1 13th of December 2018 — F. Poloni

1.1 Lanczos algorithm

$\mathbf{\Omega}$

Do you recall?

In last lecture we saw how to factorize a matrix $A \in M(m, \mathbb{R})$ with Arnoldi (i.e. $AQ_n = Q_{n+1}\underline{H}_n = Q_n\underline{H}_n + h_{n+1,n}q_{n+1}e_1^T$).

If A is **symmetric**, something special happens: $\underline{H}_n = Q_m^T A Q_m$ is symmetric as well, so it is a **tridiagonal** matrix. This improves the complexity of the Arnoldi process, because many iterations of the for loop (j = 1 : n) are not needed anymore, we need only two iterations.

This symmetric variant of Arnoldi is called *Lanczos algorithm*, and such algorithm reduces the cost to n matrix products + O(mn).

Suppose $A = A^T$ is positive definite. Then, we can find the solution to Ax = b by minimizing the (strictly convex) function $f(x) = \frac{1}{2}x^T Ax - b^T x$.

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

The pseudocode of such algorithm can be found in Algorithm 1.1, where x_k is the current iterate, $r_k = b - Ax_k = -\nabla f(x_k)$ is the residual and d_k is the search direction.

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ALGORITHM 1.1 Pseudocode for the conjugate gradient method.
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1: procedure CG_ITERATION
            x_0 \leftarrow 0;
            r_0 \leftarrow b;
 3:
 4:
            d_0 \leftarrow b;
           for k = 1:n do
 5:
                 \alpha_k \leftarrow (r_{k-1}^T r_{k-1}) / (d_{k-1}^T A d_{k-1});
 6:
                  x_k \leftarrow x_{k-1} + \alpha_k d_{k-1};
 7:
                 r_k \leftarrow r_{k-1} - \alpha_k A d_{k-1};
 8:
                 \beta_k \leftarrow (r_k^T r_k) / (r_{k-1}^T r_{k-1});
 9:
                  d_k \leftarrow r_k + \beta_k d_{k-1};
10:
           end for
11:
12: end procedure
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Notice that the search direction (line 10) is modified adding a multiple of the previous search direction to the residual and β_k is chosen such that d_k and d_{k-1} are A-orthogonal (formally, $d_k^T A d_{k-1} = 0$).

Conversely, the next point is chosen in order to minimize the objective function $f(x_{k-1}\alpha_k d_{k-1})$. As far as the complexity is concerned, the space complexity is constant (three vectors) and the time complexity is O(mn). **Theorem 1.1.** $K_k(A, b) = span(x_1, x_2, \dots, x_k) = span(d_0, d_1, \dots, d_{k-1}) = span(r_0, r_1, \dots, r_{k-1}).$

Theorem 1.2. The residuals are orthogonal and the search directions are A-orthogonal. Formally, $r_i^T r_k = d_i^T A d_k = 0$, $\forall i < k$.

Proof. By induction: Let us assume we proved the thesis $r_j^T r_k = 0$ for $k-1, k-2, \ldots, 0$. Since $x_k = x_{k-1}\alpha_k d_{k-1}$, the residual $r_k = b - Ax_k = b - A(x_{k-1} + \alpha_k d_{k-1}) = b - Ax_{k-1} - \alpha_k Ad_{k-1} = r_{k-1} - \alpha_k Ad_{k-1}$.

Let us compute $r_j^T r_k = r_j^T (r_{k-1} - \alpha_k A d_{k-1})$.

- If i < k-1 then $r_j^T r_{k-1} r_j^T \alpha_k A d_{k-1} = 0$, because the first term is 0 from induction hypothesis and $r_j^T \alpha_k A d_{k-1} = 0$, because $r_j \in K_{k-1}(A, b) = span(d_0, d_1, \dots, d_{k-2})$.
- If i = k 1, $0 = r_{k-1}^T (r_{k-1} \alpha_k A d_{k-1}) = r_{k-1}^T r_{k-1} \alpha_k r_{k-1}^T A d_{k-1}$ holds if $\alpha_k = \frac{r_{k-1}^T r_{k-1}}{r_{k-1}^T A d_{k-1}}$. We are left with proving that $\alpha_k = \frac{r_{k-1}^T r_{k-1}}{r_{k-1}^T A d_{k-1}} = \alpha_k = \frac{r_{k-1}^T r_{k-1}}{d_{k-1}^T A d_{k-1}}$.

This is true, since $d_{k-1} = r_{k-1} + \beta_{k-1} d_{k-2}$, so $d_{k-1}^T A d_{k-1} = (r_{k-1} + \beta_{k-1} d_{k-2})^T A d_{k-1} = r_{k-1}^T A d_{k-1} + \beta_{k-1} d_{k-2} A d_{k-1}^T$ and the second part is equal to 0 by induction.

Notice that this base is not orthonormal, we need to rescale it to obtain an orthonormal one, moreover, $\frac{1}{\|r_i\|}r_i$ coincides (up to a sign) with the q_i obtained with Arnoldi.

We are left with writing the equation we need to solve at each iteration, namely we need to ensure that $r_k = b - Ax_k$ is orthogonal to all vectors of $K_k(A, b)$ which is equivalent to requiring $Q_k^T(b - Ax_k) = 0$ or, equivalently, $||b|| e_1 = H_k y_k$.

In figure Figure 1.1 we can see a comparison between Arnoldi algorithm and the conjugate gradient.

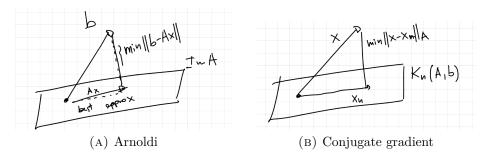


FIGURE 1.1: Traditional orthogonality (Arnoldi) leads to the minimization of the 2-norm, while in the conjugate gradient we impose A-orthogonality and we get a good approximation in several norms.

As far as convergence speed is concerned,

Theorem 1.3. x_k is the best approximation of the exact (and unknown) solution x to Ax = b in $K_k(A, b)$ in the A-norm, i.e. $x_k = arg \min_{z \in K_k(A, b)} (x - z)^T A(x - z)$

Theorem 1.4. Let λ_{max} , λ_{min} be the maximum/minimum eigenvalue of A; then, CG converges with rate:

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

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 $1 - \frac{2}{\sqrt{\kappa(A)}}$, while if $\kappa(A) \approx 1$ the convergence speed is very high.

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 λ then fast convergence is implied.