

Schur decomposition

matrix A can be expressed as similar transform of matrix U :

$$A = QUQ^{-1}$$

where U is upper triangular and is called the Schur form of A . Q is unitary, and U has all zeros below the diagonal, which means the eigenvalues of A will be on the diagonal of U . Find Schur decomp = expose eigenvalues.

Real Schur Decomposition

If A has real eigenvalues it is a pure schur form as above. $[2 \times 2]$ -block schur form for complex (conjugate) eigenvalues. $[2 \times 2]$ block i on the diagonal has eigenvalues $\lambda_{i,1}$ and $\lambda_{i,2} = \lambda_{i,1}^*$, where λ_* is eigenvalues of the original matrix A .

Real analog of power method (7.3.1)

(7.4.1) is a real analog to the power method of finding eigenvalues. At a stage ($H_k = R_k U_k$), the machinery breaks down for real matrices having complex eigenvalues (no schur form can be achieved). However, calculating this term in a different way can yield a *real Schur decomposition* (eigenvalues come in complex conjugate pairs) and save the day.

Hessenberg QR step

The term $H_k = R_k U_k$ from the first section can be calculated efficiently if we initialize the calculation with a matrix (H_0) on reduced hessenberg form (order of magnitude $O(2)$ instead of $O(3)$). if H_0 is on reduced hessenberg form, it will stay that way if we calculate $H_k = R_k U_k$ by a **Hessenberg QR step**.

The Hessenberg reduction

It reads

$$U_0^T A U_0 = H$$

everything below subdiagonal is zero in H . U_0 is a product of matrices P_k who's purpose is to zero the k 'th column below the subdiagonal.

Code for solving this:

```
#include <armadillo>
#include <iostream>
#include <math.h>

using namespace std;
using namespace arma;
```

```

void house(double & beta, colvec & v, colvec & x) {
    double sigma, mu;

    int n = x.n_rows;

    colvec xspan = x(span(1, n - 1));
    sigma = dot(xspan, xspan);

    v = zeros<colvec> (n);
    v(span(1, n - 1)) = xspan;

    //zero if sigma is zero
    if (sigma == 0) {
        beta = 0;
    } else {
        mu = sqrt(x(0) * x(0) + sigma);

        if (x(0) <= 0) {
            v(0) = x(0) - mu;
        } else {
            v(0) = -sigma / (x(0) + mu);
        }

        beta = 2 / (sigma / (v(0) * v(0)) + 1);
        v /= v(0);
    }
}

void houseRedHessenberg(mat & A) {

    double beta;
    colvec v, x;
    mat ImBvvT;

    int n = A.n_cols;
    mat I = eye<mat> (n - 1, n - 1);

    for (int k = 0; k < n - 2; k++) {
        x = A(span(k + 1, n-1), k);
        house(beta, v, x);
        A(span(k + 1, n - 1), k) = x;

        ImBvvT = I - beta * v * strans(v);
        A(span(k + 1, n - 1), span(k, n - 1)) = ImBvvT * A(span(k + 1, n - 1), span(k, n - 1))
        A(span(), span(k + 1, n - 1)) = A(span(), span(k + 1, n - 1)) * ImBvvT;
    }
}

int main() {

```

```

mat A;
A << 1 << 5 << 7 << endr
      << 3 << 0 << 6 << endr
      << 4 << 3 << 1 << endr;

cout << A << endl;
houseRedHessenberg(A);
cout << A << endl;

return 0;
}

```

Runtime:

1.0000	5.0000	7.0000
3.0000	0	6.0000
4.0000	3.0000	1.0000
1.0000	8.6000	-0.2000
5.0000	4.9600	-0.7200
0	2.2800	-3.9600

which is the example on page 345.

Flops: $10n^3/3$.

Hessenberg matrix properties

Hessenberg decomposition is not unique. However, if there are no subdiagonal zero elements it is unique. Then it is called *unreduced*.

Companion matrix form

Non unitary analog of hessenberg decomposition. Using *Krylov matrices* K ($K^{-1} \neq K^H$), we can express the companion matrix C as

$$\det(I\lambda - K^{-1}AK) = \det(I\lambda - C) = c_0 + c_1\lambda_A + \dots + c_{n-1}\lambda_A^{n-1} + \lambda^n$$

which is the characteristic polynomial of A , which can be solved for the eigenvalues λ .

Hessenberg reduction via Gauss Transform

Uses only half the flops contra Householder. Chance that it uses more since it, like Gaussian elimination's partial pivoting, has a chance of 2^n growth.

Eigenvalue condition numbers $s(\lambda)^{-1}$ are not preserved, which gives a complicated error estimation process. This is the case of non-orthogonal similarities. (Flipping rows, multiplying rows etc. changes the signs and factors of the eigenvalues, whilst orthogonal similarities preserve everything).