Jacobi

March 7, 2023

Peter Chu, all code written between 8:05 PM March 6th to 1:20 AM March 7th

0.1 Jacobi Method

```
[1]: # import necessary libraries
import math as math
import numpy as np
```

0.1.1 Part a

```
[2]: # Input A = matrix, b = sol vector, x = quess vector, tol = tolerance, n = max_{\square}
     → iterations, start writing code 8:05
     def jacobi(A, b, x, tol, n):
         # Create necessary variables for formula
         \# D = diag \ matrix
         \# D_1 = inverse \ of \ D
         # U = strictly upper matrix * -1 to make entries negative
         \# L = strictly lower matrix * -1 to make entries negative
         \# T = D^{-1} * (L + U)
         # count is a ticker variable to keep track of number of iterations
         D = np.diag(np.diag(A))
         D_1 = np.linalg.inv(D)
         U = (np.triu(ls1) - D).dot(-1)
         L = (np.tril(ls1) - D).dot(-1)
         T = D 1.dot(L + U)
         count = 0
         # Do first iteration
         x_k = x
         x_k1 = (T.dot(x_k.T)) + (D_1.dot(b.T))
         count += 1
         # If x_0 = 0, //X_0//_inf = 0 we will have a divide by zero error.
         # Iterate one more time to avoid this and then enter loop
```

```
if(x_k.all() == 0):
       x_k = x_k1.T
       x_k1 = (T.dot(x_k.T)) + (D_1.dot(b.T))
       count += 1
   # Create stop value for exit condition
   StopVal = np.linalg.norm(abs(x_k1 - x_k), np.inf) / np.linalg.norm(x_k, np.
→inf)
   # Loop until we hit exit condition or we do n interations
   while(StopVal > tol and count < n):</pre>
       #Do jacobi method
       x_k = x_k1.T
       x_k1 = (T.dot(x_k.T)) + (D_1.dot(b.T))
       #Update stop value and add 1 to count
       StopVal = np.linalg.norm(abs(x_k1 - x_k), np.inf) / np.linalg.norm(x_k,_
\rightarrownp.inf)
       count += 1
   # Expect x^k vector and the number of iterations it took
   return x_k1, 'iterations:', count
```

0.1.2 Part b

```
# tolerance
tolVal = 0.00005
```

0.1.3 Part c

```
[5]: # Calculate spectral radius for ls1 and store value as lambda1
     D = np.diag(np.diag(ls1))
     U = (np.triu(ls1) - D).dot(-1)
     L = (np.tril(ls1) - D).dot(-1)
     D_1 = np.linalg.inv(D)
     T = D_1.dot(L + U)
     lambda1 = max(abs(np.linalg.eig(T)[0]))
     # Calculate spectral radius for ls2 and store value as lambda2
     D = np.diag(np.diag(1s2))
     U = (np.triu(ls2) - D).dot(-1)
     L = (np.tril(ls2) - D).dot(-1)
     D_1 = np.linalg.inv(D)
     T = D_1.dot(L + U)
     lambda2 = max(abs(np.linalg.eig(T)[0]))
     print('Spectral radius of linear system 1:', lambda1, '\n')
     print('Spectral radius of linear system 2:', lambda2)
```

Spectral radius of linear system 1: 1.0813325779705841e-05

Spectral radius of linear system 2: 1.1180339887498956

The Jacobi method gave a good approximation for linear system 1. Looking at $\rho(T_j)$ for linear system 1, we have $\rho(T_g) < 1$. Thus T_j is convergent and converges to certain values. As a result we get a good approximation because x^k converges as well. For linear system 2, we have that $\rho(T_j) > 1$ which results in T_j being divergent and thus x^k does not converge to some value. This results in a bad approximation.

0.2 Gauss-Seidel Method

0.2.1 Part a

```
[6]: # Input A = matrix, b = sol vector, x = quess vector, tol = tolerance, n = max_{\square}
      \rightarrow iterations
     def GaussSeidel(A, b, x, tol, n):
         # Create necessary variables for formula
         \# D = diag \ matrix
         # U = strictly upper matrix * -1 to make entries negative
         # L = strictly lower matrix * -1 to make entries negative
         \# DL_1 = (D - L)^-1, T = (D - l)^-1 * U
         # count is a ticker variable to keep track of number of iterations
         D = np.diag(np.diag(A))
         U = (np.triu(A) - D).dot(-1)
         L = (np.tril(A) - D).dot(-1)
         DL_1 = np.linalg.inv(D - L)
         T = DL 1.dot(U)
         count = 0
         # Do first iteration
         x_k = x
         x_k1 = T.dot(x_k.T) + DL_1.dot(b.T)
         # Update number of iterations
         count += 1
         #If x_0 = 0, //X_0//_inf = 0 we will have a divide by zero error.
         #Iterate one more time to avoid this and then enter loop
         if(x_k.all() == 0):
             x k = x k1.T
             x_k1 = T.dot(x_k.T) + DL_1.dot(b.T)
             count += 1
         # Create stop value for exit condition
         StopVal = np.linalg.norm(abs(x_k1 - x_k), np.inf) / np.linalg.norm(x_k, np.
      \hookrightarrowinf)
```

0.2.2 Part b

```
[7]: # Use np.newaxis on vectors to make them transposable correctly
     # Python does not know how to transpose 1 dimensional arrays, need to make 21
      \rightarrow dimensional to do so
     # ex. example = np.array([1,1,1])
     \# example.T = np.array([1,1,1]) != np.array([1],[1],[1]) <- desired transposed_{\sqcup}
      \rightarrowvector
     # linear system 1
     ls1 = np.array([[1,2,-2], [1,1,1], [2,2,1]])
     b1 = np.array([7,2,5])[np.newaxis]
     # Linear system 2
     ls2 = np.array([[2,-1,1], [2,2,2], [-1,-1,2]])
     b2 = np.array([-1,4,-5])[np.newaxis]
     # initial quess
     x_0 = np.array([0,0,0])[np.newaxis]
     # tolerance
     tolVal = 0.00005
```

```
[8]: # Run Gauss Seidel using defined variables

aprox1 = GaussSeidel(ls1, b1, x_0, tolVal, 25)
```

0.2.3 Part c

```
[9]: # Calculate spectral radius for ls1 and store value as lambda1, finish code 1:20
     D = np.diag(np.diag(ls1))
     U = (np.triu(ls1) - D).dot(-1)
     L = (np.tril(ls1) - D).dot(-1)
     DL_1 = np.linalg.inv(D - L)
     T = DL_1.dot(U)
     lambda1 = max(abs(np.linalg.eig(T)[0]))
     # Calculate spectral radius for ls2 and store value as lambda2
     D = np.diag(np.diag(1s2))
     U = (np.triu(1s2) - D).dot(-1)
     L = (np.tril(ls2) - D).dot(-1)
     DL_1 = np.linalg.inv(D - L)
     T = DL_1.dot(U)
     lambda2 = max(abs(np.linalg.eig(T)[0]))
     print('Spectral radius of linear system 1:', lambda1, '\n')
     print('Spectral radius of linear system 2:', lambda2)
```

Spectral radius of linear system 1: 2.0

Spectral radius of linear system 2: 0.5

The Gauss-Seidel method gave a good approximation for linear system 2. Looking at $\rho(T_g)$ for linear system 2, we have $\rho(T_g) = 0.5 < 1$. Thus T_g is convergent and converges to certain values. As a result we get a good approximation because x^k converges as well. For linear system 1, we have that $\rho(T_j) = 2 > 1$ which results in T_j being divergent and thus x^k does not converge to some value. This results in a bad approximation.