

Matrix Solver
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Lecture – 50
Lanczos Biorthogonalization and BCG Algorithm

Welcome. So, we are looking into Krylov subspace based method and we have looked into a number of methods involving Arnoldi's, method full orthogonal method, generalized minimum residual method. And, then we looked into Krylov subspace methods which is same as Arnoldi's or full orthogonalization method, but specifically for symmetric matrices. And we saw that there are a number of interesting things happen for symmetric matrices especially we get a conjugacy of the auxiliary vector and the auxiliary vector comes due to the fact that the h matrix becomes a triangular matrix or symmetric matrix. And from there we have looked into Lanczos algorithm which has been modified as direct Lanczos and further into conjugate gradient method, which is a very efficient and very simple method for solving symmetric matrix problems.

Now, the goal of the present lecture will be and of course, this will be continued in subsequent lectures that whether conjugate gradient type simple and efficient solver can be designed for general any general matrix not only for the symmetric matrices. Earlier you have seen that most of the matrix solvers has certain limitations. In a sense that Jacobi gauss Seidel or (Refer Time: 01:45) are applicable only for diagonally dominant or irreducibly diagonally dominant systems. And conjugate gradient steepest descent are applicable for symmetric matrices.

If we look into symmetric positive definite matrices especially steepest descent; if we look into GMRES that is probably the only method which have looked into right now is applicable for more wider classes of matrix, even the matrices asymmetric GMRES can be applied; if any it is not strictly diagonally domain and GMRES can be applied. However, so our present target will be looking into general purpose matrix as the matrix may not be diagonally dominant, may not be symmetric, but the matrix has to be nonsingular for having a solution.

So, for any nonsingular matrix can we develop a method and we will explore Krylov space subspace methods more to look into that part. So, what we will look in this

particular class is Lanczos bi orthogonalization analyzation which is a variant of Krylov subspace method and then how a conjugate gradient type of methods can be developed for general matrices for non symmetric matrices using Lanczos by orthogonalization.


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Krylov subspace

The projection method seeks an approximate solution x_m from an affine subspace $x_0 + K_m$ by imposing condition $b - Ax_m \perp L_m$. L_m is another subspace of dimension m , x_0 is the initial guess.

In the case of Krylov subspace methods, $K_m = K_m(A, r_0)$, $r_0 = (b - Ax_0)$ in R^n

$$K_m = \text{span}\{r_0, Ar_0, A^2 r_0, \dots, A^{m-1} r_0\}$$



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We have earlier done with Krylov subspace methods and the projection method for solving $Ax = b$ using Krylov subspace method, is a method that seeks an approximate solution x_m from an affine subspace $x_0 + K_m$ by imposing the condition that the residual $b - Ax_m$ is orthogonal to L_m .

L_m is another subspace of dimension m , x_0 is initial guess, K_m is also a subspace of dimension n in case of Krylov subspace method K_m is the Krylov subspace of A and r_0 , A is the matrix and r_0 is the initial residual $b - Ax_0$ in R^n . And K_m the Krylov subspace is defined as span of $r_0, Ar_0, A^2 r_0, \dots, A^{m-1} r_0$ that is A^m dimensional space in R^n .

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Krylov subspace methods for matrix solver

The different versions of Krylov subspace methods arise from different choice of L_m and from the way in which the system is preconditioned. Two broad choices for L_m give rise to the best known techniques:

$L_m = K_m$ FOM or Full orthogonal

$L_m = AK_m$ GMRES, MINRES

Conjugate Gradient
Method for symmetric
matrices using
Lanczos
orthogonalization

Lanczos method finds the approximate solution $x_m = x_0 + V_m y_m$

Using inversion of a tridiagonal matrix algorithm $y_m = T_m^{-1}(\beta e_1)$

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The different versions of Krylov subspace methods arise from different choice of L_m and the way the system is preconditioned. Preconditioning will look in look later we have earlier discussed that this will come later. There are true broad two broad choices for L_m which you have seen till now in the earlier examples of Krylov subspace methods; one is L_m is equal to K_m which is full orthogonal or method or FOM another is L_m is equal to AK_m which is GMRES or minimum residual method.

This is a complete orthogonal projection and the second one is an oblique projection. Now if we look into orthogonal projection in detail then we will see that for symmetric matrix we can get Lanczos orthogonalization and can get conjugant method, gradient method from full orthogonal method. Lanczos method will find the approximate solution as x_m is equal to x_0 plus $V_m y_m$ where V_m is the basis of Krylov subspace. And y_m can be obtained by inverting the tri diagonal matrix T_m which comes from Lanczos orthogonalization of the Heisenberg matrix upper part of the Heisenberg matrix basically assumes a tri diagonal form we have looked into that.

So, y_m is equal to T_m inverse beta e_1 . And we have seen that inversion of a tri diagonal matrix is simple and as well as TDMA type of algorithms can be used for direct inversion; there can be there are recursive relations which come and this makes the problem simpler.

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Conjugate Gradient Method

- A faster method in terms of number of iterations (convergence rate) and operations in each iteration step
- Uses A -conjugacy of auxiliary vector P ($p_i^T A p_j = 0$ if $i \neq j$):

$$\begin{aligned}
 T_m &= L_m U_m \\
 x_m &= x_0 + V_m (L_m U_m)^{-1} y \quad \Rightarrow P_m^T A P_m = (V_m U_m^{-1})^T A V_m U_m^{-1} \\
 P_m &= V_m U_m^{-1} \quad \quad \quad = U_m^{-T} (V_m^T A V_m) U_m^{-1} = U_m^{-T} T_m U_m^{-1} = \boxed{U_m^{-T} L_m} \quad \text{Diagonal}
 \end{aligned}$$

A symmetric

- Applicable only for Symmetric A matrices
- Computational steps are reduced using recursive relations for r and p

Conjugate gradient method is a faster method in terms of number of iterations or convergence rate it's the convergence rate is a function of root over of condition numbers. So, the convergence is faster and also the number of operations in each iteration step is smaller, because we can use something like a recursive relation for tri diagonal matrices here. A conjugacy of auxiliary vector P $p_i^T A p_j$ is equal to 0 if i is not equal to j is illustrated we will quickly see what is an auxiliary vector.

So, tri diagonal matrix T_m can be decomposed as a lower and upper triangular matrix and the $V_m U_m$ inverse this becomes an orthogonal this becomes the auxiliary matrix auxiliary vector matrix P_m . And all the columns of P are mutually A conjugate to each other that is $p_i^T A p_j$ is equal to 0 if i not is equal to j . So, we can get $P_m^T A P_m$ is equal to $U_m^{-T} L_m$ and we can show that this is a diagonal matrix and that is called the A conjugacy of the auxiliary vectors. This is however, this A conjugacy arises only when A is a symmetric matrix asymmetric only for asymmetric matrices.

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Conjugate Gradient Method

- A faster method in terms of number of iterations (convergence rate) and operations in each iteration step
- Uses A -conjugacy of auxiliary vector P ($p_i^T A p_j = 0$ if $i \neq j$):

$$\begin{aligned} T_m &= L_m U_m \\ x_m &= x_0 + V_m (L_m U_m)^{-1} y \quad \Rightarrow P_m^T A P_m = (V_m U_m^{-1})^T A V_m U_m^{-1} \\ P_m &= V_m U_m^{-1} \quad \quad \quad = U_m^{-T} (V_m^T A V_m) U_m^{-1} = U_m^{-T} T_m U_m^{-1} = \boxed{U_m^{-T} L_m} \quad \text{Diagonal} \end{aligned}$$

- Applicable only for **Symmetric A matrices** Can there be similar methods for non-symmetric matrices??
- Computational steps are reduced using recursive relations for r and p

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So, with all its advantages conjugate gradient is only applicable for symmetric matrices and the computational steps are reduced using recursive relations for r and p for residual as well as for the auxiliary vector we can use the recursive relations. So, it is only applicable for symmetric A matrices; and the question is that can there be similar methods for non-symmetric matrices also.

And for that we explore the other variants of Krylov subspace method in which we can some way handle the asymmetric method. And one idea is that that if we have A in the space we use the Krylov space of A for K m , can you use the Krylov space of A transpose as a L m so, that A plus A transpose is a symmetric matrix. So, that the asymmetry of A is some way taken care of by the asymmetry of A transpose A plus A transpose a symmetric matrix. So, can we can we pose it like the a problem like this and we will we will explore it.

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Lanczos Biorthogonalization

Lanczos method for nonsymmetric matrices builds a pair of biorthogonal bases using the two Krylov subspaces:

$$K_m(A, v_1) = \text{span}\{v_1, Av_1, A^2v_1, \dots, A^{m-1}v_1\}$$

And

$$K_m(A^T, w_1) = \text{span}\{w_1, A^T w_1, (A^T)^2 w_1, \dots, (A^T)^{m-1} w_1\}$$

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Lanczos by orthogonalization exactly looks into the Krylov space to Krylov subspaces; one Krylov subspace of one is of A another is of A transpose. So, it builds a pair of biorthogonal basis using the two Krylov subspaces came $A v_1$ which is span of $v_1, A v_1, A^2 v_1, \dots, A^{m-1} v_1$ and K_m of A transpose w_1 which is span of $w_1, A^T w_1, (A^T)^2 w_1, \dots, (A^T)^{m-1} w_1$.

Now, if by Lanczos biorthogonalization method, we get bases of k and K_m a v_1 and K_m A transpose v_1 Krylov subspace of A and A transpose and these bases are biorthogonal to each other. In that that is a sense that we take one basis of this v_i and we take one bases of this w_j ; $v_i^T w_j$ will be 0 if $i \neq j$ or $w_j^T v_i$ will be 0 if i is not equal to j . So, one v_i is conjugate to all other w_s except that particular element of w_i all vector of w all other w_s ; there is a biconjugacy between v and w .

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Algorithm: Lanczos Biorthogonalization Procedure

1. Choose two vectors v_1, w_1 such that $(v_1, w_1) = 1$.
2. Set $\beta_1 = \delta_1 \equiv 0, w_0 = v_0 \equiv 0$
3. For $j = 1, 2, \dots, m$ Do:
4. $\alpha_j = (Av_j, w_j)$
5. $\hat{v}_{j+1} = Av_j - \alpha_j v_j - \beta_j v_{j-1}$
6. $\hat{w}_{j+1} = A^T w_j - \alpha_j w_j - \delta_j w_{j-1}$
7. $\delta_{j+1} = |(\hat{v}_{j+1}, \hat{w}_{j+1})|^{1/2}$. If $\delta_{j+1} = 0$ Stop
8. $\beta_{j+1} = (\hat{v}_{j+1}, \hat{w}_{j+1}) / \delta_{j+1}$
9. $w_{j+1} = \hat{w}_{j+1} / \beta_{j+1}$
10. $v_{j+1} = \hat{v}_{j+1} / \delta_{j+1}$
11. EndDo

v_j and w_j are the biorthonormal basis for krylov subspaces of A and A^T respectively

Handwritten notes:
 $v_i^T w_j = 1 \quad i=j$
 $= 0 \quad i \neq j$

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And there is an algorithm for that by which we can get it which is called Lanczos biorthogonalization algorithm it initially starts with any guess vectors v_1 and w_1 with their dot product being 1. So, their orthonormal in that sense the dot product is normal. And sets beta one is equal to delta 1 is equal to 0 and w_0 is equal to v_0 is equal to 0. And then uses something like a recursive relationship for v_{j+1} and w_{j+1} and gets the delta $j+1$ which is a dot product between v_{j+1} and w_{j+1} root of that, and then divides then beta $j+1$ which is this dot product divided by delta $j+1$ and divide w by that beta and v by delta and if delta is equal to 0 this $j+1$ stop. So, if the dot product between v and w is 0 this algorithm stops.

So, it starts with one take one particular v_1, w_1 and for next v is obtained as v is equal to Av minus alpha into v alpha is $A v \cdot w$ minus beta into v_{j-1} . So, from v the a certain amount of v_{j-1} of v is v_j is subtracted as well as alpha $j v_j$ is subtracted where $v \alpha_j$ comes by dot product of v_j and w_j and the similarly w_{j+1} comes. So, through this method it is seen that v_j and w_j are biorthogonal to each other which is v_j and w_j are bi orthonormals bases of Krylov subspace A and A^T . So, $v_j^T w_j = 1$ and $v_j^T w_i = 0$ if $i \neq j$.

That is the property of biorthogonality or we can write $v_i^T w_j$ is equal to one if i is equal to j is equal to 0 if i is not equal to j and that is hence ascertained by this particular algorithm.

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Algorithm: Lanczos Biorthogonalization Procedure

1. Choose two vectors v_1, w_1 such that $(v_1, w_1) = 1$.
2. Set $\beta_1 = \delta_1 \equiv 0, w_0 = v_0 \equiv 0$
3. For $j = 1, 2, \dots, m$ Do:
4. $\alpha_j = (Av_j, w_j)$
5. $\hat{v}_{j+1} = Av_j - \alpha_j v_j - \beta_j v_{j-1}$
6. $\hat{w}_{j+1} = A^T w_j - \alpha_j w_j - \delta_j w_{j-1}$
7. $\delta_{j+1} = |(\hat{v}_{j+1}, \hat{w}_{j+1})|^{1/2}$. If $\delta_{j+1} = 0$ Stop \leftarrow break down
8. $\beta_{j+1} = (\hat{v}_{j+1}, \hat{w}_{j+1}) / \delta_{j+1}$
9. $w_{j+1} = \hat{w}_{j+1} / \beta_{j+1}$
10. $v_{j+1} = \hat{v}_{j+1} / \delta_{j+1}$
11. EndDo

v_j and w_j are the biorthonormal basis for krylov subspaces of A and A^T respectively

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And the process breaks down once we get the product between v and w for the particular j to be 0 that means, there is no further independent basis of the Krylov subspaces which can be generated all the independent bases vector all the independent vectors have been found out or we have been we have calculated the entire bases of these two spaces A and A transpose. This took the Krylov spaces of A and A transpose.

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Lanczos Biorthogonalization- propositions

If the lanczos biorthogonalization algorithm (for a non-symmetric matrix, A) does not break down before the step m , then the vectors $v_i, i=1, \dots, m$ and $w_j, j=1, \dots, m$, form a biorthogonal system; i.e.,



$$v_i^T w_j = \begin{cases} 1 & \text{if } i=j \\ 0 & \text{otherwise} \end{cases} \quad 1 \leq i, j \leq m$$

Moreover, $v_i, i=1, \dots, m$ is a basis of $K_m(A, v_1)$ and $w_j, j=1, \dots, m$ is a basis of $K_m(A^T, w_1)$, and the following relations hold

$$AV_m = V_m T_m + \delta_{m+1} v_{m+1} e_m^T$$

$$A^T W_m = W_m T_m^T + \beta_{m+1} w_{m+1} e_m^T$$

with $W_m^T AV_m = T_m \leftarrow T_m$ is a tridiagonal matrix

Lanczos biorthogonalization follows this proposition if the Lanczos bi-orthogonalization algorithm for a non-symmetric matrix A does not break down before the step m ; that means, before the step m $V_m^T W_m$ is non-zero. Then for vectors $v_i, i=1, \dots, m$ and $w_j, j=1, \dots, m$ form a biorthonormal system, that is $v_i^T w_j = 1$ if $i=j$ and otherwise it is 0, and this is ascertained by the way the Lanczos algorithm has been devised.

So, it is the algorithm for finding out basis of A transpose v and A transpose w and Krylov subspace between A transpose and w and A and v and there are these Krylov subspaces can have any bases any m bases, but the basis our vectors are found using Lanczos by orthogonalization algorithm in a way that, $v_i^T w_j = 1$ or 0 if $i=j$ they are 1 otherwise there is 0 and v and w are biorthonormal bases of this two Krylov subspaces. This looks little abstract, but we will see that this is a great utility when you will try to define/derive an algorithm from here.

Moreover $v_i, i=1, \dots, m$ is a basis of $K_m(A, v_1)$ plus 1 Krylov subspace of A and v_1 and $w_j, j=1, \dots, m$ is the basis of $K_m(A^T, w_1)$ and the following relations hold that $AV_m = V_m T_m + \delta_{m+1} v_{m+1} e_m^T$ where δ_{m+1} is defined to be the first unique vector.

A transpose w_m is w_m^T m transpose β_m w_m plus 1 e_m transpose T_m is a tri diagonal matrix. So, its transpose is also a triangular matrix with that now the T_m is defined such that $W_m^T A V_m$ is equal to T_m . W_m is the matrix containing as its columns all the vectors which are bases of the Krylov subspace of A transpose all the w s.

Similarly, V_m is the matrix which contains all the bases of Krylov subspace of A all the v s. And $W_m^T A V_m$ is a tri diagonal matrix and this is also this is also a very evidently very easily a apparent because w and v are biorthonormal $w_m^T V_m$ must be a $W_m^T A V_m$ must be a tridiagonal matrix. So, T_m is a tri diagonal matrix.

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Two-sided Lanczos algorithm for linear systems

1. Compute $r_0 = b - Ax_0$ and $\beta = \|r_0\|_2$
2. Run m steps of the nonsymmetric Lanczos Algorithm
3. Start with $v_1 = r_0/\beta$ and any w_1 such that $(v_1, w_1) = 1$
4. Generate Lanczos vectors $v_i, i=1, \dots, m$ and $w_j, j=1, \dots, m$, and
5. The tridiagonal matrix $T_m = W_m^T A V_m$ using Lanczos biorthogonalization algorithm
6. Compute $y_m = T_m^{-1}(\beta e_1)$ and $x_m := x_0 + V_m y_m$

Handwritten notes:
 FOM: $y_m = H_m^{-1}(\beta e_1)$
 D-Lanczos: $T_m = V_m^T A V_m$
 D-Lanczos $y_m = T_m^{-1}(\beta e_1)$
 $H_m = T_m$

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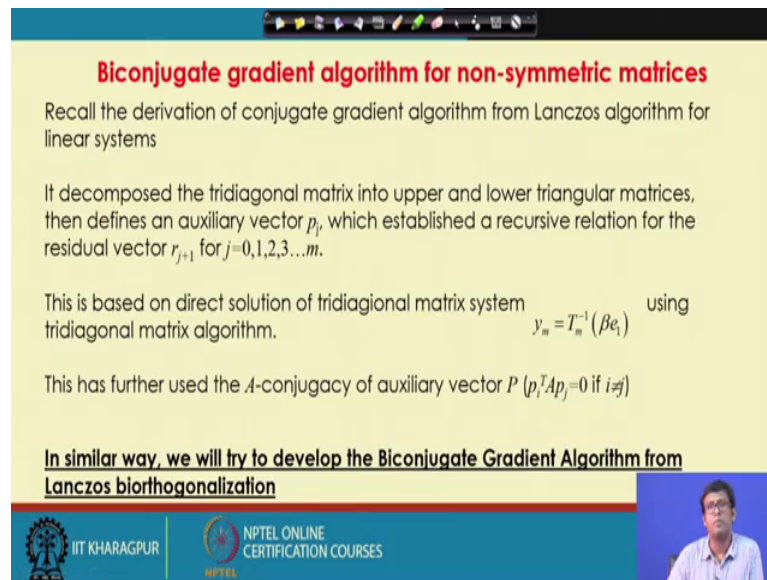
Two sided Lanczos algorithm is found is a device for linear systems or for solving $Ax = b$ compute $r_0 = b - Ax_0$ and β is equal to $\|r_0\|_2$ norm of r_0 . Run m steps of non symmetric Lanczos algorithm start with $v_1 = r_0/\beta$ and any w_1 such that $(v_1, w_1) = 1$. So, that $v_1 \cdot w_1 = 1$. Generate Lanczos vectors using Lanczos algorithm v_i and w_i find the tri diagonal matrix T_m and then compute y_m is equal to $T_m^{-1}(\beta e_1)$. So, T_m is the again the same as the Heisenberg matrix upper part of the Heisenberg matrix which is a tri diagonal matrix we found out in full orthogonal method.

So, similarly you write T_m is equal to $W_m^T A V_m$ and compute y_m is equal to $T_m^{-1} \beta e_1$ and x_m is equal to $x_0 + V_m y_m$. Now if we can remember that in the relationship for full orthogonal method was Y_m is equal to $H_m^{-1} \beta e_1$ and this and in Lanczos say D-Lanczos source or Lanczos for linear systems will say that Y_m is equal to $T_m^{-1} \beta e_1$ because H_m and $T_m H_m$ is T_m a min Lanczos method. The Heisenberg upper part of the Heisenberg matrix is same as that as is a tri diagonal matrix here how was full orthogonal method coming? It was appearing due to the fact that our K_m and L_m are same.

Here we are getting a different tridiagonal matrix, but we are getting a similar relationship due to the fact and here T_m was if we look into the conjugate gradient method, T_m was defined in a different way. But, it was coming from the fact that L_m the Krylov subspace vectors or T_m was defined in FOM or in rather not I will write Lanczos, the Lanczos for symmetric matrix T_m was defined as $V_m^T A V_m$ and this is because V_m was orthogonal to the residual vectors. V_m is where they x is updated V_m is also the subspace V_m is also the bases of the L_m , V_m is also orthogonal to the residual vector.

Here instead we get a w_m^T . So, w_m^T must be orthogonal to the residual vector here; or what we can say or w_m must be orthogonal to residue vector here or what we should see say is that that the Krylov subspace of a m transpose r_0 will be the a L_m for Lanczos biorthogonalization methods or a variant of that. We will look it into a little more elaborative way.

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Biconjugate gradient algorithm for non-symmetric matrices

Recall the derivation of conjugate gradient algorithm from Lanczos algorithm for linear systems

It decomposed the tridiagonal matrix into upper and lower triangular matrices, then defines an auxiliary vector p_j , which established a recursive relation for the residual vector r_{j+1} for $j=0,1,2,3,\dots,m$.

This is based on direct solution of tridiagonal matrix system $y_m = T_m^{-1}(\beta e_1)$ using tridiagonal matrix algorithm.

This has further used the A -conjugacy of auxiliary vector P ($p_i^T A p_j = 0$ if $i \neq j$)

In similar way, we will try to develop the Biconjugate Gradient Algorithm from Lanczos biorthogonalization

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That recall biconjugate gradient algorithm for non symmetric matrices. Let us recall the derivation of conjugate gradient algorithm for Lanczos method. If it decomposed the tri diagonal matrix into upper and lower triangular matrices then defines an auxiliary vector p . So, tri diagonal matrix is decomposed in $L m u m$ or then auxiliary vector p is defined, and a recursive relation is established for the residue vector $r j$ plus 1.

This is based on the direct solution of tri diagonal matrix system $Y m$ is equal to $T m$ inverse $\beta e 1$ using something like a TDMA type of algorithm which is a direct algorithm. This is further used a conjugacy of orthogonal auxiliary vector p transpose $A p$ is equal to an identity vector if i is not is equal to j $p i$ transpose $a p j$ is equal to 0.

In similar way we will try to now we will now try to develop a bi conjugate gradient algorithm from Lanczos biorthogonal narration. The idea is that $y m$ is calculated in similar way and there is an e conjugacy now there is a conjugacy here there were conjugacy between the bases vectors of $v v i$ transpose $v j$ is equal to 0 if i is not equal to j . Here there is a conjugation between w and v , we will utilize these facts.

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Biconjugate gradient algorithm for non-symmetric matrices

The Biconjugate Gradient (BCG) algorithm is a projection process onto

$$K_m = \text{span}\{v_1, Av_1, A^2v_1, \dots, A^{m-1}v_1\}$$


orthogonally to:



$$L_m = \text{span}\{w_1, A^T w_1, (A^T)^2 w_1, \dots, (A^T)^{m-1} w_1\}$$

Starting with as usual $v_1 = r_0 / \|r_0\|_2$ with any w_1 such that $(v_1, w_1) \neq 0$. Usually it is chosen as v_1 .

This method is equivalent to solving a dual system $A^T x^* = b^*$ to solve with A ; in that case w_1 is obtained by scaling the initial residual $b^* - A^T x_0^*$.

However, the dual system is not solved explicitly!



The biconjugated gradient algorithm is a projection process on to the Krylov subspace K_m which is a span of $v_1, Av_1, A^2v_1, \dots, A^{m-1}v_1$ and it starts with v_1 is the initial residual vector $b - r_0$ a $n \times 1$ zeros unit vector on that. Orthogonal to thus (Refer Time: 20:59) the residue vector must be orthogonal to L_m which is span of $w_1, A^T w_1, (A^T)^2 w_1, \dots, (A^T)^{m-1} w_1$.

So, now L_m is span of Krylov subspace of A^T and that is a way we are taking any general matrix which is non symmetric matrix, and trying to build an algorithm which was earlier built for a symmetric matrix. For conjugate gradient L_m was Krylov subspace of A only, but for biconjugate gradient we will see for non symmetric matrix. So, similar algorithm will come keeping in mind that L_m is now not the Krylov subspace of A rather it is the Krylov subspace of A^T .

So, we will start with as usual v_1 is equal to $r_0 / \|r_0\|_2$, and that is the unit vector along the first residue direction and take a w_1 such that $(v_1, w_1) \neq 0$. As their orthonormal usually take w_1 is equal to v_1 . The method is equivalent to solving a dual system $A^T x^* = b^*$ along with $Ax = b$. And to solve with A^T in that case w_1 is obtained by scaling the initial residual $b^* - A^T x_0^*$.

Instead of if we actually have to solve the dual system, we are not solving the dual system, but in the back of this processes this dual system is also being solved. Krylov

subspace of a is my $K \times m$ space here Krylov sub space of A^T is the $L \times m$ here. If I take another equation $A^T x = b^*$ $A^T x^*$ is equal to b star the element $K \times m$ is just reversed.

So, if I can solve one equation it is identical of solving the other equation also, which is equation for the transpose of a matrix. If we as we can solve that both the equations together, but you we, but as the given problem is $Ax = b$ for us we do not solve the transpose equation, which is type of solved in the back of the algorithm.

But if we actually have to solve the transpose equation also, able to solve a dual problem $A^T x = b$ and $Ax = b^*$ $A^T x^* = b$ star. We have to start with $w = b^* - A^T x_0$ zeros. However, as we are not solving here, it is well we can take any w_1 to start with usually it is chosen as v_1 . However the dual system is not solved explicitly.

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Biconjugate Gradient Method- derivation

Start with the LU decomposition of the tridiagonal matrix into lower triangular and upper triangular matrix:


$$T_m = L_m U_m$$



Define the auxiliary matrix, P_m as: $P_m = V_m U_m^{-1}$

The solution is expressed as:

$$\begin{aligned} x_m &= x_0 + V_m T_m^{-1} (\beta e_1) \\ &= x_0 + V_m U_m^{-1} L_m^{-1} (\beta e_1) \\ &= x_0 + P_m L_m^{-1} (\beta e_1) \end{aligned}$$

b



So, for the derivation we start with the LU decomposition of the tri diagonal matrix into lower triangular and upper triangular matrix $T_m = L_m U_m$. We define an auxiliary matrix P_m as $P_m = V_m U_m^{-1}$. It is very same as it we have defined it in the conjugate gradient case. The solution is expressed as $x_m = x_0 + V_m T_m^{-1} \beta e_1$; now $T_m^{-1} = U_m^{-1} L_m^{-1}$ and $V_m U_m^{-1}$ inverse is P_m . So, $x_0 + P_m L_m^{-1} \beta e_1$ so, this is inversion of a lower triangular matrix multiplied by a auxiliary vector matrix P .

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Biconjugate Gradient Method- derivation (contd)

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

Note: this update is similar as CG.
 Like CG algorithm, the vectors r_j and r_j^* are in same direction of v_{j+1} and w_{j+1} respectively
 As v_i and w_i are biorthogonal ($i \leq m$) hence r_j and r_j^* also form a biorthogonal system

Similarly, define the matrix $P_m^* = W_m U_m^{-1}$

Now: $(P_m^*)^T A P_m = L_m^{-1} W_m A V_m U_m^{-1} = L_m^{-1} T_m U_m^{-1} = I$

$P_m = V_m U_m^{-1}$
 $T_m = L_m U_m$

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x_m is equal to x_0 plus $P_m L_m^{-1} \beta e_1$ which is the this one. This update is very similar as conjugate gradient update. Like conjugate gradient algorithm the vector r_j and r_j^* are in same direction of v_{j+1} and w_{j+1} right. Because the residue vector is orthogonal to w and what is orthogonal to w ? Is v . So, residue vector is along v_{j+1} r_j is orthogonal to w_{j+1} . So, it should be along v_{j+1} r_j^* similarly will be along w_{j+1} .

So, as v_i and w_i are biorthogonal for $i \leq m$, hence r_j and r_j^* should also form a biorthogonal system. Similarly we can define a matrix P^* which is $W_m U_m^{-1}$. P was defined as P_m was defined as $V_m U_m^{-1}$. So, the star matrix that is the matrix for the transpose equation part P_m^* is $W_m U_m^{-1}$. And very interestingly we can show that $P_m^* A P_m = L_m^{-1} W_m A V_m U_m^{-1} = L_m^{-1} T_m U_m^{-1} = I$. $W_m A V_m W_m A V_m$ this is nothing, but the tri diagonal matrix T_m .

So, $L_m^{-1} T_m U_m^{-1}$ of a lower triangular matrix with a tri diagonal matrix and inverse of the upper triangular matrix and now we can we all we have also seen that T_m is equal to $L_m U_m$ T_m is decomposed as $L_m U_m$. So, you multiply this we will get the identity matrix. So, P_m^* and P_m the auxiliary vector matrix for A and for A^T are a conjugate. Earlier for conjugate gradient we have seen that P_m is itself a conjugate, but here we are getting P_m^* and P_m there a conjugate.

So, we got biconjugacy a (Refer Time: 26:41) a by conjugacy biconjugacy of r_j and r_j^* star residue vectors and we got by a conjugacy of auxiliary filters. Now the problem is exactly same as conjugate gradient method we can use similar type of iterate recursive relations and form the algorithm.

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Biconjugate Gradient Method- derivation (contd)

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

Note: this update is similar as CG.
 Like CG algorithm, the vectors r_j and r_j^* are in same direction of v_{j+1} and w_{j+1} respectively
 As v_i and w_i are biorthogonal ($i \leq m$) hence r_j and r_j^* also form a biorthogonal system

Similarly, define the matrix $P_m^* = W_m U_m^{-1}$

Now: $(P_m^*)^T A P_m = L_m^{-1} W_m A V_m U_m^{-1} = L_m^{-1} T U_m^{-1} = I$

So, the Auxiliary vectors are biorthogonal
 A CG like algorithm can hence be designed

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So, auxiliary vectors are biorthogonal as CG like algorithm hence can be designed.

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Biconjugate Gradient (BCG) Algorithm

ALGORITHM 7.3: Biconjugate Gradient (BCG)

1. Compute $r_0 := b - Ax_0$. Choose r_0^* such that $(r_0, r_0^*) \neq 0$.
2. Set $p_0 := r_0, p_0^* := r_0^*$
3. For $j = 0, 1, \dots$, until convergence Do:
4. $\alpha_j := (r_j, r_j^*) / (Ap_j, p_j^*)$
5. $x_{j+1} := x_j + \alpha_j p_j$
6. $r_{j+1} := r_j - \alpha_j Ap_j$
7. $r_{j+1}^* := r_j^* - \alpha_j A^T p_j^*$
8. $\beta_j := (r_{j+1}, r_{j+1}^*) / (r_j, r_j^*)$
9. $p_{j+1} := r_{j+1} + \beta_j p_j$
10. $p_{j+1}^* := r_{j+1}^* + \beta_j p_j^*$
11. EndDo

This algorithm works for any non-singular matrix A

If a dual system with A^T is being solved, then in line 1 r_0^* should be defined as $r_0^* = b^* - A^T x_0^*$ and the update $x_{j+1}^* := x_j^* + \alpha_j p_j^*$ to the dual approximate solution must be inserted after line 5. The vectors produced by this algorithm satisfy a few biorthogonality properties stated in the following proposition.

$(r_j, r_i^*) = 0, \text{ for } i \neq j,$
 $(Ap_j, p_i^*) = 0, \text{ for } i \neq j.$

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And we get the biconjugate algorithm biconjugate gradient algorithm the initial residual vector r_0 is equal to b minus x_0 we choose r_0^* such that the dot product for the

initial vector r_0 and r_0^* is non zero. Then p_0 is equal to r_0 p_0^* is equal to r_0^* zero. So, r_0 and r_0^* are non 0 dot product, but r_0 and r_1^* will be 0 that that is the idea by conjugacy. So, j is equal to 0 or two convergence find α_j which is $r_j^T A p_j$ start by $A p_j^T p_j^*$ these are biorthogonal and these are biconjugate update x_{j+1} as $x_j + \alpha_j p_j$ r_{j+1} as $r_j - \alpha_j A p_j$ r_{j+1}^* is equal to $r_j^* - \alpha_j A^T p_j^*$ get β_j and get p_{j+1} using β_j and $p_j^* + 1$ p_{j+1}^* using β_j and r_{j+1} .

So, similar recursive relations as conjugate gradient, but now it is for both r^* and r and p^* . But for x we have only finding out x_{j+1} because we not interested in finding the solution of the transpose equation you are not finding out x_{j+1} ; and after certain iterations it should converge and we get the converse solution.

The algorithm works for any nonsingular matrix A and the convergence etcetera can be shown as same as the conjugate gradient type of algorithm which; that means, these are first converging algorithms. If the dual system A^T is being solved, then in the line 1 r_0^* should be defined as $r_0^* = b^* - A^T x^*$ and then x_{j+1}^* has to be updated from x_j^* for dual approximate solution after the line 5.

The vectors produced by this algorithm satisfy about five orthogonality properties like $r_j^T r_i^* = 0$ if $i \neq j$ $A p_j^T p_i^* = 0$ if $i \neq j$, these are the base of those are proposition based on which this if this method is developed actually. The recursive relations come and then using this we can update the vectors these vectors the updated vectors also must satisfy this particular properties.

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Biconjugate Gradient (BCG) Algorithm – Computational Issues

ALGORITHM 7.3: Biconjugate Gradient (BCG)

1. Compute $r_0 := b - Ax_0$. Choose r_0^* such that $(r_0, r_0^*) \neq 0$.
2. Set, $p_0 := r_0, p_0^* := r_0^*$
3. For $j = 0, 1, \dots$, until convergence Do:
4. $\alpha_j := (r_j, r_j^*) / (Ap_j, p_j^*)$
5. $x_{j+1} := x_j + \alpha_j p_j$
6. $r_{j+1} := r_j - \alpha_j Ap_j$
7. $r_{j+1}^* := r_j^* - \alpha_j A^T p_j^*$
8. $\beta_j := (r_{j+1}, r_{j+1}^*) / (r_j, r_j^*)$
9. $p_{j+1} := r_{j+1} + \beta_j p_j$
10. $p_{j+1}^* := r_{j+1}^* + \beta_j p_j^*$
11. EndDo

Matrix vector multiplication

Matrix transpose vector multiplication

Now, there is a particular issue with biconjugate gradient algorithm, which we will discuss before we finish this lecture and we look into how to solve this issue in the subsequent lectures. That is if we look into the BiCG algorithm, that there are three matrix vector multiplication. If the first one is before the initial step, but during the iterations also there are two matrix vector multiplication Ax_0 is the first one, Ap_j is one multiplication and then $A^T p_j$ is again used another matrix vector multiplication. On top of that there is one more matrix vector multiplication with $A^T p_j$ and transpose multiplications are difficult in terms of communicating with the memory and the processor.

So, number of at least $A^T p_j$ a p_j three matrix vector multiplication then p_j is to be multiplied with p_j^* , the number of calculations operational steps are usually more and multiplication with the auxiliary vector and α is also more and we have to also multiply with the transpose vector. And for that the as the number of operational steps are more the in during each iteration, the number of the amount of round off error is also more. And due to this round of error in the convergence of this BiCG or biconjugate gradient method we see there are irregularity.

It does not converge in a smooth way or monotonically there are fluctuations during convergence and sometimes these fluctuations can be large enough. So, that the convergence is disturbed for certain case of the problems they therefore, small

perturbations we can see lot of change in the result due to this round off error related fluctuations.

So, to look into more stabilized versions of BiCG method, where this many matrix vector multiplications can be avoided and BiCG stabilized method is one of that method which we will discuss now in that the subsequent classes. And there are few other methods using some polynomial formulation of the auxiliary vector and residual vector polynomial expansion type of formulation of the auxiliary vector we can minimize the matrix vector products, and can get good recursive relations and much simple operations for in during each iteration and better algorithms can be devised.

In the next classes we will look into the much developed algorithms. Keeping in mind that BiCG is the basic algorithm for any non symmetric matrix, which will be now developed using certain polynomial expansions to BiCG stabilized type of algorithms. We will look into it in the next lesson.

Thank you.