

## Lanczos = symmetric Arnoldi

If  $A$  is symmetric ( $A = A^T$ ), then  $H_n$  is symmetric for each  $n$  and hence it is **tridiagonal**.

```
A = bucky(); b = randn(size(A, 1), 1);
[Q, H] = arnoldi(bucky, b, 40);
spy(H);
```

Indeed,  $H_{ji} = q_j^T A q_i = (q_j^T A q_i)^T = q_i^T A q_j = H_{ij}$ .

So we can shorten the orthogonalization loop:

```
w = A * Q(:, j);
for i = j-1:j % only 2 vectors to check
    %and actually we already know H(j-1,j)=H(j,j-1)
    H(i,j) = Q(:, i)' * w;
    w = w - H(i,j) * Q(:, i);
end
```

Reduces the cost to  $n$  matrix products +  $O(mn)$

**Lanczos iteration**, used to compute eigenvalues/vectors.

$$Aq_j = \cancel{\beta_1 q_1} + \cancel{\beta_2 q_2} + \dots + \cancel{\beta_{j-2} q_{j-2}} + \beta_{j-1} q_{j-1} + \beta_j q_j + \beta_{j+1} q_{j+1}$$

"short recurrence"

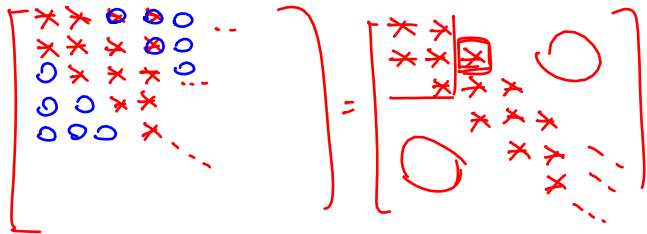
$H_{ij}$  contains  $\beta_i = q_i^T A q_j$

If  $A = A^T$ , then  $H_{ij} = q_i^T A q_j =$

$$= (q_i^T A q_j)^T = q_j^T A^T q_i = q_j^T A q_i = H_{ji}$$

because it's  $1 \times 1$

$$A = A^T$$





# Conjugate gradient | AF, optimization

Suppose  $A = A^T$  is **positive definite**. Then, we can find the solution to  $Ax = b$  by minimizing the (strictly convex) function

$$f(x) = \frac{1}{2} x^T A x - b^T x.$$

Surprisingly, **conjugate gradient** on this problem can be interpreted as a Krylov subspace method.

Three ingredients: current iterate  $x_k$ , residual

$$r_k = b - Ax_k = -\nabla f(x_k), \text{ and search direction } d_k.$$

## CG iteration

$$x_0 = 0, r_0 = d_0 = b;$$

**for**  $k = 1:n$  **do**

$$\alpha_k = (r_{k-1}^T r_{k-1}) / (d_{k-1}^T A d_{k-1});$$

$$x_k = x_{k-1} + \alpha_k d_{k-1};$$

$$r_k = r_{k-1} - \alpha_k A d_{k-1};$$

$$\beta_k = (r_k^T r_k) / (r_{k-1}^T r_{k-1});$$

$$d_k = \underline{r_k} + \underline{\beta_k d_{k-1}};$$

**end**

solution update  
residual/gradient update

// check it!

// A-orthogonalize search dir

$$d_{k-1}^T A d_k = 0$$

1)

$$X_k = X_{k-1} + \alpha_k d_{k-1}$$

We can choose  $\alpha$  by imposing for instance

$$v^T (AX_k - b) = 0 \quad \text{for a certain vector } v,$$

or by minimizing  $f(X_{k-1} + \alpha d_{k-1})$

2) Once we update  $X_{k-1}$  to  $X_k = X_{k-1} + \alpha_k d_{k-1}$

we need to compute  $r_k = b - AX_k$

$$r_k = b - AX_k = b - A(X_{k-1} + \alpha_k d_{k-1}) =$$

$$= \underbrace{b - AX_{k-1}}_{r_{k-1}} - \alpha_k A d_{k-1}$$

CG: choose search direction  $d_k$  as

$$d_k = r_k - \beta_k d_{k-1}$$

Choose  $\beta_k$  so that the search direction  $d_k$

is "A-conjugate" to  $d_{k-1}$ , i.e.

$$d_{k-1}^T A d_k = 0$$

$$f(x) = \frac{1}{2} x^T A x - b^T x$$

$$\nabla f(x) = Ax - b$$

$\nabla^2 f = A$ , if we assume  $A$  psdef it's strictly convex

one global minimum in  $\mathbb{R}^n$ , where  $\nabla f(x) = Ax - b = 0$



## CG — Considerations

// **Storage:** 3 vectors:  $x_k, r_k, d_k$ . No need to keep whole 'history'.

**Cost:**  $n$  mat-vec products ( $Ad_{k-1}$ ) plus  $O(nm)$ .

Where are the Krylov subspaces? At each step  $k$ ,

$$\begin{aligned} \underline{K}_k(A, b) &= \text{span}(x_1, x_2, \dots, x_k) = \text{span}(d_0, d_1, \dots, d_{k-1}) \\ &= \text{span}(r_0, r_1, \dots, r_{k-1}). \end{aligned}$$

(We sketch what happens in the first steps.)

$$x_0 = 0 \quad d_0 = r_0 = b$$

$$\text{span}(d_0) = \text{span}(r_0) = b$$

$$x_1 = x_0 + \alpha d_0$$

$$\text{span}(x_1) = b$$

$$r_0 = b - Ax_0 \quad r_1 - r_0 = -A(x_1 - x_0) = -A\alpha d_0$$

$$r_1 = b - Ax_1$$

$$r_1 = r_0 - \alpha Ad_0$$

$$\in \text{span}(b, Ab)$$

$$d_1 = r_1$$



## CG — Orthogonality

Where are the orthogonal bases?

At each step  $r_k^T r_i = d_k^T A d_i = 0$  for all  $i < k$ .

I.e., the  $r_i$  are orthogonal, and the  $d_i$  are  $A$ -orthogonal.

**Proof** (sketch) Suppose (induction) it holds up to step  $k - 1$ .

From  $r_k = r_{k-1} - \alpha_k A d_{k-1}$  it follows that

$$r_i^T r_k = r_i^T r_{k-1} - \alpha_k r_i^T A d_{k-1}.$$

If  $i < k - 1$ , then  $r_i^T r_{k-1}$  is zero by induction and  $r_i \in \text{span}(d_0, d_1, \dots, d_i)$  is  $A$ -orthogonal to  $d_{k-1}$ .

If  $i = k - 1$ , then the RHS is zero provided  $\alpha_k = \frac{r_{k-1}^T r_{k-1}}{r_{k-1}^T A d_{k-1}}$ . The formula we gave was slightly different: at the denominator we had  $d_{k-1}^T A d_{k-1}$  instead. But since  $d_{k-1} = r_{k-1} + \beta_{k-1} d_{k-2}$ , the difference between the two denominators is  $\beta_{k-1} d_{k-2}^T A d_{k-1} = 0$  (by induction). The second part is similar.

## CG — orthogonal bases

The  $r_i$  are an orthogonal (but not orthonormal!) basis of  $K_k(A, b)$  at each step  $k$ . In particular,  $\frac{1}{\|r_i\|} r_i$  coincides (up to a sign) with the  $q_i$  obtained with Arnoldi.

Last ingredient: what do we solve at each step? Not

$\min_{z \in K_k(A, b)} \|Az - b\|$  like in GMRES, but

$$x_k \text{ is the solution of } \min_{z \in K_k(A, b)} (x - z)^T A(x - z).$$

where  $x$  is the true solution of  $Ax = b$ .

Indeed, let  $z = x_k + y$ , and note that  $A(x - x_k) = b - Ax_k = r_k$  is orthogonal to each  $y \in K_k(A, b)$ :

$$\begin{aligned}(x - z)^T A(x - z) &= (x - x_k)^T A(x - x_k) + (x - x_k)^T Ay + y^T A(x - x_k) \\ &\quad + y^T Ay \\ &= (x - x_k)^T A(x - x_k) + y^T Ay \geq (x - x_k)^T A(x - x_k).\end{aligned}$$

## Convergence of CG

First result: let  $\lambda_{\max}$ ,  $\lambda_{\min}$  be the maximum/minimum eigenvalue of  $A$ ; then, CG converges with rate

$$\|x - x_k\| \leq \left( \frac{\sqrt{\lambda_{\max}} - \sqrt{\lambda_{\min}}}{\sqrt{\lambda_{\max}} + \sqrt{\lambda_{\min}}} \right)^k \|x - x_0\|.$$

(we won't see a proof).

We can rewrite it in terms of a more familiar quantity: for a positive definite matrix, eigenvalues and singular values coincide, hence

$$\frac{\sqrt{\lambda_{\max}} - \sqrt{\lambda_{\min}}}{\sqrt{\lambda_{\max}} + \sqrt{\lambda_{\min}}} = \frac{\sqrt{\sigma_1} - \sqrt{\sigma_m}}{\sqrt{\sigma_1} + \sqrt{\sigma_m}} = \frac{\sqrt{\frac{\sigma_1}{\sigma_m}} - 1}{\sqrt{\frac{\sigma_1}{\sigma_m}} + 1} = \frac{\sqrt{\kappa(A)} - 1}{\sqrt{\kappa(A)} + 1}.$$

For large values of  $\kappa(A)$ , this is approximately  $1 - \frac{2}{\sqrt{\kappa(A)}}$ .

## Convergence of CG

Again, that is a worst-case result — usually, convergence is faster and depends on the **eigenvalues of  $A$**  (like in GMRES).

Let  $\|v\|_A := v^T A v$  “ $A$ -norm” of a vector; the quantity that is minimized at each step. It makes sense if  $A$  is posdef.

$x_n \in K_n(A, b) \iff x_n = p(A)b$  for a polynomial  $p$  of degree  $< n$ .

$$\begin{aligned} \min_{x_n \in K_n(A, b)} \|x - x_n\|_A &= \min_{p(x)} \|x - p(A)Ax\|_A \\ &= \min_{\substack{q(x)=1-xp(x) \\ \text{of degree } \leq n}} \|q(A)x\|_A. \end{aligned}$$

As for GMRES, if  $A$  has only  $n$  different eigenvalues, then this minimum reaches 0 after  $n$  steps. If the eigenvalues of  $A$  are ‘clustered’, one can construct polynomials such that  $q(\lambda)$  is small for each  $\lambda \implies$  fast convergence.