# Homework-2

1. Householder Reflector 
$$F = I - 299*$$

Where  $9 = |hx||_2 e_1 - \epsilon$ .

$$y - \frac{2y9*}{y*y}y = \lambda y$$

$$(1-\lambda)y = \left(\frac{2y9*}{y*y}\right)y$$

$$(1-\lambda)y = 2\left(\frac{y*y}{y}\right)y$$

So one possible solution is  $\chi = 1$  and the corresponding eigenvector y is perpendicular to 9 (or the RHS, 9 \* y = 0)

If  $\lambda \neq 1$ , we can see that y is a scalar multiple of 9:

If  $y=x^y$ ,  $x\in C$ , then  $(1-\lambda)x=2x$  or  $\lambda=-1$ the corresponding set of eigenvectors are scalar multiples of  $\nu$ .

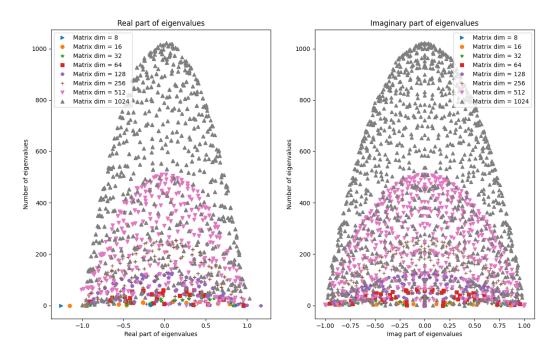
ⓑ We know 
$$F^2 = I$$
  
(I-29, of) (I-29,9\*)  
= I-49,9\*+49,9\*ay9\* = I  
So det (F) = ±1

Now, consider det  $(I + xy^*)$  can be written as  $det(I - y^*)$ . (where  $xy^* = I$ ) = I - xty (using the determinant expression for a 2x2 matrix). If 29 = Cy, then det(F) = det(I - 20/9\*) = det(I - 20/9) = -1 C F\*= (I-200\*) = I - 200 = F Since F is Hermitian, it has real eighnvalues and the eigendecomposition of F= D. A. D. yields a Orthonormal matrix Q(Q=Q-1) So, we can write  $F = Q | \Delta | sign(\Delta) Q,$ sign (A) is a diagonal matrix with signs of each of the eigenvalues in 1. Since Sign (1) &\* is unitary, we have ourselves a singular value decomposition for F. => The singular values of F = | ± 1 | = 1

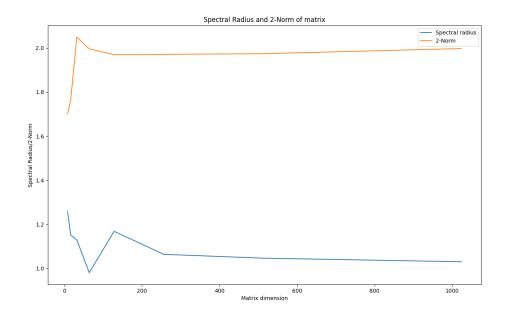
(Since the eigenvalues of F= ±1).

### Problem 2

A. Eigenvalues superimposed in a single plot for different values of m (matrix dimension). The real and imaginary parts of the eigenvalues lie between -1.0 and 1.0 for all matrix dimensions.

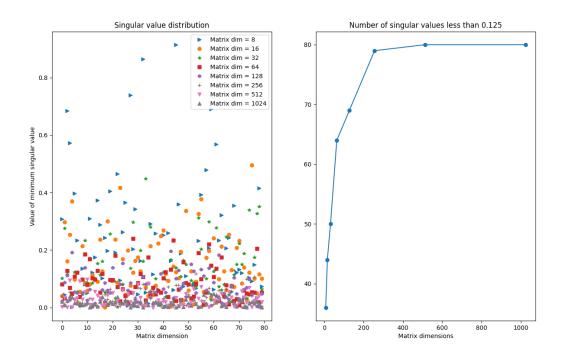


B. Spectral radius and 2-Norm of the matrix
 The spectral radius tends to 2 as m tends to infinity. The 2-Norm of the matrix tends to 1 as m tends to infinity.



### C. Smallest singular value of the matrix

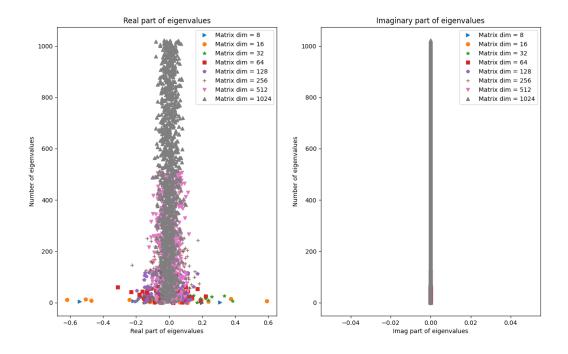
As the matrix dimension increases, the number of minimum singular values less than  $(\frac{1}{2})^n$  where  $n = \{-1, -2, ...\}$  increases. For example, m = 1024, the number of singular values less than  $\frac{1}{8}$  is higher than m = 16.



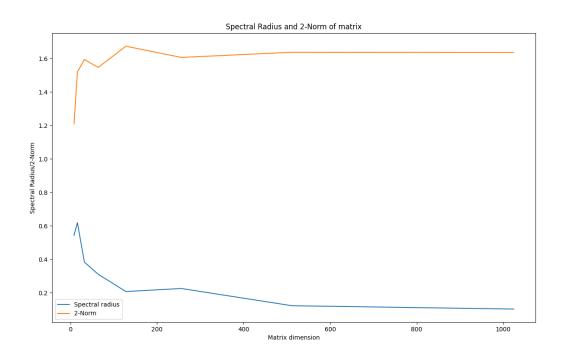
### D. Upper Triangular Matrix with normally distributed entries

### a. Eigenvalue distribution

We can see that the real part has a gaussian distribution and imaginary parts of the eigenvalues are 0 for all matrix dimensions.

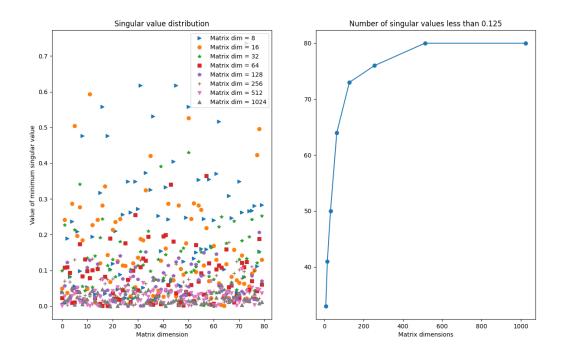


# b. Spectral Radius and 2-Norm plotThe spectral radius tends to 1.6 and the 2-norm tends to 0 as m tends to infinity.



### c. Smallest singular value of the matrix

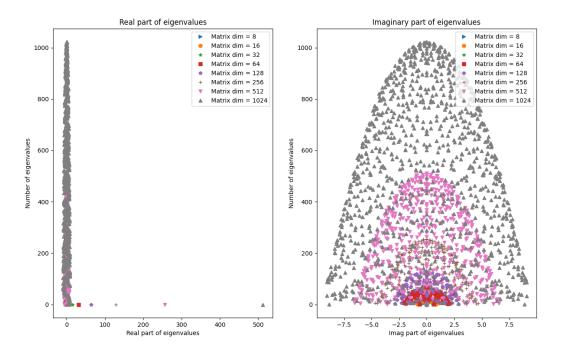
As the matrix dimension increases, the number of minimum singular values less than  $(\frac{1}{2})^n$  where  $n = \{-1, -2, ...\}$  increases. For example, m = 1024, the number of singular values less than  $\frac{1}{8}$  is higher than m = 16. However, the curve on the right is smoother than the curve for the above case.



### Problem 3

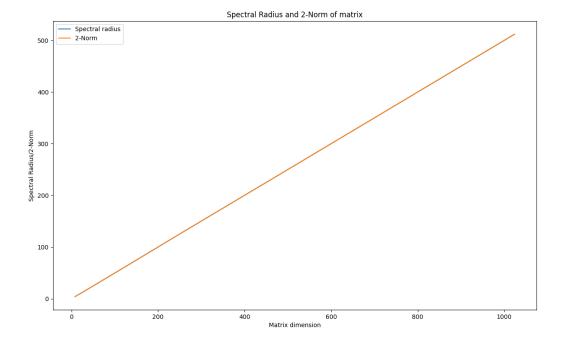
### A. Distribution of Eigenvalues

The real part of the eigenvalues lie close to 0 and imaginary parts of the eigenvalues lie between -8.0 and 8.0 for all matrix dimensions.

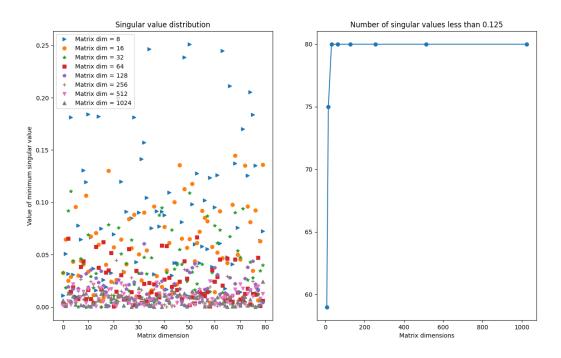


### B. Spectral Radius and 2-Norm

The spectral radius and 2-norm curves overlap each other for all matrix dimensions and increase linearly with matrix dimension.



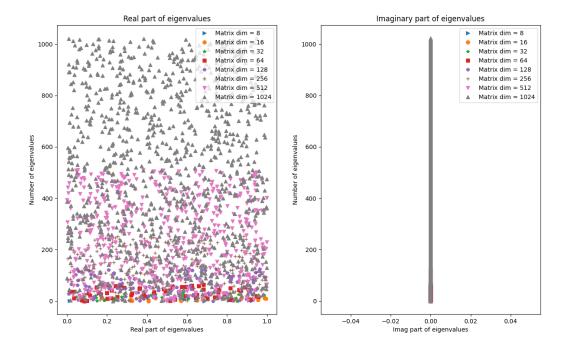
# C. Distribution of minimum singular values The minimum singular value distribution for matrices with uniformly distributed entries shows that higher numbers of singular values are close to 0 as m tends to infinity.



# D. Upper triangular uniform distribution

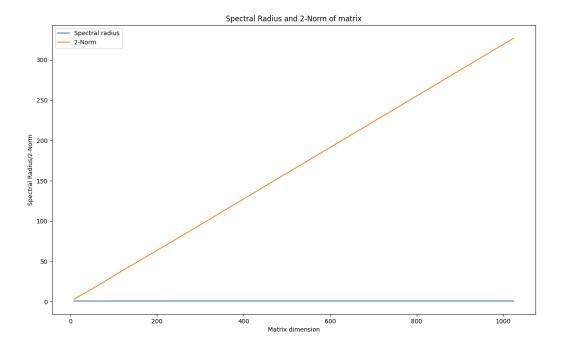
## a. Eigenvalue distribution

The real part of eigenvalues lie between 0 and 1 and the imaginary part is 0.



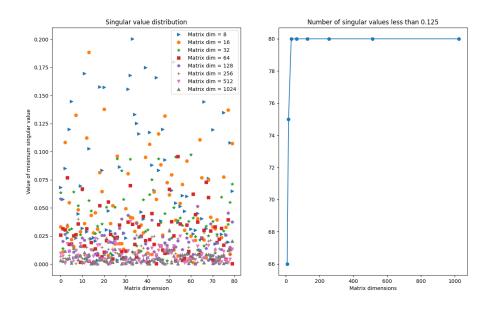
### b. Spectral radius and 2-norm

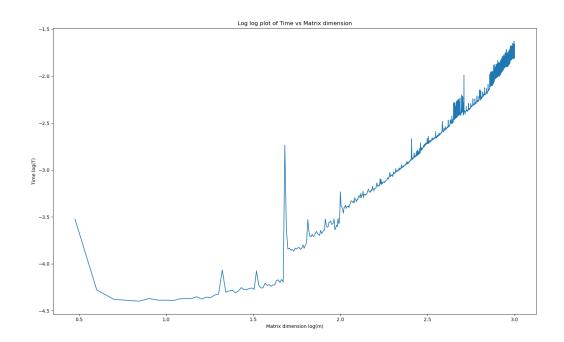
The 2-norm is similar to the curve in the case for the full matrix, but the spectral radius is 0.



## c. Minimum singular value distribution

The minimum singular value distribution for matrices with uniformly distributed entries shows that higher numbers of singular values are close to 0 as m tends to infinity.





The slope of this curve = 1.023844779280685 (end to end slope)
As the matrix dimension increases, the time taken first reduces and then starts increasing.

### b. M = 1000

Spectral Radius = 3.9999901501133026 Max Singular Value = 3.999990150113325 Min Singular Value = 9.849886676613993e-06 Condition Number = 406095.04265772586 Frobenius Norm = 77.44675590365293 2 Norm = 77.44675590365293

### g. Summarizing times for all methods

From the table, we can see that the SVD takes the longest time to decompose the matrix. We also see that LU and Cholesky have the least residual for b-Ax, and so are most accurate for this matrix.

Also, np.linalg.solve has the same residual as LU, which means that the numpy library uses LU decomposition underneath.

Type of decomposition	Time taken	b-Ax
QR	2.823699951171875 s	5.3652346348568966e-05
LU	0.61257004737854 s	2.720789863768137e-05

SVD	18.694941997528076 s	0.0007673158470886349
Cholesky	0.49833083152770996 s	3.899493143684322e-05
np.linalg.solve	0.44628095626831055 s	2.720789863768137e-05

### Code for Problem 2

```
from cProfile import label
import numpy as np
import matplotlib.pyplot as plt
from torch import norm, threshold
def generate random matrix(m):
  mat = np.random.randn(m, m)
   return mat
def make_upper_traingular(mat):
   return np.triu(mat)
def normalize_matrix(mat, m):
  mat = mat / np.sqrt(m)
   return mat
def get_eigen_values(mat):
  v, _ = np.linalg.eig(mat)
  return v
def get_spectral_radius(eig_vals):
   return np.max(np.absolute(eig_vals))
def get_norm_2(mat):
   return np.linalg.norm(mat, 2)
def get_smallest_sigma(mat):
   sg_min = np.min(np.linalg.svd(mat, compute_uv=False))
  return sg min
def plot_eig_vals(eig_values, m_sizes, markers):
  plt.clf()
   plt.cla()
   fig, (ax1, ax2) = plt.subplots(1, 2)
```

```
num_plots = len(m_sizes)
   for i in range(num plots):
       ei = eig values[i]
       ei real = np.real(ei)
       ei complex = np.imag(ei)
       m = m_sizes[i]
       y = np.arange(m)
       ax1.plot(ei real, y, markers[i], label='Matrix dim = '+str(m))
       ax2.plot(ei complex, y, markers[i], label='Matrix dim = '+str(m))
   ax1.legend()
   ax1.set_xlabel('Real part of eigenvalues')
   ax1.set ylabel('Number of eigenvalues')
   ax1.set title('Real part of eigenvalues')
  ax2.legend()
  ax2.set xlabel('Imag part of eigenvalues')
  ax2.set_ylabel('Number of eigenvalues')
   ax2.set title('Imaginary part of eigenvalues')
   # plt.legend()
   plt.show()
def plot_spectral_radius(sr, nm, m):
  plt.clf()
  plt.cla()
   plt.plot(m, sr, label='Spectral radius')
  plt.plot(m, nm, label='2-Norm')
  plt.title("Spectral Radius and 2-Norm of matrix")
  plt.xlabel("Matrix dimension")
   plt.ylabel("Spectral Radius/2-Norm")
   plt.legend()
   plt.show()
def plot min singular distribution(m list, threshold, markers):
  plt.clf()
   plt.cla()
  fig, (ax1, ax2) = plt.subplots(1, 2)
  sg_min_tail = []
   j = 0
   for m in m list:
       sg_min_list = []
      count = 0
       for i in range(80):
```

```
mat = generate_random_matrix(m)
           sg min = get smallest sigma(mat)
           sg min list.append(sg min)
           if (sg min < threshold):</pre>
               count += 1
       sg_min_tail.append(count)
       ax1.plot(sg min list, markers[j], label='Matrix dim = {}'.format(m))
   ax1.set title('Singular value distribution')
   ax1.legend()
   ax1.set_xlabel("Matrix dimension")
   ax1.set ylabel("Value of minimum singular value")
   ax2.plot(m list, sg min tail, 'o-')
   ax2.set title('Number of singular values less than {}'.format(threshold))
   ax2.set xlabel("Matrix dimensions")
   plt.show()
   return
def run_experiment(m_list):
  eig values = []
  spectral_radius = []
  norm 2 = []
   threshold = 1./np.power(2, 3)
   for m in m_list:
       mat = generate random matrix(m)
       mat = make upper traingular(mat)
       mat = normalize matrix(mat, m)
       ei = get eigen values(mat)
       eig_values.append(ei)
       sr = get spectral radius(ei)
       spectral radius.append(sr)
       n2 = get norm 2 (mat)
       norm 2.append(n2)
   markers = ['>', 'o', '*', 's', "p", "+", "v", "^", "x"]
   plot eig vals(eig values, m list, markers)
   plot spectral radius(spectral radius, norm 2, m list)
   # m copy = [m list[4]]
   plot min singular distribution(m list, threshold, markers)
if __name__ == '__main__':
   m \text{ list} = [8, 16, 32, 64, 128, 256, 512, 1024]
```

```
run_experiment(m_list)
```

### Code for Problem 3

```
from cProfile import label
import numpy as np
import matplotlib.pyplot as plt
from torch import norm, threshold
def generate_random_matrix(m):
  mat = np.random.rand(m, m)
   return mat
def make_upper_traingular(mat):
  return np.triu(mat)
def normalize matrix(mat, m):
  mat = mat / np.sqrt(m)
   return mat
def get_eigen_values(mat):
  v, _ = np.linalg.eig(mat)
  return v
def get_spectral_radius(eig_vals):
   return np.max(np.absolute(eig_vals))
def get_norm_2(mat):
  return np.linalg.norm(mat, 2)
def get smallest sigma(mat):
  sg_min = np.min(np.linalg.svd(mat, compute_uv=False))
  return sg_min
def plot eig vals(eig values, m sizes, markers):
  plt.clf()
   plt.cla()
  fig, (ax1, ax2) = plt.subplots(1, 2)
  num_plots = len(m_sizes)
   for i in range(num plots):
```

```
ei = eig_values[i]
       ei real = np.real(ei)
       ei complex = np.imag(ei)
       m = m \text{ sizes}[i]
       y = np.arange(m)
       ax1.plot(ei_real, y, markers[i], label='Matrix dim = '+str(m))
       ax2.plot(ei complex, y, markers[i], label='Matrix dim = '+str(m))
   ax1.legend()
  ax1.set xlabel('Real part of eigenvalues')
  ax1.set ylabel('Number of eigenvalues')
  ax1.set_title('Real part of eigenvalues')
  ax2.legend()
  ax2.set xlabel('Imag part of eigenvalues')
  ax2.set ylabel('Number of eigenvalues')
  ax2.set title('Imaginary part of eigenvalues')
   # plt.legend()
  plt.show()
def plot_spectral_radius(sr, nm, m):
  plt.clf()
  plt.cla()
  plt.plot(m, sr, label='Spectral radius')
  plt.plot(m, nm, label='2-Norm')
  plt.title("Spectral Radius and 2-Norm of matrix")
  plt.xlabel("Matrix dimension")
  plt.ylabel("Spectral Radius/2-Norm")
  plt.legend()
  plt.show()
def plot min singular distribution(m list, threshold, markers):
  plt.clf()
  plt.cla()
  fig, (ax1, ax2) = plt.subplots(1, 2)
  sg_min_tail = []
  j = 0
  for m in m list:
       sg min list = []
       count = 0
       for i in range(80):
           mat = generate_random_matrix(m)
           sg min = get smallest sigma(mat)
```

```
sg_min_list.append(sg_min)
           if (sg min < threshold):</pre>
               count += 1
       sg min tail.append(count)
       ax1.plot(sg min list, markers[j], label='Matrix dim = {}'.format(m))
       j += 1
  ax1.set title('Singular value distribution')
   ax1.legend()
  ax1.set xlabel("Matrix dimension")
  ax1.set ylabel("Value of minimum singular value")
  ax2.plot(m_list, sg_min_tail, 'o-')
  ax2.set title('Number of singular values less than {}'.format(threshold))
  ax2.set_xlabel("Matrix dimensions")
  plt.show()
  return
def run experiment(m list):
  eig values = []
  spectral radius = []
  norm 2 = []
  threshold = 1./np.power(2, 3)
  for m in m list:
       mat = generate random matrix(m)
       mat = make_upper_traingular(mat)
       # mat = normalize matrix(mat, m)
       ei = get_eigen_values(mat)
       eig values.append(ei)
       sr = get spectral radius(ei)
       spectral_radius.append(sr)
       n2 = get norm 2 (mat)
       norm 2.append(n2)
  markers = ['>', 'o', '*', 's', "p", "+", "v", "^", "x"]
  plot eig vals(eig values, m list, markers)
  plot_spectral_radius(spectral_radius, norm_2, m_list)
   # m copy = [m list[4]]
  plot min singular distribution(m list, threshold, markers)
if name == ' main ':
  m = [8, 16, 32, 64, 128, 256, 512, 1024]
  run experiment(m list)
```

### Code for Problem 4

```
import numpy as np
import matplotlib.pyplot as plt
import scipy as scp
from scipy import linalg as scplinalg
from torch import permute
import time
def construct 1D laplace(m):
  diagonal = -2. * np.ones(m)
  off diag = np.ones(m-1)
  return np.diag(diagonal, 0) + np.diag(off diag, 1) + np.diag(off diag, -1)
def construct rhs(m):
  b = np.arange(m)
  b = b.astype(np.float64)
  return b
def get spectral radius(mat):
  ei, _ = np.linalg.eig(mat)
  sr = np.abs(ei)
  sr = np.max(sr)
   return sr
def get singular value min max(mat):
   sg = np.linalg.svd(mat, compute_uv=False)
   return np.max(sg), np.min(sg)
def get frobenius norm(mat):
  return np.linalg.norm(mat, 'fro')
def get_2_norm(mat):
   return np.linalg.norm(mat, 2)
def compute condition number(mat):
  return np.linalg.cond(mat)
# Get Spectral radius, condition number, sigma max/min, matrix norm fro, 2
def get matrix characteristics(mat):
  sr = get_spectral_radius(mat)
```

```
sg_max, sg_min = get_singular_value_min_max(mat)
  norm_f = get_frobenius_norm(mat)
  norm_2 = get_frobenius_norm(mat)
   cnum = compute condition number(mat)
   return sr, sg max, sg min, cnum, norm f, norm 2
# QR Decomposition
def get QR decomposition(mat):
   q, r = np.linalg.qr(mat)
   return q, r
def check qr correctness(mat, q, r):
  diff = np.linalg.norm(mat - np.matmul(q, r))
   print("Diff in QR = {}".format(diff))
def solve_using_qr(q, r, b):
  start = time.time()
  y = np.matmul(np.transpose(q), b)
  end = time.time() - start
  x = scplinalg.solve triangular(r, y)
  return x, end
# LU Decomposition
def get_lu_decomposition(mat):
   p, l, u = scp.linalg.lu(mat, permute l=False)
   return p, 1, u
def check lu correctness(mat, p, 1, u):
   diff = np.linalg.norm(mat - p @ 1 @ u)
   print("Diff in LU = {}".format(diff))
def solve using lu(p, l, u, b):
  b1 = np.matmul(np.transpose(p), b)
  y = scplinalg.solve_triangular(1, b1, lower=True)
   x = scplinalg.solve triangular(u, y, lower=False)
   return x
# SVD Decomposition
def get_svd_decomposition(mat):
  u, sigma, vh = np.linalg.svd(mat)
  return u, sigma, vh
```

```
def check_svd_correctness(mat, u, sigma, vh):
   diff = np.linalg.norm(mat - u @ np.diag(sigma) @ vh)
   print("Diff in SVD = {}".format(diff))
def solve_using_svd(u, sigma, vh, b):
   c = np.dot(np.transpose(u),b)
   w = np.dot(np.diag(1./sigma),c)
   \# Vh x = w <=> x = Vh.H w (where .H stands for hermitian = conjugate transpose)
   x = np.dot(np.transpose(vh),w)
   return x
# Cholesky Decomposition
def get cholesky decomposition(mat):
  L = np.linalg.cholesky(mat)
   return L
def check cholesky correctness(mat, L):
   diff = np.linalg.norm(mat - np.matmul(L, np.transpose(L)))
   print("Diff in Cholesky = {}".format(diff))
def solve using cholesky(L, b):
  y = scplinalg.solve triangular(L, b, lower=True)
   x = scplinalg.solve_triangular(np.transpose(L), y, lower=False)
   return x
# Solve using numpy.linalg.solve
def solve using np solve(mat, b):
   x = np.linalg.solve(mat, b)
   return x
# Verify norm of |b-Ax|
def verify solution(mat, x, b):
   diff = np.linalg.norm(np.matmul(mat, x) - b)
   print("|b - Ax| = {}".format(diff))
def log plot qr time():
  times = []
  m list = []
  for m in range(3, 1001):
```

```
mat = construct_1D_laplace(m)
       b = construct rhs(m)
       time qr = time.time()
       q, r = get QR decomposition(mat)
       y = np.matmul(np.transpose(q), b)
       time_qr = time.time() - time_qr
       y = y + 1
       times.append(np.log10(time qr))
       m list.append(np.log10(m))
   print(np.array(times[1:]) - np.array(times[0:-1])/(np.array(m list[1:]) -
np.array(m_list[0:-1])))
  plt.clf()
  plt.cla()
  plt.plot(m list, times)
  plt.title("Log log plot of Time vs Matrix dimension")
  plt.xlabel("Matrix dimension log(m)")
   plt.ylabel("Time log(T)")
   plt.show()
def run experiment(m):
  mat = construct_1D_laplace(m)
   b = construct rhs(m)
   sr, sg_max, sg_min, cnum, norm_f, norm_2 =
get matrix characteristics(construct 1D laplace(1000))
   print("Spectral Radius = {}".format(sr))
   print("Max Singular Value = {}".format(sg max))
   print("Min Singular Value = {}".format(sg min))
   print("Condition Number = {}".format(cnum))
   print("Frobenius Norm = {}".format(norm f))
   print("2 Norm
                           = {}".format(norm 2))
   # Log time vs Log dimension
   log_plot_qr_time()
   # Solve using QR
   print("Solve using QR")
   time qr = time.time()
   q, r = get_QR_decomposition(mat)
   time qr = time.time() - time qr
   check qr correctness(mat, q, r)
```

```
x_qr, time_qr1 = solve_using_qr(q, r, b)
  time qr += time qr1
  print("Time taken for QR decomposition and y=Q*b=\{\} s".format(time qr))
  verify solution(mat, x qr, b)
   # Solve using LU
  print("Solve using LU")
  time lu = time.time()
  p, l, u = get lu decomposition(mat)
  time lu = time.time() - time lu
  check_lu_correctness(mat, p, 1, u)
  time lu solve = time.time()
  x lu = solve using lu(p, l, u, b)
  time lu += time.time() - time lu solve
  print("Time taken for LU decomposition and solve = {} s".format(time_lu))
  verify_solution(mat, x_lu, b)
   # Solve using SVD
  print("Solve using SVDs")
  time svd = time.time()
  u, sigma, vh = get_svd_decomposition(mat)
  time svd = time.time() - time svd
  check svd correctness(mat, u, sigma, vh)
  time_svd_solve = time.time()
  x \text{ svd} = \text{solve using svd}(u, \text{sigma, vh, b})
  time_svd += time.time() - time_svd_solve
  print("Time taken for SVD decomposition and solve = {} s".format(time svd))
   verify solution(mat, x svd, b)
   # Solve using Cholesky
  print("Solve using Cholesky")
  time cholesky = time.time()
  L = get cholesky decomposition(-1.0 * mat)
  time_cholesky = time.time() - time_cholesky
  check cholesky correctness(-1.0 * mat, L)
  time cholesky solve = time.time()
  x cholesky = solve using cholesky(L, -1.0 * b)
  time cholesky += time.time() - time cholesky solve
  print("Time taken for Cholesky factorization and Solve = {}
s".format(time_cholesky))
  verify solution(mat, x cholesky, b)
```

```
# Solve using np.linalg.solve
print("Solve using numpy.linalg.solve")
time_np = time.time()
x = solve_using_np_solve(mat, b)
time_np = time.time() - time_np
print("Time taken for Numpy factorization and Solve = {} s".format(time_np))
verify_solution(mat, x, b)

if __name__ == '__main__':
m = 4000
run_experiment(m)
```