

Iterative methods for solving  $Ax = b$  where  $A$  is non-symmetric.

GMRES (Generalized Minimum Residual) method

The  $k$ th iteration of GMRES is the solution  $x_k$   
to the least squares problem

$$\min_{x \in x_0 + K_k} \|b - Ax\|_2^2 \Rightarrow \nabla \|b - Ax\|_2^2 = 2A^T (Ax - b) = 0$$

Step2 in CG  $\Rightarrow \underbrace{(b - Ax_k)^T}_{r_k} \underset{\in K_k}{A} J = 0 \Rightarrow r_k$  is no longer  $\perp$  to  $K_k$

Step3 in CG  $\overset{x_{k+1}=x_k+\alpha_{k+1}p_{k+1}}{\Rightarrow} \alpha_k (JAp_{k+1}) = J(r_k - r_{k+1}) \neq 0$   
 $\Rightarrow$  no A-orthogonality property for  $p_{k+1}$

Consider  $y \in K_k$  and  $K_k = \text{span}\{v_1 \cdots v_k\} = V_k$

One must have  $x_k = x_0 + V_k \cdot y$  for some  $y$  (coefficient vector)

The minimization problem can be rewritten as following:

$$\begin{aligned} \min_{x \in x_0 + K_k} \|b - Ax\|_2 &\equiv \min_{y \in R^k} \|b - A(x_0 + V_k y)\| \\ &\equiv \min_{y \in R^k} \|r_0 - AV_k \cdot y\| \end{aligned}$$

This is a standard least Square problem.

One can solve it by  $QR$  or the Gram-Schmidt orthogonalization.

# Gram-Schmidt orthogonalization procedure:

(Arnoldi process)  $V_K = \text{span}\{v_1, v_2, \dots, v_k\}$

1. Define  $r_0 = b - Ax_0$ ,  $v_1 = \frac{r_0}{\|r_0\|_2}$

2. For  $i = 1 \sim k - 1$ , (restart with new  $r_0 = b - Ax_k$  when needed)

$$v_{i+1} = \frac{Av_i - \sum_{j=1}^i \overbrace{\left[ (Av_i)^T v_j \right]}^{h_{j,i}} \cdot v_j}{\underbrace{\left\| Av_i - \sum_{j=1}^i \left[ (Av_i)^T v_j \right] \cdot v_j \right\|_2}_{h_{i+1,i}}}$$

(the algorithm breaks down when  $h_{i+1,i} = 0$ )  
 $\Rightarrow$  restart is needed)

$$\Rightarrow A_{n \times n} V_{n \times k-1} = V_{n \times k} \cdot H_{k \times k-1}; H = \begin{bmatrix} * & & & h_{1,i} & * & * \\ * & \ddots & & \vdots & * & * \\ & \ddots & \ddots & \vdots & & * \\ & & \ddots & \vdots & & \vdots \\ & & & \vdots & & \vdots \\ & & & h_{i+1,i} & * & * \\ & & & & * & * \\ & & & & 0 & * \end{bmatrix}_{k \times k-1}$$

$\leftarrow$  upper hessenberg matrix.

Now suppose we are at  $k$ th step. We have

$$\begin{aligned}\|r_k\|_2 &= \|b - Ax_k\|_2 = \|A(x - x_0 + x_0 - x_k)\|_2 \\ &= \left\| r_0 - A \underbrace{(x_k - x_0)}_{=V_k y} \right\|_2\end{aligned}$$

Since  $r_0 = \beta v_1 e_1$ ,  $\beta = \|r_0\|_2$ , and  $V_i$  is an orthonormal basis

$$\begin{aligned}\|r_k\|_2 &= \|V_k^T (r_0 - AV_k y)\|_2 \\ &= \left\| \beta e_1 - V_k^T \cdot V_{k+1} H_{k+1 \times k} y \right\|_2 = \left\| \beta e_1 - \begin{matrix} [H]_{[1:k] \times [1:k]} \\ \parallel \\ \bar{H}_k \end{matrix} y \right\|_2\end{aligned}$$

Our goal now is to find  $y$  that satisfies

$$\min_{x \in x_0 + K_k} \|b - Ax\| \equiv \min_{y \in R^k} \|r_0 - AV_k y\| \equiv \min_{y \in R^k} \|\beta e_1 - \bar{H}_k y\|_2 \quad -(+++)$$

Now the Gmres algorithm can be read as following

ALGORITHM 3.4.2. *gmresa*( $x, b, A, \epsilon, kmax, \rho$ )

1.  $r = b - Ax$ ,  $v_1 = r / \|r\|_2$ ,  $\rho = \|r\|_2$ ,  $\beta = \rho$ ,  $k = 0$
2. While  $\rho > \epsilon \|b\|_2$  and  $k < kmax$  do
  - (a)  $k = k + 1$
  - (b) for  $j = 1, \dots, k$   
 $h_{jk} = (Av_k)^T v_j$
  - (c)  $v_{k+1} = Av_k - \sum_{j=1}^k h_{jk} v_j$
  - (d)  $h_{k+1,k} = \|v_{k+1}\|_2$
  - (e)  $v_{k+1} = v_{k+1} / \|v_{k+1}\|_2$
  - (f)  $e_1 = (1, 0, \dots, 0)^T \in R^{k+1}$   
 Minimize  $\|\beta e_1 - H_k y^k\|_{R^{k+1}}$  over  $R^k$  to obtain  $y^k$ .
  - (g)  $\rho = \|\beta e_1 - H_k y^k\|_{R^{k+1}}$ .
3.  $x_k = x_0 + V_k y^k$ .

The convergence analysis is similar to the convergence analysis for CG. We leave the analysis to readers.

Theorem 1: Let  $A = VAV^{-1}$  be a nonsingular diagonalizable matrix and  $x_k$  be the  $k$ th GMRES iteration. Then for all  $p_k \in P_k$

$$\frac{\|r_k\|_2}{\|r_r\|_2} \leq K_2(v) \max_{z \in \sigma(A)} |p_k(z)|$$

Remark 1:

A is normal  $\Rightarrow$  V is orthonormal

$\Rightarrow$  the condition number  $K_2(v) = 1$

If A is not normal, it is not clear how to estimate  $K_2(v)$

Remark2: Proconditioning is generally needed to accerlatethe convergence rate. Moreover, due to possible lose of orthogonality in the Gram-Schmidt process.

$\left( \begin{array}{l} \text{no guarantee that } Av_i \notin \text{span}\{v_1 \cdots v_n\} \Rightarrow \text{breakdown} \\ \Rightarrow \text{Arnoldi process can't generate } v_{i+1} \end{array} \right)$

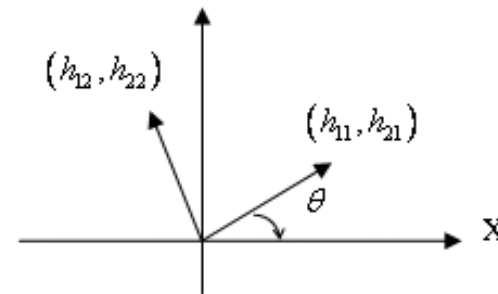
"Restart" is also generally needed in GMRES.

Remark3: The storage cost for the upper hessnberg martix H grows linearly with the number of iteration. Restarting strategy also can reduce the storage cost of H matrix.

By applying rotation matrices to  $H$ , one can easily solve the above minimization problem.

let  $H = \left[ \begin{array}{cc|cc} h_{1,1} & h_{1,2} & & \\ h_{2,1} & h_{2,2} & & \\ \hline & h_{3,2} & h_{3,3} & \\ & & \dots & \\ & & & h_{k-1,k-1} \\ & & & h_{k,k-1} & h_{k,k} \\ & & & & h_{k+1,k} \end{array} \right]$

observation:



$$\cos \theta = \frac{h_{11}}{\sqrt{h_{11}^2 + h_{21}^2}}$$

$$\sin \theta = \frac{h_{21}}{\sqrt{h_{11}^2 + h_{21}^2}}$$

$$\begin{bmatrix} \cos \theta & \sin \theta \\ -\sin \theta & \cos \theta \end{bmatrix} \begin{bmatrix} h_{11} \\ h_{21} \end{bmatrix} = \begin{bmatrix} \sqrt{h_{11}^2 + h_{21}^2} \\ 0 \end{bmatrix}$$

From the block  $B_1$ , we can form  $R_1 = \left[ \begin{array}{cc|c} c_1 & s_1 & \\ -s_1 & c_1 & \\ \hline & & I \end{array} \right]$ ,  $\|R_1\| = 1$ . We have  $R_1 H = \begin{bmatrix} * & \widetilde{h}_{12} & & \\ 0 & \widetilde{h}_{22} & \widetilde{h}_{23} & \dots & \dots \\ 0 & \widetilde{h}_{32} & h_{33} & \dots & \dots \\ \vdots & & & \ddots & \\ 0 & & & & \ddots \end{bmatrix}$



From the block  $B_2$ , we can obtain  $R_2 = \begin{bmatrix} & & \\ & c_2 & s_2 \\ & -s_2 & c_2 \\ & & & I \end{bmatrix}$ ,  $\|R_2\| = 1$

By repeatedly applying the rotation matrices, we have

$$\beta e_1 - H_k y = \underbrace{\overline{R}_k^*}_{\text{rotation matrices}} \cdot \underbrace{\left( \beta \overline{R}_k e_1 - \begin{bmatrix} \cdot & \cdot & * \\ 0 & \cdot & \cdot \end{bmatrix} y \right)}_{\text{upper-triangular linear system}}, \text{ here } \overline{R}_k = R_k R_{k-1} \cdots R_2 R_1$$

Now one can apply the backward substitution to solve the upper triangular linear system.

Moreover,

$$\|\beta e_1 - H_k y\| = \left| \beta \underbrace{\left( \overline{R}_k e_1 \right)_k}_{\text{(k+1)th component of } \overline{R}_k e_1} \right|,$$

the quantity can be used in checking the stopping tolerance.

# The GMRES algorithm can be rewritten as following

ALGORITHM 3.5.1.  $\text{gmres}(x, b, A, \epsilon, kmax, \rho)$

- $r = b - Ax$ ,  $v_1 = r/\|r\|_2$ ,  $\rho = \|r\|_2$ ,  $\beta = \rho$ ,  
 $k = 0$ ;  $g = \rho(1, 0, \dots, 0)^T \in R^{kmax+1}$
- While  $\rho > \epsilon\|b\|_2$  and  $k < kmax$  do
  - $k = k + 1$
  - $v_{k+1} = Av_k$   
for  $j = 1, \dots, k$ 
    - $h_{jk} = v_{k+1}^T v_j$
    - $v_{k+1} = v_{k+1} - h_{jk}v_j$
  - $h_{k+1,k} = \|v_{k+1}\|_2$
  - Test for loss of orthogonality and reorthogonalize if necessary.
  - $v_{k+1} = v_{k+1}/\|v_{k+1}\|_2$
  - If  $k > 1$  apply  $Q_{k-1}$  to the  $k$ th column of  $H$ .
    - $\nu = \sqrt{h_{k,k}^2 + h_{k+1,k}^2}$
    - $c_k = h_{k,k}/\nu$ ,  $s_k = -h_{k+1,k}/\nu$   
 $h_{k,k} = c_k h_{k,k} - s_k h_{k+1,k}$ ,  $h_{k+1,k} = 0$
    - $g = G_k(c_k, s_k)g$ .
  - $\rho = |(g)_{k+1}|$ .
- Set  $r_{i,j} = h_{i,j}$  for  $1 \leq i, j \leq k$ .  
Set  $(w)_i = (g)_i$  for  $1 \leq i \leq k$ .  
Solve the upper triangular system  $Ry^k = w$ .
- $x_k = x_0 + V_k y^k$ .