

Biostatistics 615/815 Lecture 20: Linear Algebra with C++

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Programming with Matrix

Why Matrix matters?

- Many statistical models can be well represented as matrix operations
 - Linear regression
 - Logistic regression
 - Mixed models
- Efficient matrix computation can make difference in the practicality of a statistical method
- Understanding C++ implementation of matrix operation can expedite the efficiency by orders of magnitude

Ways for Matrix programming in C++

- Implementing Matrix libraries on your own
 - Implementation can well fit to specific need
 - Need to pay for implementation overhead
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 - Low-level Fortran/C API
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 - Used in many statistical packages including R
 - Not user-friendly interface use.
 - boost supports C++ interface for BLAS
- Using a third-party library, Eigen package
 - A convenient C++ interface
 - Reasonably fast performance
 - Supports most functions BLAS/LAPACK provides

Using a third party library

Downloading and installing Eigen package

- Download at <http://eigen.tuxfamily.org/>
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Using Eigen package

- Add `-I [PARENT_PATH_OF Eigen/]` option when compile
- No need to install separate library. Including header files is sufficient

Example usages of Eigen library

```
#include <iostream>
#include <Eigen/Dense> // For non-sparse matrix
using namespace Eigen; // avoid using Eigen::
int main()
{
    Matrix2d a;           // 2x2 matrix type is defined for convenience
    a << 1, 2,
        3, 4;
    MatrixXd b(2,2);     // but you can define the type from arbitrary-size matrix
    b << 2, 3,
        1, 4;
    std::cout << "a + b =\n" << a + b << std::endl; // matrix addition
    std::cout << "a - b =\n" << a - b << std::endl; // matrix subtraction
    std::cout << "Doing a += b;" << std::endl;
    a += b;
    std::cout << "Now a =\n" << a << std::endl;
    Vector3d v(1,2,3);           // vector operations
    Vector3d w(1,0,0);
    std::cout << "-v + w - v =\n" << -v + w - v << std::endl;
}
```

More examples

```
#include <iostream>
#include <Eigen/Dense>

using namespace Eigen;
int main()
{
    Matrix2d mat;           // 2*2 matrix
    mat << 1, 2,
          3, 4;
    Vector2d u(-1,1), v(2,0); // 2D vector
    std::cout << "Here is mat*mat:\n" << mat*mat << std::endl;
    std::cout << "Here is mat*u:\n" << mat*u << std::endl;
    std::cout << "Here is u^T*mat:\n" << u.transpose()*mat << std::endl;
    std::cout << "Here is u^T*v:\n" << u.transpose()*v << std::endl;
    std::cout << "Here is u*v^T:\n" << u*v.transpose() << std::endl;
    std::cout << "Let's multiply mat by itself" << std::endl;
    mat = mat*mat;
    std::cout << "Now mat is mat:\n" << mat << std::endl;
    return 0;
}
```

More examples

```
#include <Eigen/Dense>
#include <iostream>
using namespace Eigen;
int main()
{
    MatrixXd m(2,2), n(2,2);
    MatrixXd result(2,2);
    m << 1,2,
        3,4;
    n << 5,6,7,8;
    result = m * n;
    std::cout << "-- Matrix m*n: --" << std::endl << result << std::endl << std::endl;
    result = m.array() * n.array();
    std::cout << "-- Array m*n: --" << std::endl << result << std::endl << std::endl;
    result = m.cwiseProduct(n);
    std::cout << "-- With cwiseProduct: --" << std::endl << result << std::endl << std::endl;
    result = (m.array() + 4).matrix() * m;
    std::cout << "-- (m+4)*m: --" << std::endl << result << std::endl << std::endl;
    return 0;
}
```

Time complexity of matrix computation

Square matrix multiplication / inversion

- Naive algorithm : $O(n^3)$
- Strassen algorithm : $O(n^{2.807})$
- Coppersmith-Winograd algorithm : $O(n^{2.376})$ (with very large constant factor)

Determinant

- Laplace expansion : $O(n!)$
- LU decomposition : $O(n^3)$
- Bareiss algorithm : $O(n^3)$
- Fast matrix multiplication algorithm : $O(n^{2.376})$

Computational considerations in matrix operations

Avoiding expensive computation

- Computation of $\mathbf{u}' \mathbf{A} \mathbf{B} \mathbf{v}$

Computational considerations in matrix operations

Avoiding expensive computation

- Computation of $\mathbf{u}'AB\mathbf{v}$
- If the order is $((\mathbf{u}'(AB))\mathbf{v})$
 - $O(n^3) + O(n^2) + O(n)$ operations
 - $O(n^2)$ overall

Computational considerations in matrix operations

Avoiding expensive computation

- Computation of $\mathbf{u}' A B \mathbf{v}$
- If the order is $((\mathbf{u}'(AB))\mathbf{v})$
 - $O(n^3) + O(n^2) + O(n)$ operations
 - $O(n^2)$ overall
- If the order is $((\mathbf{u}' A)B)\mathbf{v}$
 - Two $O(n^2)$ operations and one $O(n)$ operation
 - $O(n^2)$ overall

Quadratic multiplication

Same time complexity, but one is slightly more efficient

- Computing $\mathbf{x}' A \mathbf{y}$.
- $O(n^2) + O(n)$ if ordered as $(\mathbf{x}' A) \mathbf{y}$.
- Can be simplified as $\sum_i \sum_j x_i A_{ij} y_j$

A symmetric case

- Computing $\mathbf{x}' A \mathbf{x}$ where $A = LL'$
- $\mathbf{u} = L' \mathbf{x}$ can be computed more efficiently than $A \mathbf{x}$.
- $\mathbf{x}' A \mathbf{x} = \mathbf{u}' \mathbf{u}$

Solving linear systems

Problem

Find \mathbf{x} that satisfies $A\mathbf{x} = \mathbf{b}$

A simplest approach

- Calculate A^{-1} , and $\mathbf{x} = A^{-1}\mathbf{b}$
- Time complexity is $O(n^3) + O(n^2)$
- A has to be invertible
- Potential issue of numerical instability

Using matrix decomposition to solve linear systems

LU decomposition

- $A = LU$, making $U\mathbf{x} = \mathbf{L}^{-1}\mathbf{b}$
- A needs to be square and invertible.
- Fewer additions and multiplications
- Precision problems may occur

QR decomposition

- $A = QR$ where A is $m \times n$ matrix
- Q is orthogonal matrix, $Q'Q = I$.
- R is $m \times n$ upper-triangular matrix, $R\mathbf{x} = Q'\mathbf{b}$.

Cholesky decomposition

- A is a square, symmetric, and positive definite matrix.
- $A = U'U$ is a special case of LU decomposition
- Computationally efficient and accurate

Solving least square

Solving via inverse

- Most straightforward strategy
- $\mathbf{y} = X\beta + \epsilon$, \mathbf{y} is $n \times 1$, X is $n \times p$.
- $\beta = (X'X)^{-1}X'\mathbf{y}$.
- Computational complexity is $O(np^2) + O(np) + O(p^3)$.
- The computation may become unstable if $X'X$ is singular
- Need to make sure that $\text{rank}(X) = p$.

Singular value decomposition

Definition

A $m \times n$ ($m \geq n$) matrix A can be represented as $A = UDV^T$ such that

- U is $m \times n$ matrix with orthogonal columns ($U'U = I_n$)
- D is $n \times n$ diagonal matrix with non-negative entries
- V^T is $n \times n$ matrix with orthogonal matrix ($V'V = VV' = I_n$).

Computational complexity

- $4m^2n + 8mn^2 + 9m^3$ for computing U, V , and D .
- $4mn^2 + 8n^3$ for computing V and D only.
- The algorithm is numerically very stable

Stable inference of least square using SVD

$$\begin{aligned} X &= UDV' \\ \beta &= (X'X)^{-1}X'\mathbf{y} \\ &= (VDU'UDV')^{-1} VDU'\mathbf{y} \\ &= (VD^2V')^{-1} VDU'\mathbf{y} \\ &= VD^{-2}V' VDU'\mathbf{y} \\ &= VD^{-1}U'\mathbf{y} \end{aligned}$$

Stable inference of least square using SVD

```
#include <iostream>
#include <Eigen/Dense>

using namespace std;
#using namespace Eigen;

int main()
{
    MatrixXf A = MatrixXf::Random(3, 2);
    cout << "Here is the matrix A:\n" << A << endl;
    VectorXf b = VectorXf::Random(3);
    cout << "Here is the right hand side b:\n" << b << endl;
    cout << "The least-squares solution is:\n"
        << A.jacobiSvd(ComputeThinU | ComputeThinV).solve(b) << endl;
}
```

Linear Regression

Linear model

- $\mathbf{y} = X\beta + \epsilon$, where X is $n \times p$ matrix
- Under normality assumption, $y_i \sim N(X_i\beta, \sigma^2)$.

Key inferences under linear model

- Effect size : $\hat{\beta} = (X^T X)^{-1} X^T \mathbf{y}$
- Residual variance : $\hat{\sigma}^2 = (\mathbf{y} - X\hat{\beta})^T (\mathbf{y} - X\hat{\beta}) / (n - p)$
- Variance/SE of $\hat{\beta}$: $\text{Var}(\hat{\beta}) = \hat{\sigma}^2 (X^T X)^{-1}$
- p-value for testing $H_0 : \beta_i = 0$ or $H_o : R\beta = 0$.

Using R to solve linear model

```
> y <- rnorm(100)
> x <- rnorm(100)
> summary(lm(y~x))
```

Call:

```
lm(formula = y ~ x)
```

Residuals:

Min	1Q	Median	3Q	Max
-2.15759	-0.69613	0.08565	0.70014	2.62488

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	0.02722	0.10541	0.258	0.797
x	-0.18369	0.10559	-1.740	0.085 .

Signif. codes: ...

Residual standard error: 1.05 on 98 degrees of freedom

Multiple R-squared: 0.02996, Adjusted R-squared: 0.02006

F-statistic: 3.027 on 1 and 98 DF, p-value: 0.08505

Dealing with large data with lm

```
> y <- rnorm(5000000)
> x <- rnorm(5000000)
> system.time(print(summary(lm(y~x))))
```

Call:

```
lm(formula = y ~ x)
```

Residuals:

Min	1Q	Median	3Q	Max
-5.1310	-0.6746	0.0004	0.6747	5.0860

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	-0.0005130	0.0004473	-1.147	0.251
x	0.0002359	0.0004473	0.527	0.598

Residual standard error: 1 on 4999998 degrees of freedom

Multiple R-squared: 5.564e-08, Adjusted R-squared: -1.444e-07

F-statistic: 0.2782 on 1 and 4999998 DF, p-value: 0.5979

user	system	elapsed
57.434	14.229	100.607

A case for simple linear regression

A simpler model

- $\mathbf{y} = \beta_0 + \mathbf{x}\beta_1 + \epsilon$
- $X = [1 \ \mathbf{x}], \beta = [\beta_0 \ \beta_1]^T.$

Question of interest

Can we leverage this simplicity to make a faster inference?

A faster inference with simple linear model

Ingredients for simplification

- $\sigma_y^2 = (\mathbf{y} - \bar{\mathbf{y}})^T(\mathbf{y} - \bar{\mathbf{y}})/(n - 1)$
- $\sigma_x^2 = (\mathbf{x} - \bar{\mathbf{x}})^T(\mathbf{x} - \bar{\mathbf{x}})/(n - 1)$
- $\sigma_{xy} = (\mathbf{x} - \bar{\mathbf{x}})^T(\mathbf{y} - \bar{\mathbf{y}})/(n - 1)$
- $\rho_{xy} = \sigma_{xy}/\sqrt{\sigma_x^2\sigma_y^2}$.

Making faster inferences

- $\hat{\beta}_1 = \rho_{xy}\sqrt{\sigma_y^2/\sigma_x^2}$
- $\text{SE}(\hat{\beta}_1) = \sqrt{(n - 1)\sigma_y^2(1 - \rho_{xy}^2)/(n - 2)}$
- $t = \rho_{xy}\sqrt{(n - 2)/(1 - \rho_{xy}^2)}$ follows t-distribution with d.f. $n - 2$

A faster R implementation

```
# note that this is an R function, not C+
fastSimpleLinearRegression <- function(y, x) {
  y <- y - mean(y)
  x <- x - mean(x)
  n <- length(y)
  stopifnot(length(x) == n)          # for error handling
  s2y <- sum( y * y ) / ( n - 1 )  # \sigma_y^2
  s2x <- sum( x * x ) / ( n - 1 )  # \sigma_x^2
  sxy <- sum( x * y ) / ( n - 1 )  # \sigma_xy
  rxy <- sxy / sqrt( s2y * s2x )   # \rho_xy
  b <- rxy * sqrt( s2y / s2x )
  se.b <- sqrt( ( n - 1 ) * s2y * ( 1 - rxy * rxy ) / (n-2) )
  tstat <- rxy * sqrt( ( n - 2 ) / ( 1 - rxy * rxy ) )
  p <- pt( abs(t), n - 2, lower.tail=FALSE )^2
  return(list( beta = b, se.beta = se.b, t.stat = tstat, p.value = p ))
}
```

Now it became must faster

```
> system.time(print(fastSimpleLinearRegression(y,x)))
$beta
[1] 0.0002358472

$se.beta
[1] 1.000036

$t.stat
[1] 0.5274646

$p.value
[1] 0.597871

  user  system elapsed
0.382   1.849   3.042
```

Dealing with even larger data

Problem

- Supposed that we now have 5 billion input data points
- The issue is how to load the data
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What we want

- As fast performance as before
- But do not store all the data into memory
- R cannot be the solution in such cases - use C++ instead

Streaming the inputs to extract sufficient statistics

Sufficient statistics for simple linear regression

- ① n
- ② $\sigma_x^2 = \hat{\text{Var}}(x) = (\mathbf{x} - \bar{x})^T(\mathbf{x} - \bar{x})/(n - 1)$
- ③ $\sigma_y^2 = \hat{\text{Var}}(y) = (\mathbf{y} - \bar{y})^T(\mathbf{y} - \bar{y})/(n - 1)$
- ④ $\sigma_{xy} = \hat{\text{Cov}}(x, y) = (\mathbf{x} - \bar{x})^T(\mathbf{y} - \bar{y})/(n - 1)$

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Extracting sufficient statistics from stream

- $\sum_{i=1}^n x = n\bar{x}$
- $\sum_{i=1}^n y = n\bar{y}$
- $\sum_{i=1}^n x^2 = \sigma_x^2(n - 1) + n\bar{x}^2$
- $\sum_{i=1}^n y^2 = \sigma_y^2(n - 1) + n\bar{y}^2$
- $\sum_{i=1}^n xy = \sigma_{xy}(n - 1) + n\bar{xy}$

Implementation : Streamed simple linear regression

```
#include <iostream>
#include <fstream>
#include <boost/math/distributions/students_t.hpp>
using namespace boost::math;      // for calculating p-values from t-statistic
int main(int argc, char** argv) {
    std::ifstream ifs(argv[1]);    // read file from the file arguments
    double x, y;                  // temporay values to store the input
    double sumx = 0, sumsqx = 0, sumy = 0, sumsqy = 0, sumxy = 0;
    int n = 0;

    // extract a set of sufficient statistics
    while( ifs >> y >> x ) { // assuming each input line feeds y and x
        sumx += x;
        sumy += y;
        sumxy += (x*y);
        sumsqx += (x*x);
        sumsqy += (y*y);
        ++n;
    }
}
```

Streamed simple linear regression (cont'd)

```
// convert the set of sufficient statistics to
double s2y = (sumsqy - sumy*sumy/n)/(n-1);           // s2y = \sigma_y^2
double s2x = (sumsqx - sumx*sumx/n)/(n-1);           // s2x = \sigma_x^2
double sxy = (sumxy - sumx*sumy/n)/(n-1);           // sxy = \sigma_{xy}
double rxy = sxy/(s2x*s2y);                          // rxy = cor(x,y)

// calculate beta, SE(beta), and p-values
double beta = rxy * s2y / s2x;
double seBeta = s2y * sqrt( (n-1) * ( 1 - rxy*rxy ) / (n-2) );
double t = rxy * sqrt( (n-2)/(1-rxy*rxy) );          // t-statistics

students_t dist(n-2);      // use student's t-distribution to compute p-value
double pvalue = 2.0*cdf(complement(dist, t > 0 ? t : (0-t) ));
```

Streamed simple linear regression (cont'd)

```
    std::cout << "Number of observations      = " << n << std::endl;
    std::cout << "Effect size - beta        = " << beta << std::endl;
    std::cout << "Standard error - SE(beta) = " << seBeta << std::endl;
    std::cout << "Student's-t statistic     = " << t << std::endl;
    std::cout << "Two-sided p-value          = " << pvalue << std::endl;
    return 0;
}
```

Summary - Simple Linear Regression

- A linear regression with one predictor and intercept
- `lm()` function in R may be computationally slow for large input
- Faster inference is possible by computing a set of summary statistics in linear time
- Streaming via C++ programming further resolves the memory overhead
- The idea can be applied in more sophisticated, large-scale analyses.