Iterative Solvers 3 - The Conjugate Gradient Method

Symmetric positive definite matrices

A very frequent type of matrices are symmetric positive definite matrices. Let $A \in \mathbb{R}^{n \times n}$ be a symmetric matrix (that is $A^T = A$). A is called symmetric positive definite if

$$x^TAx > 0, \forall x \neq 0.$$

This is equivalent to the condition that all eigenvalues of A are larger than zero (remember that symmetric matrices only have real eigenvalues).

One application of symmetric positive definite matrices are energy functionals. The expression $x^T A x$ arises when discretising functional involving kinetic energies (e.g. energies of the from $E = \frac{1}{2}m|\nabla f|^2$ for f a given function).

For linear systems involving symmetric positive definite matrices we can derive a special algorithm, namely the Method of Conjugate Gradients (CG).

Lanczos - Arnoldi for symmetric matrices

Let us start with the Arnoldi recurrence relation

$$AV_m = V_m H_m + h_{m+1,m} v_{m+1} e_m^T$$

We know that H_m is an upper Hessenberg matrix (i.e. the upper triangular part plus the first lower triangular diagonal can only be nonzero). Also, we know from the orthogonality of the v_k vectors that

$$V_m^T A V_m = H_m.$$

Let A now be symmetric. From the symmetry of A an even nicer structure for H_m arises. H_m is upper Hessenberg, but now it is also symmetric. The only possible type of matrices to satisfy this condition are tridional matrices. These are matrices, where only the diagonal and the first upper and lower super/subdiagonals are nonzero.

Let us test this out. Below you find our simple implementation of Arnoldi's method. We then plot the resulting matrix H_m .

```
import numpy as np
def arnoldi(A, r0, m):
    """Perform m-1 step of the Arnoldi method."""
    n = A.shape[0]
    V = np.empty((n, m + 1), dtype=np.float64)
    H = np.zeros((m+1, m), dtype=np.float64)
    V[:, 0] = r0 / np.linalg.norm(r0)
    for index in range(m):
        # Multiply the previous vector with A
        tmp = A @ V[:, index]
        # Now orthogonalise against the previous basis vectors
        h = V[:, :index + 1].T @ tmp # h contains all inner products against previous vectors
       H[:index + 1, index] = h
        w = tmp - V[:, :index + 1] @ h # Subtract the components in the directions of the
previous vectors
        # Normalise and store
       H[index + 1, index] = np.linalg.norm(w)
        V[:, index + 1] = w[:] / H[index + 1, index]
    return V, H
```

The following code creates a random symmetric positive definite matrix.

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A numerical example

```
from numpy.random import RandomState

n = 500

rand = RandomState(0)
Q, _ = np.linalg.qr(rand.randn(n, n))
D = np.diag(rand.rand(n))
A = Q.T @ D @ Q
```

Now let's run Arnoldi's method and plot the matrix H. We are adding some artificial noise so as to ensure for the log-plot that all values are nonzero. The colorscale shows the logarithm of the magnitude of the entries.

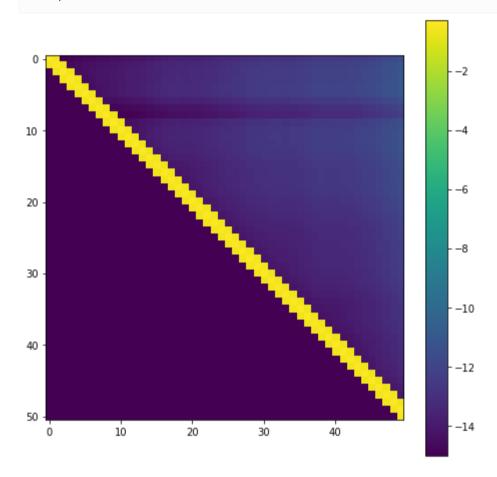
```
%matplotlib inline
from matplotlib import pyplot as plt

m = 50
r0 = rand.randn(n)
V, H = arnoldi(A, r0, m)

fig = plt.figure(figsize=(8, 8))
ax = fig.add_subplot(111)

im = ax.imshow(np.log10(1E-15 + np.abs(H)))
fig.colorbar(im)
```

<matplotlib.colorbar.Colorbar at 0x7ff7a79672b0>



It is clearly visible that only the main diagonal and the first upper and lower off-diagonal are nonzero, as expected. This hugely simplifies the Arnoldi iteration. Instead of orthogonalising Av_m against all previous vectors we only need to orthogonalise against v_m and v_{m-1} . All other inner products are already zero. Hence, the main orthogonalisation step now takes the form

$$w = Av_m - (v_m^T A v_m) v_m - (v_{m-1}^T A v_m) v_{m-1}.$$

Since the new vector w is composed of only 3 vectors. This is also called a 3-term recurrence. The big advantage is that in addition to Av_m we only need to keep v_m and v_{m-1} in memory. Hence, no matter how many iterations we do, the memory requirement remains constant, in contrast to Arnoldi for nonsymmetric matrices, where we need to keep all previous vectors in memory.

Arnoldi with a short recurrence relation for symmetric matrices has a special name. It is called Lanczos method.

Solving linear systems of equations with Lanczos

We can now proceed exactly as in the Full orthogonalisation method and arrive at the linear system of equations

$$T_m y_m = \|r_0\|_2 e_1,$$

where $x_m=x_0+V_my_m$ and $T_m=V_m^TAV_m$ is the tridiagonal matrix obtained from the Lanczos method.

The conjugate gradient method is an implementation of this approach. A very good derivation from Lanczos to CG is obtained in the beautiful book by Yousef Saad "<u>Iterative Methods for Sparse Linear Systems</u>", which is available online for free. Here, we will briefly motivate another approach to CG, which is a bit more intuitive and reveals more about the structure of the method, namely CG as an optimisation algorithm for a quadratic minimisation problem. One of the most beautiful summaries of this approach is contained in the paper <u>An introduction to the Conjugate Gradient Method Without the Agonizing Pain</u> by Jonathan Shewchuk.

A quadratic optimisation problem

We consider the quadratic minimisation problem

$$\min_{x \in \mathbb{R}^n} f(x)$$

with $f(x) = \frac{1}{2}x^TAx - b^Tx$. We have

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

and hence the only stationary point is the solution of the linear system Ax=b. Furthermore, it is really a minimiser since f''(x) is positive definite for all $x\in\mathbb{R}^n$ as A is positive definite.

The Method of Steepest Descent

Our first idea is the method of steepest descent. Remember that the negative gradient is a descent direction. Given a point x_k . We have

$$-\nabla f(x_k) = b - Ax_k := r_k.$$

The negative gradient is hence just the residual. Hence, we need to minimise along the direction of the residual, that is we will have $x_{k+1}=x_k+\alpha_k r_k$ for some value α_k . To compute α_k we just solve

$$rac{d}{dlpha}f(x_k+lpha r_k)=0$$

Since $\frac{d}{d\alpha}f(x_k+\alpha r_k)=r_{k+1}^Tr_k$ we just need to choose α_k such that r_{k+1} is orthogonal to r_k . The solution is given by $\alpha_k=\frac{r_k^Tr_k}{r_k^TAr_k}$. This gives us a complete method consisting of three steps to get from x_k to x_{k+1} .

$$egin{aligned} r_k &= b - A x_k \ lpha_k &= rac{r_k^T r_k}{r_k^T A r_k} \ x_{k+1} &= x_k + lpha_k r_k \end{aligned}$$

We are not going to derive the complete convergence analysis here but only state the final result. Let $\kappa:=\frac{\lambda_{max}}{\lambda_{min}}$, where λ_{max} and λ_{min} are the largest, respectively smallest eigenvalue of A (remember that all eigenvalues are positive since A is symmetric positive definite). The number κ is called the condition number of A. Let $e_k=x_k-x^*$ be the difference of the exact solution x^* satisfying $Ax^*=b$ and our current iterate x_k . Note that $r_k=-Ae_k$.

We now have that

$$\|e_k\|_A \leq \left(rac{\kappa-1}{\kappa+1}
ight)^k \|e_0\|_A,$$

where
$$\|e_k\|_A := \left(e_k^T A e_k\right)^{1/2}$$
 .

This is an extremely slow rate of convergence. Let $\kappa=10$, which is a fairly small number. Then the error reduces in each step only by a factor of $\frac{9}{11}\approx 0.81$ and we need 11 iterations for each digit of accuracy.

The method of conjugate directions

The steepest descent approach was not bad. But we want to improve on it. The problem with the steepest descent method is that we have no guarantee that we are reducing the error e_{k+1} as much as possible along our current direction r_k when we minimize. But we can fix this.

Let us pick a set of directions d_0,d_1,\ldots,d_{n-1} , which are mutually orthogonal, that is $d_i^Td_j=0$ for $i\neq j$. We now want to enforce the condition that

$$e_{k+1}^T d_k = 0.$$

This means that the remaining error is orthogonal to d_k and hence is a linear combination of all the other search directions. We have therefore exhausted all the information from d_k . Let's play this through.

We know that $e_{k+1}=x_{k+1}-x^*=x_k-x^*+lpha_k d_k=e_k+lpha_k d_k$.

It follows that

$$egin{aligned} e_{k+1}^T d_k &= d_k^T (e_k + lpha_k d_k) \ &= d_k^T e_k + lpha_k d_k^T d_k = 0 \end{aligned}$$

and therefore $lpha_k = -rac{d_k^T e_k}{d_k^T d_k}.$

Unfortunately, this does not quite work in practice as we don't know e_k . But there is a solution. Remember that $r_k=-Ae_k$. We just need an A in the right place. To achieve this we choose **conjugate directions**, that is we impose the condition that

$$d_i^T A d_i = 0$$

for i
eq j . We also impose the condition that $e_{k+1}^T A d_k = 0$. Writing this out we obtain

$$lpha_k = rac{d_k^T r_k}{d_k^T A d_k}.$$

This expression is computable if we have a suitable set of conjugate directions d_k . Moreoever, it guarantees that the method converges in at most n steps since in every iteration we are annihiliating the error in the direction of d_k and there are only n different directions.

Conjugate Gradients - Mixing steepest descent with conjugate directions

The idea of conjugate gradients is to obtain the conjugate directions d_i by taking the r_i (the gradients) and to Aorthogonalise (conjugate) them against the previous directions. We are leaving out the details of the derivation and
refer to the Shewchuk paper. But the final algorithm now takes the following form.

$$egin{aligned} d_0 &= r_0 = b - A x_0 \ lpha_i &= rac{r_i^T r_i}{d_i^T A d_i} \ x_{i+1} &= x_i + lpha_i d_i \ r_{i+1} &= r_i - lpha_i A d_i \ eta_{i+1} &= rac{r_{i+1}^T r_{i+1}}{r_i^T r_i} \ d_{i+1} &= r_{i+1} + eta_{i+1} d_i \end{aligned}$$

Conjugate Gradients has a much more favourable convergence bound than steepest descent. One can derive that

$$\|e_i\|_A \leq 2 \left(rac{\sqrt{\kappa}-1}{\sqrt{\kappa}+1}
ight)^i \|e_0\|_A.$$

If we choose again the example that $\kappa=10$ we obtain

$$||e_i||A \lesssim 0.52^i ||e_0||_A$$
.

Hence, we need around 4 iterations for each digits of accuracy instead of 11 for the method of steepest descent.

A numerical example

The following code creates a symmetric positive definite matrix.

We now solve the associated linear system with CG and plot the convergence.

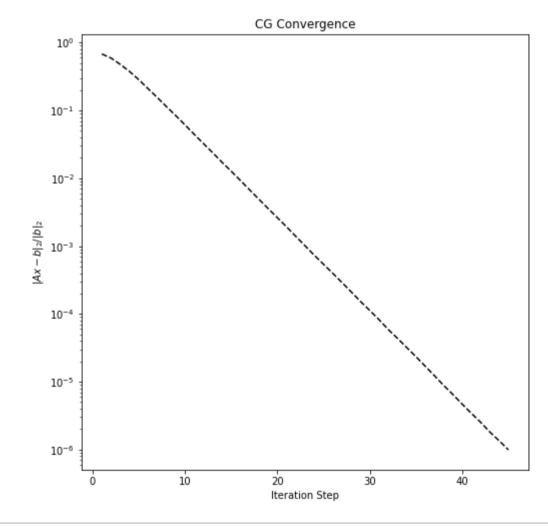
```
from scipy.sparse.linalg import cg

b = rand.randn(n)
residuals = []

callback = lambda x: residuals.append(np.linalg.norm(b - A @ x) / np.linalg.norm(b))

sol, _ = cg(A, b, tol=1E-6, callback=callback, maxiter=1000)

fig = plt.figure(figsize=(8, 8))
    ax = fig.add_subplot(111)
    ax.semilogy(1 + np.arange(len(residuals)), residuals, 'k--')
    ax.set_title('CG Convergence')
    ax.set_title('CG Convergence')
    ax.set_ylabel('$\|Ax-b\|_2 / \|b\|_2$')
```



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