 5])
b = np.array([6, 3, 4, 8, 9, 7, 1])
pair_max(a,b)
```

[16]: array([6, 7, 9, 8, 9, 7, 5])

```
[17]: # Question: Swap columns 1 and 2 in the array arr.

# Input:
arr = np.arange(9).reshape(3,3)

print('Original array')
arr

# Solution

print("\nModified array")
arr[:, [1,0,2]]
```

Original array

Modified array

```
[17]: array([[1, 0, 2],
             [4, 3, 5],
             [7, 6, 8]])
[18]: # Question: Swap rows 1 and 2 in the array arr:
      # Input:
      arr = np.arange(9).reshape(3,3)
      print('Original array')
      arr
      # Solution
      print("\nModified array")
      arr[[1,0,2], :]
     Original array
     Modified array
[18]: array([[3, 4, 5],
             [0, 1, 2],
             [6, 7, 8]])
[19]: # Question: Reverse the rows of a 2D array arr.
      # Input:
      arr = np.arange(9).reshape(3,3)
      print('Original array')
      arr
      # Solution
      print("\nModified array")
      arr[::-1, :]
     Original array
     Modified array
[19]: array([[6, 7, 8],
             [3, 4, 5],
             [0, 1, 2]])
```

```
[20]: # Question: Reverse the columns of a 2D array arr.
      # Input: arr = np.arange(9).reshape(3,3)
      # Solution
      arr = np.arange(9).reshape(3,3)
      print('Original array')
      arr
      print("\nModified array")
      arr[:, ::-1]
     Original array
     Modified array
[20]: array([[2, 1, 0],
             [5, 4, 3],
             [8, 7, 6]])
[21]: # Question: Create a 2D array of shape 5x3 to contain random decimal numbers.
       \hookrightarrow between 5 and 10.
      # Solution:
      rand_arr = np.random.uniform(5,10, size=(5,3))
      rand_arr
[21]: array([[6.87002046, 9.61643417, 6.06037858],
             [6.68709556, 9.55509519, 8.84574836],
             [7.77841326, 7.0560554, 5.36966421],
             [5.13337423, 9.14736216, 9.00253587],
             [5.4988653, 5.93242979, 8.27820228]])
[22]: # Pretty print rand_arr by suppressing the scientific notation (like 1e10)
      # Input:
      # Create the random array
      np.random.seed(100)
      rand_arr = np.random.random([3,3])/1e3
      np.set_printoptions(suppress=False)
      rand_arr
      # Output:
      #> array([[ 0.000543, 0.000278, 0.000425],
```

```
#>
                [ 0.000845, 0.000005, 0.000122],
                [ 0.000671, 0.000826, 0.000137]])
      #>
     np.set_printoptions(suppress=True)
     rand_arr
     #> array([[ 0.000543, 0.000278, 0.000425],
     #>
                [ 0.000845, 0.000005, 0.000122],
                [ 0.000671, 0.000826, 0.000137]])
      #>
[22]: array([[0.0005434, 0.00027837, 0.00042452],
             [0.00084478, 0.00000472, 0.00012157],
             [0.00067075, 0.00082585, 0.00013671]])
[23]: # Question: Limit the number of items printed in python numpy array a to a L
      ⇔maximum of 6 elements.
     a = np.arange(15)
     np.set_printoptions(threshold=6)
     a
[23]: array([0, 1, 2, ..., 12, 13, 14])
[24]: # Question: Print the full numpy array a without truncating.
      # Input: np.set printoptions(threshold=6)
      \# a = np.arange(15)
      # a
      # Output: a
     #> array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14])
     # Solution
     a = np.arange(15)
     np.set_printoptions(threshold=15)
[24]: array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14])
[25]: # Question: Import the iris dataset keeping the text intact.
      # Solution:
     iris_data = np.genfromtxt('Iris.csv', delimiter=',', skip_header=1,
                               usecols = [0,1,2,3,4,5], dtype = object)
     iris_data
```

```
[25]: array([[b'1', b'5.1', b'3.5', b'1.4', b'0.2', b'Iris-setosa'],
             [b'2', b'4.9', b'3.0', b'1.4', b'0.2', b'Iris-setosa'],
             [b'3', b'4.7', b'3.2', b'1.3', b'0.2', b'Iris-setosa'],
             [b'148', b'6.5', b'3.0', b'5.2', b'2.0', b'Iris-virginica'],
             [b'149', b'6.2', b'3.4', b'5.4', b'2.3', b'Iris-virginica'],
             [b'150', b'5.9', b'3.0', b'5.1', b'1.8', b'Iris-virginica']],
           dtype=object)
[26]: # Question-26: Extract the text column species from the 1D iris imported in
      ⇔previous question.
      data = np.genfromtxt('Iris.csv', delimiter=',', skip_header=1,
                                usecols = [-1], dtype = object)
      data
[26]: array([b'Iris-setosa', b'Iris-setosa', b'Iris-setosa', ...,
            b'Iris-virginica', b'Iris-virginica', b'Iris-virginica'],
            dtype=object)
[27]: # Question: Convert the 1D iris to 2D array iris 2d by omitting the species.
      \rightarrow text field.
      iris_data = np.genfromtxt('Iris.csv', delimiter=',', skip_header=1,__
      ⇔dtype='float', usecols=[0,1,2,3])
      iris_data
[27]: array([[ 1., 5.1, 3.5, 1.4],
             [ 2., 4.9,
                             3., 1.4],
             [ 3., 4.7,
                            3.2,
                                   1.3],
             [148., 6.5,
                             3.,
                                   5.2],
             [149., 6.2,
                             3.4,
                                   5.4],
                     5.9, 3., 5.1]])
             [150.,
[28]: # Question: Find the mean, median, standard deviation of iris's sepallength.
      \hookrightarrow (1st column)
      iris data = np.genfromtxt('Iris.csv', delimiter=',', skip header=1, usecols = 1
      [1])
      print('Mean', np.mean(iris_data))
      print('Median', np.median(iris_data))
      print('Standard Deviation', np.std(iris_data))
     Mean 5.843333333333334
     Median 5.8
```

Standard Deviation 0.8253012917851409

```
[29]: # Question: Create a normalized form of iris's sepallength whose values range
                  exactly between 0 and 1 so that the minimum has value 0 and maximum_
      ⇔has value 1.
      # Solution
      iris_data = np.genfromtxt('Iris.csv', delimiter=',', dtype='float',_
      ⇔usecols=[1], skip_header=1)
      (iris_data - np.min(iris_data))/(np.max(iris_data) - np.min(iris_data))
[29]: array([0.22222222, 0.16666667, 0.111111111, ..., 0.611111111, 0.52777778,
             0.4444444])
[30]: # Question: Compute the softmax score of sepallength.
      iris_data = np.genfromtxt('Iris.csv', delimiter=',', dtype='float',_
      →usecols=[1], skip_header=1)
      softmax = np.exp(iris_data)/sum(np.exp(iris_data))
      softmax.sum() # it must sum 1
      softmax
[30]: array([0.00221959, 0.00181724, 0.00148783, ..., 0.00900086, 0.006668 ,
             0.004939781)
[31]: | # Question. Find the 5th and 95th percentile of iris's sepallength
      iris_data = np.genfromtxt('Iris.csv', delimiter=',', dtype='float',__

susecols=[1], skip_header=1)

      np.percentile(iris_data, q=[5, 95])
[31]: array([4.6 , 7.255])
[32]: # Question: Insert np.nan values at 20 random positions in iris 2d dataset
      # Solution
      iris_data = np.genfromtxt('Iris.csv', delimiter=',', dtype='float',u

susecols=[1,2,3,4], skip_header=1)

      for i in np.random.randint(0, len(iris_data), 20):
          iris data[i]=np.nan
      iris_data
[32]: array([[5.1, 3.5, 1.4, 0.2],
             [4.9, 3., 1.4, 0.2],
             [4.7, 3.2, 1.3, 0.2],
             [6.5, 3., 5.2, 2.],
```

```
[33]: # Question: Find the number and position of missing values in iris 2d's
       ⇔sepallength (1st column)
      iris_data = np.genfromtxt('Iris.csv', delimiter=',', dtype='float',_
       \subsetequsecols=[1,2,3,4], skip_header=1)
      iris_data[np.random.randint(len(iris_data), size=20),np.random.
       ⇒randint(4,size=20)] = np.nan
      # Find total mising value in complete data
      print("Number of missing values in Iris data: \n", np.isnan(iris_data[:, :]).
       ⇒sum())
      # Find total mising value in 1D data
      print("Number of missing values in any one feature of Iris data: \n", np.
       sisnan(iris_data[:, 0]).sum())
      print("Position of missing values: \n", np.where(np.isnan(iris_data[:, 0])))
     Number of missing values in Iris data:
      20
     Number of missing values in any one feature of Iris data:
     Position of missing values:
      (array([ 38, 80, 106, 113, 121], dtype=int64),)
[34]: # Question: Filter the rows of iris 2d that has petallength (3rd column) > 1.5_{\sqcup}
       \hookrightarrow and sepallength (1st column) < 5.0
      iris_data = np.genfromtxt('Iris.csv', delimiter=',', dtype='float',u
       \subsetequsecols=[1,2,3,4], skip_header=1)
      # Solution
      iris_data[(iris_data[:, 2] > 1.5) & (iris_data[:, 0] < 5.0)]</pre>
[34]: array([[4.8, 3.4, 1.6, 0.2],
             [4.8, 3.4, 1.9, 0.2],
             [4.7, 3.2, 1.6, 0.2],
             [4.8, 3.1, 1.6, 0.2],
             [4.9, 2.4, 3.3, 1.],
             [4.9, 2.5, 4.5, 1.7]
[35]: # Question: Select the rows of iris_2d that does not have any nan value.
      diabetes_data = np.genfromtxt('diabetes.csv', delimiter=',',
```

[6.2, 3.4, 5.4, 2.3], [5.9, 3., 5.1, 1.8]])

```
dtype='float', usecols=[0,1,2,3,4,5,6,7],__
       ⇒skip_header=1)
      diabetes_data[np.random.randint(150, size=20), np.random.randint(4, size=20)] = __
      diabetes_data[np.sum(np.isnan(diabetes_data), axis = 1) == 0][:5]
[35]: array([[ 1.
                                 66.
                     , 85.
                                        , ...,
                                             26.6 ,
                                                       0.351, 31.
             [ 8.
                                             23.3 ,
                                                       0.672, 32.
                                                                      ],
                     , 183.
                                 64.
                                        , ...,
             [ 1.
                     , 89.
                                 66.
                                             28.1 ,
                                                        0.167, 21.
                                                                      ],
                                                                      ],
             [ 0.
                     , 137.
                                 40.
                                             43.1 ,
                                                        2.288,
                                                                33.
                                                                      ]])
             [ 3.
                                                        0.248, 26.
                      78.
                                 50.
                                             31.
                                        , ...,
[36]: # question: Find the correlation between SepalLength(1st column) and
      →PetalLength(3rd column) in iris_2d
      # insted or using iris data I am going to used pima diabetes data and going to \sqcup
       ⇔find corelation
      # between BP(1st column) and BMI (5th column).
      diabetes_data = np.genfromtxt('diabetes.csv',
                                    delimiter=',', dtype='float',u
       \cupusecols=[0,1,2,3,4,5,6,7], skip_header=1)
      print(np.corrcoef(diabetes_data[:, 1], diabetes_data[:, 5]))
      print('\n')
      # you can get correlation by getting value at index [0,1] or [1,0]
      print(np.corrcoef(diabetes_data[:, 1], diabetes_data[:, 5])[0,1])
                  0.22107107]
     [[1.
      [0.22107107 1.
                            ]]
     0.2210710694589828
[37]: # Question: What is the value of second longest petallength of species setosa
      # For this question I am going to find second highest bloodpressure (2nd_{
m L}
       ⇔column) where outcome is 1
      diabetes_data = np.genfromtxt('diabetes.csv',
                                    delimiter=',', dtype=object,⊔
       \cupusecols=[0,1,2,3,4,5,6,7,8], skip_header=1)
      # Solution
      bloodpressure= diabetes_data[diabetes_data[:, 8]==b'1', [2]].astype('float')
      np.unique(np.sort(bloodpressure))[-2]
```

[37]: 110.0

```
[38]: # Question: Sort the iris dataset based on sepallength column.
      # In this problem, I am going to sort the diabetes dataset based on Glucose_{\sqcup}
       \hookrightarrow (1th column)
      diabetes data = np.genfromtxt('diabetes.csv',
                                      delimiter=',', dtype=object,⊔
       \cupusecols=[0,1,2,3,4,5,6,7,8], skip_header=1)
      diabetes_data[diabetes_data[:,1].argsort()]
[38]: array([[b'1', b'0', b'74', ..., b'0.299', b'21', b'0'],
              [b'5', b'0', b'80', ..., b'0.346', b'37', b'1'],
             [b'1', b'0', b'68', ..., b'0.389', b'22', b'0'],
             [b'4', b'99', b'72', ..., b'0.294', b'28', b'0'],
             [b'4', b'99', b'76', ..., b'0.223', b'21', b'0'],
             [b'2', b'99', b'60', ..., b'0.453', b'21', b'0']], dtype=object)
[39]: # Question: Find the most frequent value of petal length (3rd column) in irisu
       \hookrightarrow dataset.
      iris_data = np.genfromtxt('Iris.csv', delimiter=',',
                                 dtype=object, usecols=[1,2,3,4,5], skip_header=1)
      v,c = np.unique(iris_data[:, 2], return_counts=True)
      v[np.argmax(c)]
[39]: b'1.5'
[40]: # Question: Find the position of the first occurrence of a value greater than 1.
      # petalwidth 4th column of iris dataset.
      iris_data = np.genfromtxt('Iris.csv', delimiter=',',
                                 dtype=object, usecols=[4], skip_header=1)
      np.argwhere(iris_data[:].astype(float) > 1.0)[0]
[40]: array([50], dtype=int64)
[41]: # Question: From the array a, replace all values greater than 30 to 30 and less
       \hookrightarrow than 10 to 10.
      # Solution
      np.set_printoptions(precision=2)
      np.random.seed(100)
      a = np.random.uniform(1,50, 20)
```

```
a[a<10]=10
      a[a>30]=30
      np.set_printoptions(threshold=20)
[41]: array([27.63, 14.64, 21.8 , 30. , 10. , 10. , 30. , 30. , 10. ,
             29.18, 30. , 11.25, 10.08, 10. , 11.77, 30. , 30. , 10. ,
             30. , 14.43])
[42]: # Question: Get the positions of top 5 maximum values in a given array a.
      np.random.seed(100)
      a = np.random.uniform(1,50, 20)
      sort = a.argsort()
      print('Positions')
      sort[-5:][::-1]
      print('Values')
      a[sort][-5:][::-1]
     Positions
     Values
[42]: array([48.95, 44.67, 42.39, 41.47, 41. ])
[43]: # Question: Compute the counts of unique values row-wise.
      # Solution
      def counts_of_all_values_rowwise(arr2d):
          # Unique values and its counts row wise
          num_counts array = [np.unique(row, return_counts=True) for row in arr2d]
          # Counts of all values row wise
          return([[int(b[a==i]) if i in a else 0 for i in np.unique(arr2d)] for a, b_{\sqcup}
       →in num_counts_array])
      np.random.seed(100)
      np.set_printoptions(threshold=10)
      arr = np.random.randint(1,11,size=(6, 10))
      arr
      print(np.arange(1,11))
      counts_of_all_values_rowwise(arr)
     [1 2 3 4 5 6 7 8 9 10]
[43]: [[1, 0, 2, 1, 1, 1, 0, 2, 2, 0],
       [2, 1, 3, 0, 1, 0, 1, 0, 1, 1],
       [0, 3, 0, 2, 3, 1, 0, 1, 0, 0],
```

```
[1, 0, 2, 1, 0, 1, 0, 2, 1, 2],
       [2, 2, 2, 0, 0, 1, 1, 1, 1, 0],
       [1, 1, 1, 1, 1, 2, 0, 0, 2, 1]]
[44]: # Question: Convert array of arrays into a flat linear 1d array.
      arr1 = np.arange(3)
      arr2 = np.arange(3,7)
      arr3 = np.arange(7,10)
      arr_2d = np.concatenate([arr1, arr2, arr3])
      print(arr_2d)
     [0 1 2 3 4 5 6 7 8 9]
[45]: # Q-51
      #1 How to get the n largest values of an array ?
      Z = np.arange(10000)
      np.random.shuffle(Z)
      n = 5
      # Slow
      print (Z[np.argsort(Z)[-n:]])
      # Fast
      print (Z[np.argpartition(-Z,n)[:n]])
     [9995 9996 9997 9998 9999]
     [9999 9998 9996 9997 9995]
[46]: #2 Given an arbitrary number of vectors, build the cartesian product
      # (every combinations of every item).
      def cartesian(arrays):
          arrays = [np.asarray(a) for a in arrays]
          shape = (len(x) for x in arrays)
          ix = np.indices(shape, dtype=int)
          ix = ix.reshape(len(arrays), -1).T
          for n, arr in enumerate(arrays):
              ix[:, n] = arrays[n][ix[:, n]]
          return ix
      print (cartesian(([1, 2, 3], [4, 5], [6, 7])))
     [[1 4 6]
```

[1 4 7]

```
[1 5 6]
      [3 4 7]
      [3 5 6]
      [3 5 7]]
[47]: #3 Consider a 16x16 array, how to get the blocksum (block size is 4x4)?
      Z = np.ones((16,16))
      k = 4
      S = np.add.reduceat(np.add.reduceat(Z, np.arange(0, Z.shape[0], k), axis=0), np.
       \Rightarrowarange(0, Z.shape[1], k), axis=1)
      print ('input array')
      print (Z)
      print ('block sum')
      print (S)
     input array
     [[1. 1. 1. ... 1. 1. 1.]
      [1. 1. 1. ... 1. 1. 1.]
      [1. 1. 1. ... 1. 1. 1.]
      [1. 1. 1. ... 1. 1. 1.]
      [1. 1. 1. ... 1. 1. 1.]
      [1. 1. 1. ... 1. 1. 1.]]
     block sum
     [[16. 16. 16. 16.]
      [16. 16. 16. 16.]
      [16. 16. 16. 16.]
      [16. 16. 16. 16.]]
[48]: #4 Compute a matrix rank.
      Z = np.random.uniform(0,1,(10,10))
      U, S, V = np.linalg.svd(Z)
                                    # Singular Value Decomposition
      rank = np.sum(S > 1e-10)
      print (rank)
     10
[49]: #5 How to find the most frequent value in an array?
      Z = np.random.randint(0,10,50)
      print (Z)
      print('rank:', np.bincount(Z).argmax())
     [3 8 7 ... 9 4 5]
     rank: 3
[50]: #6 Extract all the contiguous 3x3 blocks from a random 10x10 matrix.
      Z = np.random.randint(0,5,(6,6))
```

```
n = 3
   i = 1 + (Z.shape[0]-3)
   j = 1 + (Z.shape[1]-3)
   C = np.lib.stride_tricks.as_strided(Z, shape=(i, j, n, n), strides=Z.strides + L tricks.as_strides + L trick

∠Z.strides)
   print(C)
[[[[2 3 1]
                   [0 3 3]
                   [4 0 2]]
            [[3 1 2]
                   [3 3 0]
                   [0 2 2]]
            [[1 2 1]
                   [3 0 0]
                   [2 2 1]]
            [[2 1 2]
                   [0 0 1]
                   [2 1 3]]]
     [[[0 3 3]
                   [4 0 2]
                   [0 1 0]]
            [[3 3 0]
                   [0 2 2]
                   [1 0 1]]
            [[3 0 0]
                   [2 2 1]
                   [0 1 1]]
            [[0 0 1]
                   [2 1 3]
                   [1 1 4]]]
     [[[4 0 2]
                   [0 1 0]
                   [1 2 0]]
            [[0 2 2]
```

[1 0 1]

- [2 0 4]]
- [[2 2 1]
- [0 1 1]
- [0 4 4]]
- [[2 1 3]
- [1 1 4]
- [4 4 2]]]
- [[[0 1 0]
 - [1 2 0]
 - [0 1 1]]
- [[1 0 1]
- [2 0 4]
- [1 1 4]]
- [[0 1 1]
- [0 4 4]
- [1 4 0]]
- [[1 1 4]
- [4 4 2]
- [4 0 2]]]