ECE 7650 (Advance Matrix Algorithm) REPORT

On

"FINAL EXAMINATION FOR FALL 2016"

By

Jamiu Babatunde Mojolagbe

(Student ID: #7804719)

antu baro

The Department of Electrical and Computer Engineering
University of Manitoba

Submitted to

312921/06

Ian Jeffrey, PhD

(Course Instructor)

Question 1(i)

This question was implemented in file named "Q1i.m". Random matrices of chosen dimension were used and the following results were obtained:

Case 1: For randomly chosen 'm'

Matrix Dimension	Size of Krylov Subspace (m)	Rank of V
10x10	5	6
20x20	12	13
50x50	35	12
100x100	98	8
200x200	131	7
300x300	211	7
400x400	313	1

Case 2: For fixed 'm' for different dimensions of input matrix A

Matrix Dimension	Size of Krylov Subspace (m)				
	Rank(V) for m=5	Rank(V) for m=10			
10x10	6	10			
20x20	6	11			
50x50	6	11			
100x100	6	11			
200x200	6	11			
300x300	6	11			
400x400	6	11			

Observation:

As it can be observed from the tables, it is obvious that choosing $V = [\underline{r} \ A\underline{r} \ A^2\underline{r} \ . \ . \ A^{m-1}\underline{r}]$ resulted in linearly dependent bases V such that, V is rank deficient. While Case 1 presented above tends to be a generic case and not totally clear, however observing Case 2 actually led to the conclusion that for a chosen size of Krylov subspace 'm', provided that $m < \dim(A)$, $\operatorname{rank}(V) = m+1$ and for $m = \dim(A)$, $\operatorname{rank}(V) = m$ for linearly independent bases.

Note: The ranks obtained are more than 'm' since dimension of V is actually nx(m+1) but in a specific case V_m of dimension nxm can be obtained from it by just removing the last column.

Question 1(ii)

Arnoldi method was implemented based on Modified Gram-Schmidt process as a function called 'arnoldi.m'. The driver program for this question is named 'Q1ii.m'. For the sake of clarity, 'Raw V' used to refer to the bases V obtained from $V = [\underline{r} \ A\underline{r} \ A^2\underline{r} \ . \ . \ A^{m-1}\underline{r}]$ while the ones obtained from Arnoldi iteration is termed 'Arnoldi V'.

Case 1: For randomly chosen 'm'

Matrix Dimension	Size of Krylov Subspace (m)	Rank of		
		Raw V	Arnoldi V	
10x10	9	10	10	
20x20	11	12	12	
50x50	41	13	42	
100x100	78	11	79	
200x200	153	8	154	
300x300	240	4	241	
400x400	297	2	298	

Case 2: For fixed 'm' for different dimensions of input matrix A

Matrix Dimension	Rank						
	V for	· m=5	V for m=10				
	Raw V	Arnoldi V	Raw V	Arnoldi V			
10x10	6	6	10	10			
20x20	6	6	11	11			
50x50	6	6	11	11			
100x100	6	6	11	11			
200x200	6	6	11	11			
300x300	6 6		11	11			
400x400	6	6	11	11			

Observation:

From the table shown above for Case 1, it can be observed that using Arnoldi method, the bases V obtained are linearly independent such the rank of V are now m+1 (for the reasons previously presented). If V_m is considered, that is V of dimension nxm, then rank(V) = m.

Hence forth, the table in Case 2 perfectly confirm the assertion made in the "Observation" section of the previous question, that is Question 1(i) above for Case 2.

Note: The ranks obtained are more than 'm' since dimension of V is actually nx(m+1) but in a specific case V_m of dimension nxm can be obtained from it by just removing the last column.

Question 2(i)

While it is necessary to say that Arnoldi process in itself does not directly access the entries of a given matrix but instead makes the matrix map vectors and as such reaches its conclusions from their images. However, it is important to show rigorously or partially rigorously that for $V_m{}^TAV_m = H_m$, when m=n that the eigenvalues of $H_m = H_n$ are equal to the eigenvalues of A.

Recall that square matrices say, A and B, are related by:

$$\mathbf{B} = \mathbf{T}^{-1}\mathbf{A}\mathbf{T} = \mathbf{T}^{T}\mathbf{A}\mathbf{T}$$

where $T^{-1} => Non-singular matrix (that is, invertible)$

The transformation represented by $T^{-1}AT$ above is known as similarity transformation or conjugation by matrix T.

Suppose:

$$\mathbf{A} = \begin{bmatrix} 0 & 0 \\ 1 & 0 \end{bmatrix} \qquad \qquad \mathbf{B} = \begin{bmatrix} 0 & 1 \\ 0 & 0 \end{bmatrix}$$

These two matrices are similar under transformation or conjugation by T, such that T is given by

$$\mathbf{T} = \begin{bmatrix} 0 & 1 \\ 1 & 0 \end{bmatrix}$$

Since a linear transformation is the same as a matrix after a basis, says bi, is chosen; then it can be shown that:

$$\mathbf{L}(\Sigma \lambda_i b_i) = \Sigma \mathbf{a}_{ii} \lambda_i b_i$$

Now, changing the basis, changes the co-efficient of the matrix such that

$$\mathbf{L}(\Sigma \gamma_i \mathbf{e}_i) = \Sigma \mathbf{a}_{ji} \gamma_i \mathbf{e}_j$$

If operator L(V) = AV uses standard basis (Euclidean basis), then L is the matrix TAT^{-1} with basis

$$b_i = Te_i$$

Note: *Notice how the eigenvalues are represented to depict that they are images of each other.*

Question 2(ii)

To proof that for the largest eigenvalue of A and H_m are (approximately) equal, a file named "Q2ii.m" was used. Random matrices were used as well as random initial vector and H_m was calculate from Arnoldi method previously implemented. The following results were obtained:

Case 1: For randomly chosen 'm'

Matrix	Size of Krylov Subspace	Maximum Eigenva	lue of
Dimension	(m)	A	H _m
10x10	8	-3.9721	-3.9752
20x20	18	-3.735+1.5751i	-3.7347+1.5762i
50x50	33	8.3546	8.3544
100x100	93	-2.03653+9.94841i	-2.03653+9.94841i
200x200	171	-2.82874+15.0757i	-2.82874+15.0757i
300x300	266	-4.39007+16.9614i	-4.39007+16.9614i
400x400	333	19.5548+6.06034i	19.5548+6.06034i

Case 2: For fixed 'm' for different dimensions of input matrix A

Matrix	Maximum Eigenvalue					
Dimension	m	=5	m=	=10		
	A H _m		A	H _m		
10x10	2.1836+2.2106i	-0.99642+2.7697i	-2.6912	-2.6912		
20x20	5.1404	4.9816	4.3142+1.3456i	-4.4952		
50x50	-8.6843	-8.8097	-2.5343+7.2262i	-5.2299+5.1851i		
100x100	-8.62056+5.87658i	-6.8939	9.28855+4.45615i	8.4777+4.5603i		
200x200	15.1315+2.55873i	-7.7372	-8.25263+12.1035i	8.94859+10.3865i		
300x300	-9.26965+15.9366i	-10.9218+7.77496i	18.424	-8.13645+11.1553i		
400x400	-7.43289+18.9942i	-9.83433+6.59414i	20.7206	-10.4521+12.1804i		

Observation:

Though Case 1 uses randomly chosen 'm' but it really does so such that it is very useful for arriving at a very good conclusion here. Why? Because the randomly chosen 'm's in Case 1 above tend to be less that the dimension of a but are not much less, which is good for our purpose here. Therefore, from table for Case 1, it can be concluded that when m < n but not much less, that is when the value of m approaches the dimension n, then the eigenvalues of m tend to mimic eigenvalues of m and in fact some cases in Case 1 above the eigenvalue of m is equal to eigenvalue of m for that given precision.

Observing Case 2 above, however, clear things up and it can be concluded that when m << n then the eigenvalues (maximums are used here) of H_m are not in any way approximately equals to eigenvalues of A.

Question 2(iii)

Give a matrix A^{nxn} , show that when m < n, $A = V_m H_m V_m^T$ is true or false.

Finally, it can be concluded that for the inequality n < m, statement $A = V_m H_m V_m^T$ is not true (that is false) for given real matrix A^{nxn} .

Question 3(i)

Deriving the directional derivative of cost functional given below in terms of the arbitrary direction vector \mathbf{h} using the real scalar parameter $\mathbf{\epsilon}$:

$$||b - Ax||_2^2$$

Recall that the directional derivative is given by:

$$\nabla_h f(x) = \frac{df(\underline{x} + \varepsilon \underline{h})}{d\varepsilon} \bigg|_{\varepsilon = 0} = \lim_{\varepsilon \to 0} \frac{df(\underline{x} + \varepsilon \underline{h}) - f(\underline{x})}{\varepsilon}$$

Now,
$$f(x) = \|b - Ax\|_{2}^{2}$$

 $= \langle b - Ax, b - Ax \rangle$

$$= \lim_{\epsilon \to 0} \frac{1}{\epsilon} \langle b - Ax - \epsilon Ah, b - Ax - \epsilon Ah \rangle - \langle b - Ax, b - Ax \rangle$$
since $r = b - Ax$

$$= \lim_{\epsilon \to 0} \frac{1}{\epsilon} \langle r - \epsilon Ah, r - \epsilon Ah \rangle - \langle r, r \rangle$$

$$= \lim_{\epsilon \to 0} \frac{1}{\epsilon} ((r - \epsilon Ah)^{T} (r - \epsilon Ah) - r^{T} r)$$

$$= \lim_{\epsilon \to 0} \frac{1}{\epsilon} ((r - \epsilon Ah)^{T} r + (r - \epsilon Ah)^{T} (-\epsilon Ah) - r^{T} r)$$

$$= \lim_{\epsilon \to 0} \frac{1}{\epsilon} (r^{T} r + (-\epsilon Ah)^{T} r + r^{T} (-\epsilon Ah) + (-\epsilon Ah)^{T} (-\epsilon Ah) - r^{T} r)$$

$$= \lim_{\epsilon \to 0} \frac{1}{\epsilon} (\langle r, -\epsilon Ah \rangle + \langle -\epsilon Ah, r \rangle + \langle -\epsilon Ah, -\epsilon Ah \rangle)$$

$$= \lim_{\epsilon \to 0} \frac{1}{\epsilon} (-\epsilon \langle r, Ah \rangle - \epsilon \langle -Ah, r \rangle - \epsilon^{2} \langle Ah, Ah \rangle)$$

$$= -\langle r, Ah \rangle - \langle Ah, r \rangle$$

$$= -2 \langle r, Ah \rangle$$

If $\mathbf{h} = \mathbf{r}$, then

$$\nabla_h f(x) = \frac{df(\underline{x} + \varepsilon \underline{h})}{d\varepsilon} \Big|_{\varepsilon=0} = -2 < r, \ Ar >$$

Question 3(ii)

For step length α , direction $\underline{\mathbf{d}}$, evaluate functional $(\underline{\mathbf{x}} + \alpha \underline{\mathbf{d}})$

$$\alpha = \frac{argmin}{\alpha} < b - Ax - \alpha Ad, \ b - Ax - \alpha Ad > \alpha$$

Recall, r = b - Ax

$$\alpha = \frac{argmin}{\alpha} < r - \alpha Ad, r - \alpha Ad >$$

Given that, d = r

$$\alpha = \frac{argmin}{\alpha} < r - \alpha Ar, \ r - \alpha Ar >$$

$$= \frac{argmin}{\alpha} ((r - \alpha Ar)^T (r - \alpha Ar))$$

$$= \frac{argmin}{\alpha} ((r - \alpha Ar)^T r + (r - \alpha Ar)^T (-\alpha Ar))$$

$$= \frac{argmin}{\alpha} (r^T r + (-\alpha Ar)^T r + r^T (-\alpha Ar) + (-\alpha Ar)^T (-\alpha Ar))$$

$$= \frac{argmin}{\alpha} (r^T r - \alpha r^T A^T r - \alpha r^T Ar + (-\alpha r^T A^T) (-\alpha Ar))$$

$$= \frac{argmin}{\alpha} (r^T r - \alpha r^T A^T r - \alpha r^T Ar + \alpha^2 r^T A^T Ar)$$

Now, derivative of the function with respect to α is equal to 0

$$= -r^{T}A^{T}r - r^{T}Ar + 2\alpha r^{T}A^{T}Ar = 0$$
$$2\alpha r^{T}A^{T}Ar = r^{T}A^{T}r + r^{T}Ar$$

$$\alpha = \frac{r^T A^T r + r^T A r}{2r^T A^T A r}$$

$$\alpha = \frac{\langle r, Ar \rangle + \langle Ar, r \rangle}{2 \langle Ar, Ar \rangle} = \frac{2 \langle r, Ar \rangle}{2 \langle Ar, Ar \rangle} = \frac{\langle r, Ar \rangle}{\langle Ar, Ar \rangle}$$

$$\alpha = \frac{\langle r, Ar \rangle}{\langle Ar, Ar \rangle}$$

Comparing the result obtain for α above with the alpha in **Minimum Residual 1-D**, shows that both $\alpha's$ has the same formulas

Question 3(iii)

From 3(i) above,

$$\nabla_h f(x) = -2 < r$$
, $Ar >$

Such that,

$$-\nabla_h f(x) = A^T r$$

Therefore:

$$\alpha = \frac{argmin}{\alpha} < b - Ax - \alpha Ad, \qquad b - Ax - \alpha Ad >$$

$$\alpha = \frac{argmin}{\alpha} < r - \alpha Ad, \quad r - \alpha Ad > \qquad \text{since } \mathbf{r} = \mathbf{b} - \mathbf{A}\mathbf{x}$$

$$= \frac{argmin}{\alpha} < r - \alpha AA^{T}r, \quad r - \alpha AA^{T}r > \qquad \text{since } \mathbf{d} = -\nabla_{\mathbf{h}}\mathbf{f}(\mathbf{x}) = \mathbf{A}^{T}\mathbf{r}$$

$$= \frac{argmin}{\alpha} ((r - \alpha AA^{T}r)^{T}(r - \alpha AA^{T}r))$$

$$= \frac{argmin}{\alpha} ((r - \alpha AA^{T}r)^{T}r + (r - \alpha AA^{T}r)^{T}(-\alpha AA^{T}r))$$

$$= \frac{argmin}{\alpha} (r^{T}r + (-\alpha AA^{T}r)^{T}r - \alpha r^{T}AA^{T}r + (-\alpha AA^{T}r)^{T}(-\alpha AA^{T}r))$$

$$= \frac{argmin}{\alpha} (r^{T}r - \alpha r^{T}AA^{T}r - \alpha r^{T}AA^{T}r + \alpha^{2}r^{T}AA^{T}AA^{T}r)$$

$$= \frac{argmin}{\alpha} (r^{T}r - 2\alpha r^{T}AA^{T}r + \alpha^{2}r^{T}AA^{T}AA^{T}r)$$

The derivative of the function with respect to α is equal to 0

$$=> (-2r^{T}AA^{T}r + 2\alpha r^{T}AA^{T}AA^{T}r) = 0$$
$$2\alpha r^{T}AA^{T}AA^{T}r = -2r^{T}AA^{T}r$$

$$\alpha = \frac{r^T A A^T r}{r^T A A^T A A^T r} = \frac{r^T r}{r^T A A^T r} = \frac{\langle r, r \rangle}{\langle Ar, Ar \rangle} =$$

$$\alpha = \frac{\langle r, Ar \rangle}{\langle Ar, Ar \rangle} = \frac{\|r\|_2^2}{\|Ar\|_2^2}$$

Finally, the obtained result is identical to alpha in Residual Norm Steepest Descent algorithm.

Question 4(i)

Recall from Arnoldi that,

$$\begin{aligned} \mathbf{H}_m &= \mathbf{V}_m^T \mathbf{A}_m \mathbf{V}_m \\ & \text{where } \mathbf{V}_m \text{ is } \mathbf{n} \mathbf{x} \mathbf{m} \\ & \mathbf{H}_m \text{ is } \mathbf{m} \mathbf{x} \mathbf{m} \\ & A \quad \text{is } \mathbf{n} \mathbf{x} \mathbf{n} \text{ and symmetric} \end{aligned}$$

Let's consider the (ij)th entry of matrix H_m

$$(H_m)_{ij} = (V_m^T)_i A(V_m)_j = h_{ij} = v_i^T A v_j$$

Recall that, orthonormal basis for

$$K_{j}(A,b) = \{v_{1} \quad v_{2} \quad \dots \quad v_{j}\}$$
Thus;
$$Av_{j} \in K_{j+1}(A,b) = span\{v_{1} \quad v_{2} \quad \dots \quad v_{j+1}\}$$
Hence,
$$h_{ij} = v_{i}^{T}Av_{j} = 0 \qquad \text{for } \mathbf{i} > \mathbf{j} + \mathbf{1}$$
Since,
$$v_{i} \perp span\{v_{1} \quad v_{2} \quad \dots \quad v_{j+1}\} \quad \textit{for } \quad i > j + 1$$
Such that,
$$h_{ij} = h_{ji} = v_{j}^{T}Av_{i}$$

Since matrix **A** given is symmetric, then we have symmetric \mathbf{H}_{m} , thus $\mathbf{h}_{ij} = \mathbf{0}$ for j > i + 1 as a result of the reason presented/proven above.

Therefore, it can be concluded that for a symmetric matrix A, H_m produce from Arnoldi process is symmetric tridiagonal.

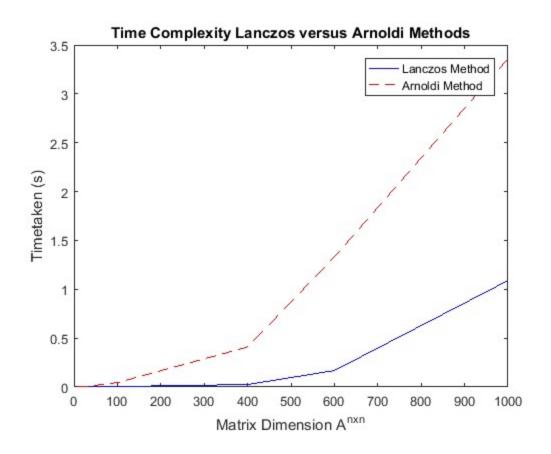
Other Details: For the purpose of confirmation and personal conviction, this proof was tested in file named "Q4i.m" for random input matrix, symmetric tridiagonal H_m was obtained.

Question 4(ii)

Lanczos method was implemented in function named "lanczos.m". The driver program for this question is named "Q4ii.m". The following results were obtained for different cases tested.

Case 1: For randomly chosen 'm'

Matrix	Size of Krylov Subspace	Time taken in seconds			
Dimension	(m)	Arnoldi Method	Lanczos Method		
10x10	8	0.00158832017805	0.00113115987732		
20x20	11	0.00084607676844	0.000893040266430		
100x100	76	0.04512201520370	0.004989871661235		
400x400	210	0.40976936152758	0.025197017377595		
600x600	490	1.33099212957816	0.168348730389529		
1000x1000	694	3.36032083407172	1.089681569138520		



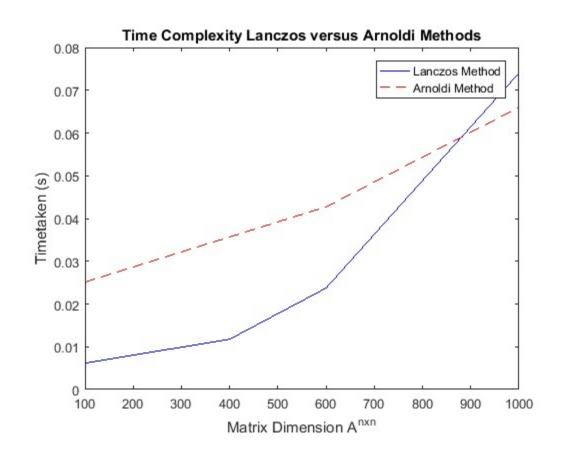
Prove that the algorithm worked was also performed and the result obtained from the various norms were as follows:

```
****** Proofs of Orthogonality of Resulting Bases *********
Norm of Km from Arnoldi:
Norm of Km from Lanczos:
Error: norm(Lanczos(Km) - Arnoldi(Km)):
                                    2.0147e-09
Inner Product (Arnoldi): <v(m), v(m)>:
Inner Product (Arnoldi): <v(m), v(m-1)>:
                                    -1.5404e-15
Inner Product (Arnoldi): \langle v(2), v(m) \rangle: -5.3257e-12
Inner Product (Lanczos): \langle v(m), v(m) \rangle: 1
Inner Product (Lanczos): \langle v1(m), v1(m-1) \rangle: -1.3878e-17
Inner Product (Lanczos): \langle v1(2), v1(m) \rangle: 6.4199e-14
******* Proofs of Orthogonality of Resulting Bases *********
Norm of Km from Arnoldi:
Norm of Km from Lanczos:
Error: norm(Lanczos(Km) - Arnoldi(Km)):
                                   2.8877e-06
Inner Product (Arnoldi): <v(m), v(m)>:
Inner Product (Arnoldi): \langle v(m), v(m-1) \rangle: -7.7716e-16
Inner Product (Arnoldi): \langle v(2), v(m) \rangle: 1.6214e-08
Inner Product (Lanczos): <v(m), v(m)>:
Inner Product (Lanczos): \langle v1(m), v1(m-1) \rangle: 4.3021e-16
Inner Product (Lanczos): <v1(2), v1(m)>: -1.3128e-14
______
******* Proofs of Orthogonality of Resulting Bases *********
Norm of Km from Arnoldi:
                                     1.5959
Norm of Km from Lanczos:
                                     1.4142
Error: norm(Lanczos(Km) - Arnoldi(Km)):
                                     2.18
Inner Product (Arnoldi): <v(m), v(m)>:
Inner Product (Arnoldi): <v(m), v(m-1)>:
                                    7.9353e-15
Inner Product (Arnoldi): <v(2), v(m)>:
                                    4.1302e-06
Inner Product (Lanczos): <v(m), v(m)>:
Inner Product (Lanczos): \langle v1(m), v1(m-1) \rangle: 8.2391e-15
Inner Product (Lanczos): \langle v1(2), v1(m) \rangle: -0.0012023
______
```

```
******* Proofs of Orthogonality of Resulting Bases **********
Norm of Km from Arnoldi:
                                     1.8428
Norm of Km from Lanczos:
                                     1.7321
Error: norm(Lanczos(Km) - Arnoldi(Km)):
                                     2.5176
Inner Product (Arnoldi): <v(m), v(m)>:
Inner Product (Arnoldi): \langle v(m), v(m-1) \rangle: 6.7521e-16
Inner Product (Arnoldi): <v(2), v(m)>:
                                   0.046588
Inner Product (Lanczos): \langle v(m), v(m) \rangle: 1
Inner Product (Lanczos): \langle v1(m), v1(m-1) \rangle: -2.8027e-14
Inner Product (Lanczos): <v1(2), v1(m)>: 0.012046
******* Proofs of Orthogonality of Resulting Bases *********
Norm of Km from Arnoldi:
                                     2.0865
Norm of Km from Lanczos:
                                     2.4495
Error: norm(Lanczos(Km) - Arnoldi(Km)):
                                     3.0614
Inner Product (Arnoldi): <v(m), v(m)>:
Inner Product (Arnoldi): <v(m), v(m-1)>:
                                     1.1645e-13
Inner Product (Arnoldi): <v(2), v(m)>:
                                    3.639e-08
Inner Product (Lanczos): <v(m), v(m)>:
Inner Product (Lanczos): <v1(m), v1(m-1)>: -2.8271e-14
Inner Product (Lanczos): \langle v1(2), v1(m) \rangle: -0.022383
******* Proofs of Orthogonality of Resulting Bases **********
Norm of Km from Arnoldi:
                                     1.9411
Norm of Km from Lanczos:
                                     2.6451
Error: norm(Lanczos(Km) - Arnoldi(Km)):
                                     3.1447
Inner Product (Arnoldi): <v(m), v(m)>:
Inner Product (Arnoldi): <v(m), v(m-1)>:
                                     -1.2157e-13
Inner Product (Arnoldi): <v(2), v(m)>:
                                     -2.7093e-07
Inner Product (Lanczos): <v(m), v(m)>:
Inner Product (Lanczos): \langle v1(m), v1(m-1) \rangle: -1.7757e-14
Inner Product (Lanczos): \langle v1(2), v1(m) \rangle: -0.034224
```

Case 2: For fixed 'm' and chosen to be 50, the following time complexity results were obtained:

Matrix	Size of Krylov Subspace	Time taken ir	1 seconds
Dimension	(m)	Arnoldi Method	Lanczos Method
100x100	50	0.0251170326700840	0.00617166405982563
400x400	50	0.0356834528150765	0.0117577520041122
600x600	50	0.0427276106109621	0.0237334439910755
1000x1000	50	0.0659936210361248	0.0740232784851043



Observation:

From the foregoing, it can be observed from the results and plots that the implementation worked and the error tends to be very small. However, a close look at the results showed also that the bases produced are truly orthogonal but there seems to be loss of orthogonality when comparing says 1^{st} and 500^{th} vector as the error obtained, for **Case 1** for matrix dimensions [$600 \times 600 \times 1000 \times 1000$] for $\mathbf{m} = [690 \times 694]$, clearly shown.

Moreover, comparing the complexity, it can be seen from the graphs and tables presented that in overall Lanczos method is fantastically faster than Arnoldi. However, when the m is fixed at 50 Arnoldi tends to be faster. Why? This misery was uncovered when a new test was performed and the size of Krylov subspace m = 500 and matrix dimension is 1000×1000 . The time obtained was 2.6839 for Arnoldi and 0.79408 for Lanczos; this thereby led to the conclusion that, when m is chosen to m << n there is possibility that Arnoldi would be faster than Lanczos.

Question 4(iii)

Recall,

$$\mathbf{H}_m = \mathbf{V}_m{}^T \mathbf{A}_m \mathbf{V}_m$$

For symmetric trdiagonal let $H_m = T_m$

$$T_m = V_m^T A V_m$$

Such that,

$$y_m = T_m^{-1} A \beta e_1$$

$$x_m = x_0 + V_m y_m$$
 ----- (iii)

Using LU decomposition,

$$T_m = L_m U_m$$

Where L_m = Unit lower bi-diagonal U_m = Upper bi-diagonal matrix

equation (iii) becomes:

$$x_m = x_0 + V_m U_m^{-1} L_m^{-1} (\beta e_1)$$

$$let P_m = V_m U_m^{-1}$$

$$z_m = L_m^{-1}(\beta e_1)$$

Therefore, $x_m = x_0 + P_m z_m$

The last column of matrix P_m is given by:

$$p_m = \eta_m^{-1}[V_m - \beta_m p_{m-1}]$$
 where $\eta_m = mth \ G. E.$

$$\lambda_m = \frac{\beta_m}{\eta_m}$$

$$\eta_m = \alpha_m - \lambda_m \beta_m$$

Recall that,

$$z_m = \begin{bmatrix} z_{m-1} \\ \zeta_m \end{bmatrix}$$

$$\zeta_m = -\lambda_m \zeta_{m-1}$$

Finally, update equation is given by:

$$x_m = x_0 + \zeta_m p_m$$

Implementation Note

The proved algorithm above was implemented in the file named "lanczosfom.m" as a function. The driver program for this function is named "Q4iii.m". As contained in the driver program, random matrices can be used or an already loaded matrix and right hand side obtained from Finite Element Method solution to a particular problem. Again, for clear comparison, Restarted FOM was implemented to serve as basis for our comparison here; meanwhile the two implementation are compared to the solution obtained from MATLAB's backslash operator which now serves as analytic solution for validation and comparison purposes. While the input matrix dimension is fixed, 157x157, random matrix generation provided a good alternative, though the former is a real time data.

Results and Discussions

The following results were obtained when the implemented function was compared against Restarted Full Orthogonalization Method (FOM):

Case 1: For Krylov subspace, m = 20 and error tolerance, tol = 1e-06

Where **x** = solution from **lanczosfom.m**, **x1** = solution from restarted FOM, **xe** = A\b **LLU** is used here to refer to the solution we obtained based on LU **RFOM** is used here to refer to the solution obtained from restarted FOM

Matrix	Time taken (s)		Proof that it works		Number of iterations	
Dimension	LLU	RFOM	norm(x-xe)	norm(x1-xe)	LLU	RFOM
50x50	0.0179180	0.0085334	6.7281e-08	6.7281e-08	8	8
100x100	0.0025287	0.0036320	9.048e-08	9.048e-08	7	7
200x200	0.0026633	0.0036283	1.2478e-08	1.2478e-08	7	7
500x500	0.005898	0.005949	2.1109e-08	2.1109e-08	6	6

Case 2: For Krylov subspace, m = 20 and error tolerance, tol = 1e-10

Matrix	Time taken (s)		Proof that i	Proof that it works		Number of iterations	
Dimension	LLU	RFOM	norm(x-xe)	norm(x1-xe)	LLU	RFOM	
50x50	0.0015623	0.0032273	2.7607e-12	2.7608e-12	12	12	
100x100	0.0031587	0.0032486	2.5796e-12	2.5798e-12	11	11	
200x200	0.0030508	0.0037120	9.0409e-13	9.0451e-13	10	10	
500x500	0.0033564	0.0027470	7.0452e-13	7.0452e-13	9	9	

Case 3: For Krylov subspace, m = 50 and error tolerance, tol = 1e-6

Matrix	Time taken (s)		Proof that it works		Number of iterations	
Dimension	LLU	RFOM	norm(x-xe)	norm(x1-xe)	LLU	RFOM
50x50	0.0015847	0.0019662	6.3966e-08	6.3966e-08	8	8
100x100	0.0023119	0.0034027	8.8003e-08	8.8003e-08	7	7
200x200	0.0025459	0.0024315	8.7779e-09	8.7779e-09	7	7
500x500	0.0043801	0.0046013	1.9508e-08	1.9508e-08	6	6

Observation:

It can be observed from the table that the implemented algorithm worked. It can be seen result wise that increasing the size of Krylov subspace **m** decreases the number of iterations and vice versa. It can also be said that, setting error tolerance very low increases the number of iterations.

Regarding comparison with restarted FOM, it can be observed that this approach tends to reduce the computational time when compared to the restarted FOM as shown in the table above. However, regard the error obtained, there was also slight but not obvious improvement in the error, this is not really obvious because of the chosen precision, though, Case 2 above somehow show this (for matrix dimension 100x100).

Question 5(i)

Solving Ax = b

Such that: $A \in \mathbb{R}^{n \times m}$ where n > m, that is, over determined system

It is what noting that over determined systems fall under least square problem.

Why QR for this system?

QR provides efficient way in solving non-square system such that Q factor provides orthogonal bases for the span of the columns of given matrix A.

How do we do this?

 \Rightarrow

Soving $Ax \approx b$ as given means, minimizing the functional $||b - Ax||_2$

It should be noted that transforming that difference by an orthogonal matrix preserve the norm (Euclidean norm).

Therefore,
$$||Q^T(b - Ax)||_2 = ||Q^Tb - Q^TAx||_2 = ||Q^Tb - Rx||_2$$

Which can be represented in block form as:

$$\left\| \begin{bmatrix} Q_1^T b \\ Q_2^T b \end{bmatrix} - \begin{bmatrix} R_1 \\ 0 \end{bmatrix} \right\|_2 = \left\| \begin{bmatrix} Q_1^T b - R_1 x \\ Q_2^T b \end{bmatrix} \right\|_2$$
$$\| Q_1^T b - R_1 x \|_2 + \| Q_2^T b \|_2$$

Since the second term above contains no 'x' then it is not relevant to the functional minimization.

Therefore, minimizing the first term is equivalent to minimizing the norm of the first term.

Given that R_1 to be non – singular (once A is not singular itself)

$$Q_1^T b - R_1 x = 0$$

$$Q_1^T b = R_1 x$$

$$R_1 x = Q_1^T b$$

Such that the solution, x, can be obtained from the above expression by applying backward substitution. Conclusively, QR factorization reduces the over determined system to a square one. It should also be stated that Q_2 need not to be computed as the QR can be obtained more efficiently if only Q_1 is calculated rather than obtaining all of the Q. This is what "economy size" QR in MATLAB does.

Question 5(ii)

Given:

$$x = x_0 + V(W^T A V)^{-1} W^T r$$
 _____(i) where, W = AV and V = V_m

Equation (i) can be written as:

$$x = x_0 + V(W^T W)^{-1} W^T r$$

Let
$$y_m = (W^T W)^{-1} W^T r$$
 such that, $x = x_0 + V_m y_m$

Thus: simplify y_m

Recall from least-square that, $(A^T A)^{-1} A = A^{-1} (A^T)^{-1} A^T = A^{-1}$

Therefore,
$$y_m = (W^T W)^{-1} W^T r = W^{-1} r$$

Since W = AV as given above, then,

$$y_m = (AV_m)^{-1}r$$

Recall from Arnoldi,

$$AV_m = V_{m+1}H_{m+1} \approx V_mH_m$$

 \Rightarrow

$$y_m = (V_m H_m)^{-1} r$$

 $y_m = H_m^{-1} (V_m^T r)$

Where H_m is mxm, V_m is nxm, H_{m+1} is (m+1)xm and V_{m+1} is nx(m+1)

This result can be said to be approximately and theoretically equal to the result of minimizing the functional $\|b - A(x^{(0)} + V_m y_m)\|_2$ which gives $x = x^{(0)} + V_m y_m$

Where

$$y_m = H_m^{-1} (V_m^T r)$$
 and letting $\beta = ||r||_2$

$$v_1 = \frac{r}{||r||_2} = \frac{r}{\beta}$$

$$V_m^T r = V_m^T \beta v_1 = \beta e_1$$

$$y_m = H_m^{-1} (\beta e_1)$$

 \Rightarrow

Question 5(iii)

Implementation Note

Restarted GMRES was implemented based on the guidelines provided, however Modified Gram-Schmidt variant of Lanczos was used. The QR decomposition is obtained via Givens Rotation – which is implemented as a function with name "givens.m" and called within the main function. The main function implementation file name is "gmresrestart.m" and the driver program for it is named "Q5iii.m". Another implementation setback is that inner residual tracking was not implemented, thereby returning inner iteration as the same value as the chosen size of Krylov subspace "m". The screen dump of the outer residual is implemented while there is no screen dump for the inner residual. The solution obtained is compared against solution calculated from MATLAB's backslash operator to confirm that the implementation worked.

Results and Discussions

Case 1: For size of Krylov subspace, m = 3, maximum iteration = 40, error tolerance, tol = $1e^{-12}$ and matrix dimension 1000x1000 the following results were obtained:

Where, $\mathbf{x} = \text{solution obtained from implemented Restarted GMRES}$ $\mathbf{xe} = \text{solution obtained from the backslash operator}$

Case 2: For size of Krylov subspace, m = 10, maximum iteration = 40, error tolerance, tol = $1e^{-12}$ and matrix dimension 1000x1000 the following results were obtained:

Case 3: For size of Krylov subspace, m = 3, maximum iteration = 40, error tolerance, tol = $1e^{-6}$ and matrix dimension 1000x1000 the following results were obtained:

Case 4: For size of Krylov subspace, m = 10, maximum iteration = 40, error tolerance, tol = $1e^{-6}$ and matrix dimension 1000x1000 the following results were obtained:

Observation:

From the screen dumps of the results presented above, it can be concluded that the implemented Restarted GMRES worked as the norm of the obtained solution compared to the solution computed by backslash operator tend to be negligible – it is to the machine precision. Having said that, it can also be observed that as the outer iteration increases the residual decreases, that is, technically speaking, the solution tends to be converging as the number of iterations increases until it finally converged.

Finally, another thing that can be noted from the result is what has been previously asserted, that number of iterations increase with the decrease in the size of "m" for a given fixed tolerance value and vice versa. On the same note, setting tolerance very small for fixed "m" and fixed matrix dimension increases the number of iterations and vice versa.

Question 5(iv)

GMRES will converge for any positive definite matrix. It's often used in practice because most practical application problems resulted in either symmetric or non – symmetric positive definite matrices. And owing to the fact that it is efficient compared to other techniques for the same problem makes it a better choice. In fact, as used in this exam, there is a matrix with its RHS included in the folder named "data" that was obtained from charge distribution on a square plate by Second Order Finite Element Method which turned out to be symmetric and positive definite. This matrix was used throughout the testing phase of this exam. For this reason, it also makes me a fan of Restarted GMRES if a problem of such is to be solved.