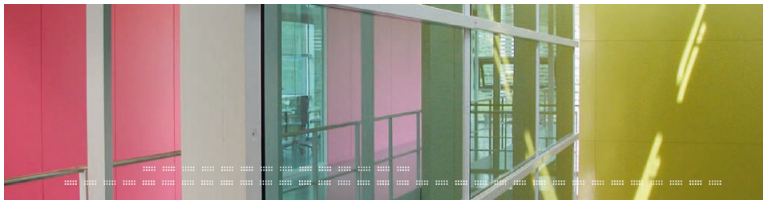


Linear Equation Systems and Interpolation

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Outline

Direct solution of linear equation systems

- Pivoting

- Cholesky Decomposition

Interpolation

- Preparatory Remarks

- Interpolation with Polynomials

- Aitken-Neville

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Pivoting

- ▶ In the last codes we quite carelessly divided by diagonal elements, e.g. $r_{j,j}$. What if $r_{j,j} = 0$?
- ▶ *partial pivoting* or *column pivoting*:
if $r_{j,j} = 0$ look for an entry $a_{i,j} \neq 0, i = j + 1, \dots, n$
 - ▶ there is no such $a_{i,j} \rightarrow$ completely vanishing column
 $\rightarrow \det(A) = 0$.
 - ▶ there are such $a_{i,j} \rightarrow$ choose the one with maximum absolute value (let it be $a_{k,j}$) and exchange the rows k and j (also right-hand side)
 - ▶ large pivot \rightarrow small elimination factors $l_{i,j}$.
- ▶ *total pivoting*:
search not only in column j but in the whole remaining submatrix. As columns may also have to be exchanged and thus \mathbf{x} may have to be reordered, it is more expensive.

Cholesky Decomposition I

For special matrices there are sometimes better algorithms. One special type are the positive definite matrices.

- ▶ A quadratic symmetric matrix $A \in \mathbb{R}^{n,n}$ is called positive definite, if all of its eigenvalues are positive or if

$$x^T A x > 0 \quad \forall x \neq 0$$

- ▶ In $A = LR$, the factor R can be further decomposed into a diagonal matrix D and a upper triangular \tilde{R} with ones on the diagonal:

$$A = L \cdot R = L \cdot D \cdot \tilde{R} \quad , D = \text{diag}(r_{1,1}, \dots, r_{n,n})$$

The symmetry of A gives us:

$$A^T = (L \cdot D \cdot \tilde{R})^T = \tilde{R}^T \cdot D \cdot L^T = L \cdot D \cdot \tilde{R} = A$$

Cholesky Decomposition II

- ▶ Because the decomposition is unique:

$$L = \tilde{R}^T \Rightarrow A = L \cdot D \cdot L^T =: \tilde{L} \cdot \tilde{L}^T$$

- ▶ In the last step D is equally distributed among L and L^T ($D = \tilde{D} \cdot \tilde{D}$ with $\tilde{D} = \text{diag}(\sqrt{r_{1,1}}, \dots, \sqrt{r_{n,n}})$).
- ▶ We rename $L = \tilde{L}$ for simpler notation:

$$A = L \cdot L^T$$

- ▶ In an LR-decomposition now the computation of the $r_{i,k}, i \neq k$ can be avoided \rightarrow half of memory requirements and computational costs.

Cholesky Decomposition Derivation I

As with the LR-decomposition let's begin with the matrix product (remember $l_{i,j} = 0$ for $i < j$ and $l_{i,j}^T = 0$ for $i > j$). We only need a lower triangular Matrix, thus $i \geq k$:

$$a_{i,k} = \sum_{j=1}^n l_{i,j} \cdot l_{j,k}^T = \sum_{j=1}^k l_{i,j} \cdot l_{j,k}^T = \sum_{j=1}^k l_{i,j} \cdot l_{k,j}, \quad i \geq k$$

For $i = k$ (the diagonal elements) this gives:

$$a_{k,k} = \sum_{j=1}^k l_{k,j}^2 \quad \rightarrow \quad l_{k,k} = \sqrt{a_{k,k} - \sum_{j=1}^{k-1} l_{k,j}^2}$$

Cholesky Decomposition Derivation II

For $i > k$:

$$a_{i,k} = \sum_{j=1}^k l_{i,j} \cdot l_{k,j} \quad \rightarrow \quad l_{i,k} = \left(a_{i,k} - \sum_{j=1}^{k-1} l_{i,j} \cdot l_{k,j} \right) / l_{k,k}$$

You need the elements in the columns with index $< k$ for the current row (i) and a previous row (k).

Thus when we compute L column by column (starting at diagonal), we are fine, as every needed value for $l_{i,k}$ has been computed.

Cholesky Decomposition Code

```
1 def cholesky(A,n):
2     L = []
3     for i in range(0,n):
4         L.append([0]*n)
5     for k in range(0,n):
6         L[k][k] = A[k][k]
7         for j in range(0,k):
8             L[k][k] = L[k][k] - pow(L[k][j],2)
9         L[k][k] = pow(L[k][k], 0.5)
10        for i in range(k+1,n):
11            L[i][k] = A[i][k]
12            for j in range(0,k):
13                L[i][k] = L[i][k] - L[i][j]*L[k][j]
14            L[i][k] = L[i][k] / L[k][k]
15    return L
```

Approximation and Interpolation

- ▶ **approximation**: approximate some function $f(x)$ by some $p(x)$ which is easy to handle, following certain rules.
- ▶ **interpolation**: special case of the approximation, where the approximant has to go through a set of given points.
 - ▶ **interpolation of a function**: The given points are taken from the function. In the following we assume, that the functions are sufficiently smooth (all necessary derivatives shall exist).
 - ▶ **interpolation of discrete data**: You are given only some data points (e.g. from a measurement).

Basics and some notation

- ▶ **nodes:** the abscissas x_j , where the interpolant is to be "fixed"
- ▶ **nodal points:** pairs (x_i, y_i) , i.e. nodes x_i with corresponding nodal values y_i
- ▶ polynomials are widespread interpolants:
 - ▶ \mathbb{P}_n : vector space of all polynomials with real coefficients of degree $\leq n$.
 - ▶ $\dim(\mathbb{P}_n) = n+1$
 - ▶ with the differential operator D^k (for $\frac{\partial^k}{\partial x^k}$):
$$D^{n+1}p = 0 \quad \forall p \in \mathbb{P}_n$$
- ▶ different variants:
 - ▶ **single nodes:** for x_i prescribe $p(x_i)$ (*Lagrange Interpolation*)
 - ▶ **multiple nodes:** also derivatives are prescribed. (*Hermite Interpolation*)

Interpolation with Polynomials

- ▶ $p \in \mathbb{P}_n$ is called *polynomial interpolant* of f with respect to the nodes $a = x_0 < x_1 < \dots < x_n = b$, if
$$p(x_i) = f(x_i) =: y_i \quad \forall i \in [0, n].$$
- ▶ The number of nodes determines the degree of the polynomial.
- ▶ Existence and Uniqueness of the solution of this interpolation problem is known from analysis.
- ▶ The difference $f(x) - p(x)$ is called *remainder* or *error term*.

Approaches



$$p(x) := \sum_{i=0}^n a_i \cdot x^i$$

Make the incidental test for each node and solve the resulting system of $n + 1$ linear equations.

- ▶ With the Lagrange polynomials

$$L_k(x) = \prod_{i=0, i \neq k}^n \frac{x - x_i}{x_k - x_i} \Rightarrow L_k(x_j) = \delta_{k,j}$$

you get:

$$p(x) = \sum_{k=0}^n y_k \cdot L_k(x)$$

Aitken-Neville's Scheme

Often, we don't need the interpolant p , but the evaluation $p(x)$ at one or several points.

- ▶ define auxiliary polynomials $p_{i,k}(x)$ of degree k , which interpolate the nodes (x_l, y_l) for $l = i, \dots, i + k$
- ▶ do this recursively with

$$p_{i,k}(x) := \frac{x_{i+k} - x}{x_{i+k} - x_i} \cdot p_{i,k-1}(x) + \frac{x - x_i}{x_{i+k} - x_i} \cdot p_{i+1,k-1}(x)$$

- ▶ Starting values are $p_{0,0}(x) = y_0, \dots, p_{n-1,0}(x) = y_{n-1}$
- ▶ The value $p_{0,n-1}(x)$ will be the result you wanted.

Aitken-Neville Code

```
1 def aitken (x,y,x0,n):
2     x0 = float(x0)
3     p = []
4     for i in range(0,n):
5         p.append([0]*n)
6
7     for i in range(0,n):
8         p[i][0] = y[i]
9     for k in range (1,n):
10        for i in range(0, n-k):
11            p[i][k] = (x[i+k]-x0) / (x[i+k]-x[i]) * p[i][k-1]
12                    (x0-x[i]) / (x[i+k]-x[i]) * p[i+1][k-1]
13    return p[0][n-1]
```

Divided Differences I

Let us denote the highest coefficient of $p_{i,k}$ as:

$$[x_i, \dots, x_{i+k}]f$$

and call it *divided differences of f of order k with respect to x_i, \dots, x_{i+k}* , i.e.

$$p_{i,k}(x) = [x_i, \dots, x_{i+k}]f \cdot x^k + r \quad , r \in \mathbb{P}_{k-1}.$$

So we have:

$$\begin{aligned} [x_i]f &= f(x_i) = y_i \\ [x_i, x_{i+1}]f &= \frac{y_{i+1} - y_i}{x_{i+1} - x_i} \end{aligned}$$

Generally (with Aitken-Neville):

$$[x_i, \dots, x_{i+k}]f := \frac{[x_{i+1}, \dots, x_{i+k}]f - [x_i, \dots, x_{i+k-1}]f}{x_{i+k} - x_i}$$

Divided Differences in Python

```
1 def divdiff (xList, yList, xVal):
2     length = len (xList)
3     diffs = []
4     for i in range (0, length):
5         diffs.append (yList[i])
6     for i in range (1,length):
7         for j in range (length-i, length):
8             lowerIndex = length - i - 1
9             diffs[j] = (diffs[j] - diffs[j-1])/(xList[j]-xList
10 retVal = diffs[0]
11 product = 1
12 for i in range (1, length):
13     product *= xVal - xList[i-1]
14     retVal += diffs[i] * product
15 return retVal
```

Newton's interpolation formula

The used basis for \mathbb{P}_n is

$$\{1, x - x_0, (x - x_0) \cdot (x - x_1), (x - x_0) \cdot \dots \cdot (x - x_{n-1})\}.$$

The approach here will be:

$$p_{0,k}(x) = \sum_{i=0}^k c_i \left(\prod_{j=0}^{i-1} (x - x_j) \right)$$

As c_k is the highest coefficient of $p_{0,k}$:

$$c_k = [x_0, \dots, x_k]f$$

Adding one node will only add another summand:

$$p_{0,k+1} = \sum_{i=0}^{k+1} c_i \left(\prod_{j=0}^{i-1} (x - x_j) \right) = p_{0,k}(x) + c_{k+1} \cdot \prod_{j=0}^{i-1} (x - x_j)$$

Newton's interpolation formula II

As adding a new node only adds a new summand, the previously calculated c_k don't change, and in the end we get *Newton's interpolation formula*:

$$\begin{aligned} p_{0,n}(x) = p(x) &= [x_0]f + \\ & [x_0, x_1]f \cdot (x - x_0) + \\ & [x_0, x_1, x_2]f \cdot (x - x_0) \cdot (x - x_1) + \\ & [x_0, \dots, x_3]f \cdot (x - x_0) \cdot (x - x_1) \cdot (x - x_2) + \\ & \vdots \\ & [x_0, \dots, x_n]f \cdot \prod_{i=0}^{n-1} (x - x_i) \end{aligned}$$

Interpolation Error I

Now we estimate the interpolation error $|f(x) - p(x)|$.

- ▶ Let $f \in C^n([x_0, x_n])$, i.e. n times continuously differentiable on $[x_0, x_n]$. Then there exists a $\xi \in [x_0, x_n]$ with:

$$[x_0, \dots, x_n]f = \frac{D^n f(\xi)}{n!}$$

- ▶ Now consider some $\tilde{x} \in [x_0, x_n] \setminus \{x_0, \dots, x_n\}$ and let $\tilde{p}(x) \in \mathbb{P}_{n+1}$ interpolate also \tilde{x} in addition to x_0, \dots, x_n :

$$\begin{aligned} f(\tilde{x}) - p(\tilde{x}) &= P(\tilde{x}) - p(\tilde{x}) = [x_0, \dots, x_n, \tilde{x}]f \cdot \prod_{i=0}^n (\tilde{x} - x_i) \\ &= \frac{D^{n+1} f(\xi)}{(n+1)!} \cdot \prod_{i=0}^n (\tilde{x} - x_i) \end{aligned}$$

Interpolation Error II

For equidistant nodes with mesh width $h := x_{i+1} - x_i$, the product can be estimated.

In the worst case the distance of \tilde{x} to the farrest node is nh ; to the second farrest $(n-1)h$ and so on.

Thus we can estimate:

$$|f(\tilde{x}) - p(\tilde{x})| \leq \frac{\max_{[x_0, x_n]} |D^{n+1} f(x)|}{n+1} \cdot h^{n+1} \in \mathcal{O}(h^{n+1})$$