

Summary for Numerical Methods for Computational Science and Engineering

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Computing with Matrices and Vectors

Machine Arithmetic

As machines can't compute in real numbers but compute with machine numbers, there are some constraints. Over- and underflow exist, the amount of numbers is finite. Thus, instead of checking for equality, we check if the difference of two numbers is smaller than some maximal relative error.

Cancellation

Cancellation occurs when Substrachting we attempt to:

- Subtract numbers of the same size.
- Divide by a small number close to zero.

To avoid cancellation, we simply avoid such kinds of subtractions and divisions by recasting expressions or use tricks like Taylor approximations.

Increasing Speed of Algorithms

For the following algorithms and formulas described, take into consideration that in many cases, the execution speed of a program can be increased significantly by simply precomputing things that would be computed multiple times, for example in a `for`-loop.

Direct Methods for Square Linear Systems of Equations

Gaussian Elimination

A is *invertible* or *regular* if and only if there exists a unique solution x for $Ax = b$, that is $x = A^{-1}b$.

In Eigen, we use the following code to solve $Ax = b$, and remember to exploit structure by using the appropriate function:

```
MatrixXd A;
VectorXd b, x;
...
// A has no special form
x = A.lu().solve(b);
// A is lower triangular
x = A.triangularView<Eigen::Lower>().solve(b);
// A is upper triangular
x = A.triangularView<Eigen::Upper>().solve(b);
// A is hermitian and positive definite
x = A.selfadjointView<Eigen::Upper>().llt().solve(b);
// A is hermitian and positive or negative definite
x = A.selfadjointView<Eigen::Upper>().ldlt().solve(b);
// Solving based on householder transformation
x = A.HouseholderQR<MatrixXd>().solve(b);
// Solving based on singular value decomposition
x = A.jacobiSvd<MatrixXd>().solve(b);
```

The generic asymptotic complexity is $O(n^3)$, but triangular linear systems can be solved in $O(n^2)$.

Solving $Ax = b$ by computing A^{-1} and calculating $x = A^{-1}b$ is unstable. Instead we use Gaussian elimination, which is stable and not affected by roundoff in practice.

Low-Rank Modifications

We assume a generic rank-one modification is applied to A :

$$\tilde{A} = A + uv^H$$

By using *block elimination*, we get:

$$\begin{bmatrix} A & u \\ v^H & -1 \end{bmatrix} \begin{bmatrix} \tilde{x} \\ \zeta \end{bmatrix} = \begin{bmatrix} b \\ 0 \end{bmatrix}$$

From this follows that:

$$\begin{aligned}
A\tilde{x} + u\zeta &= b \\
\tilde{x} &= A^{-1}(b - u\zeta) \\
&\dots \\
(1 + v^H A^{-1}u)\zeta &= v^H A^{-1}b \\
&\dots \\
A\tilde{x} &= b - \frac{uv^H A^{-1}}{1 + v^H A^{-1}u}b
\end{aligned}$$

Sparse Matrices

A matrix A is called *sparse*, if the number of non-zero entries $nnz(A)$ is much smaller than the size of A .

Storage Formats

Sparse matrices should be solved with `x = A.SparseLU<SparseMatrixType>().solve(b)` to further reduce the asymptotic complexity.

Also note that the product of sparse matrices isn't necessarily sparse.

Triplet/Coordinate List (COO) Format

Stores triplets with row indices, column indices and the associated values.

Compressed Row-Storage (CRS) Format

Stores the values, the corresponding column indices and row pointers that indicate the start of a new row.

```
SparseMatrix<double, RowMajor> A(rows, cols);
```

Compressed Column-Storage (CCS) Format

Same as CRS, but with column pointers and row indices instead.

```
SparseMatrix<double, ColMajor> A(rows, cols);
```

Direct Methods Linear Least Square Problems

Example: Linear Parameter Estimation

Assume we take m measurements and find the pairs (x_i, y_i) . We can express these measurements in the following overdetermined linear system:

$$\begin{bmatrix} x_1 & 1 \\ x_2 & 1 \\ \vdots & \vdots \\ x_m & 1 \end{bmatrix} \begin{bmatrix} \alpha \\ \beta \end{bmatrix} = \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_m \end{bmatrix}$$

Solution Concepts

For given A, b , the vector x is a least squares solution of $Ax = b$, if:

$$x \in \operatorname{argmin}_y \|Ay - b\|_2^2$$

For a least squares solution x , the vector Ax is the unique orthogonal projection of b onto the image of $\operatorname{Image}(A)$. From this follows that:

$$b - Ax \perp \operatorname{Image}(A)$$

$$A^T(b - Ax) = 0$$

$$A^T Ax = A^T b$$

If $m \geq n$ and $\operatorname{Kernel}(A) = 0$, then the linear system of equations $Ax = b$ has a unique least squares solution $x = (A^T A)^{-1} A^T b$.

We want to solve the *normal equation* $A^T Ax = A^T b$. To do this, we use the following procedure:

1. Compute the regular matrix $C = A^T A$.
2. Compute the right hand side vector $c = A^T b$.
3. Solve the symmetric positive definite linear system of equations $Cx = c$.

```
MatrixXd C = A.transpose() * A;  
VectorXd c = A.transpose() * b;  
VectorXd x = C.llt().solve(c);
```

The asymptotic complexity is $O(n^2 m + n^3)$.

Moore-Penrose Pseudoinverse

As there are many potential least square solutions, we define the *generalized solution* x^+ of a linear system of equations as:

$$x^+ = \operatorname{argmin}\{\|x\|_2, x \in \operatorname{lsq}(A, b)\}$$

The generalized solution x^+ of the linear systems of equations $Ax = b$ is given by:

$$x^+ = V(V^T A^T A V)^{-1} (V^T A^T b)$$

Where the matrix $V(V^T A^T A V)^{-1} V^T$ is called the *Moore-Penrose pseudoinverse* of A .

Orthogonal Transformation Methods

Transform $Ax = b$ to $\tilde{A}\tilde{x} = \tilde{b}$ such that $\operatorname{lsq}(A, b) = \operatorname{lsq}(\tilde{A}, \tilde{b})$ and $\tilde{A}\tilde{x} = \tilde{B}$ is easy to solve.

QR-Decomposition

Householder

Data Interpolation and Data Fitting in 1D

Given some data points (t_i, y_i) , $i = 0, \dots, n$, find a interpolant function f satisfying $f(t_i) = y_i$ and belonging to a set V of specific functions.

Global Polynomial Interpolation

The set of functions is $V = \mathcal{P}_k = \{f \mid f(t) = \alpha_k t^k + \dots + \alpha_0 t^0, \alpha_j \in \mathbb{R}\}$.

Polynomials are useful because ...

- ... they span a finite-dimensional vector space.
- ... they are smooth.
- ... they are easy to evaluate, differentiate and integrate.

Notice that $\dim \mathcal{P}_k = k + 1$. For n data points, we construct a $n + 1$ -dimensional Polynomial from \mathcal{P}_n . The polynomial will be unique.

We introduce the Lagrange polynomials as a helpful tool:

$$L_i(t) = \frac{(t - t_0) \dots (t - t_{i-1})(t - t_{i+1}) \dots (t - t_n)}{(t_i - t_0) \dots (t_i - t_{i-1})(t_i - t_{i+1}) \dots (t_i - t_n)}$$

Note that:

$$L_i(t_j) = \begin{cases} 1, & j = i \\ 0, & \text{else} \end{cases}$$

The polynomial is built by summing up the parts:

$$p(x) = \sum_{j=0}^n y_j L_j(x)$$

We can use the *Aitken-Neville* algorithm for increased efficiency, where l is the first and k is the last point included in the interpolation, such that $p_{0,n} = p$.

$$p_{k,k}(x) = y_k$$

$$p_{k,l}(x) = \frac{(x - t_k)p_{k+1,l}(x) - (x - t_l)p_{k,l-1}(x)}{t_l - t_k}$$

Another advantage is the update-friendliness of this algorithm, because there is no need to recompute the whole formula if another point is added.

Because adding another point affects all Lagrange polynomials, we introduce the new *Newton basis* with the purpose of being update-friendlier.

$$N_0(t) = 1, N_1(t) = (t - t_0), \dots, N_n(t) = \prod_{i=0}^{n-1} (t - t_i)$$

$$p(t) = \sum_{i=0}^{i \leq n} a_i N_i$$

To find a_i , we have to solve the following equations:

$$p(t_i) = y_i$$

Which leads us a triangular linear system to solve:

$$\begin{bmatrix} 1 & 0 & \dots & 0 \\ 1 & (t_1 - t_0) & & \vdots \\ \vdots & \vdots & & 0 \\ 1 & (t_n - t_0) & \dots & \prod_{i=0}^{n-1} (t_n - t_i) \end{bmatrix} \begin{bmatrix} a_0 \\ a_1 \\ \vdots \\ a_n \end{bmatrix} = \begin{bmatrix} y_0 \\ y_1 \\ \vdots \\ y_n \end{bmatrix}$$

We observe a similar recursion like before, and we can define analogous to $p_{l,m}$:

$$a_{l,m} = \frac{a_{l+1,m} - a_{l,m-1}}{t_m - t_l}$$

Instead of solving the triangular linear system, we use the *divided differences* algorithm.

$$y[t_i] = y_i$$

$$y[t_i, \dots, t_{i+k}] = \frac{y[t_{i+1}, \dots, t_{i+k}] - y[t_i, \dots, t_{i+k-1}]}{t_{i+k} - t_i}$$

$$a_i = y[t_0, \dots, t_i]$$

Shape Preserving Interpolation

Data (t_i, y_i) is called *monotonic* if $y_i \geq y_{i-1}$ or $y_i \leq y_{i-1}$ for $i = 1, \dots, n$.

Data (t_i, y_i) is called *convex* (*concave*) if $s_j \leq (\geq) s_{j+1}$ for $i = 1, \dots, n-1$, where $s_j = \frac{y_j - y_{j-1}}{t_j - t_{j-1}}$.

Interpolation is *shape preserving* if the shape features of the interpolant are the same as the shape features for the data.

If the shape is preserved (positivity/negativity, increasing/decreasing, convexity/concavity) for every piece of the interpolation, we say that the interpolant is *locally shape preserving*.

Cubic Hermite Interpolation

Given some data points $(t_j, y_j), j = 0, \dots, n$ and slopes $c_j \in \mathbb{R}$, find a piecewise cubic hermite interpolant s such that $s_{[t_{i-1}, t_i]} \in \mathcal{P}_3, i = 1, \dots, n$ and further $s(t_i) = y_i, s'(t_i) = c_i, i = 0, \dots, n$.

Splines

Trigonometric Interpolation

Least Squares Data Fitting