

# HWI - Summing

- $1e-8$  smaller than machine epsilon (float)
- Forward sum fails utterly
- Backward sum does better (why not correctly?)

```
$ ./part1
Left sum: 1
Right sum: 1.25
```

# HW1 - Summing

- Lots of ways around this:

```
float pairwisesum(float *list, const int n) {  
  
    if (n == 1) return list[0];  
  
    const int newn = n/2 + n%2;  
    float *sums = new float[newn];  
  
    for (int i=0; i<n/2; i++)  
        sums[i] = list[2*i] + list[2*i+1];  
  
    if (n%2 == 1)  
        sums[n/2] = list[n-1];  
  
    return pairwisesum(sums, newn);  
}
```

# HW1 - Summing

- Lots of ways around this:

```
float kahensum(float *list, const int n) {  
  
    float tot = 0.;  
    float comp = 0.;  
  
    for (int i=0; i<n; i++) {  
        float y = list[i] - comp;  
        float t = tot + y;  
        comp = (t - tot) - y;  
        tot = t;  
    }  
  
    return tot;  
}
```

# HW1 - Summing

- Lots of ways around this:

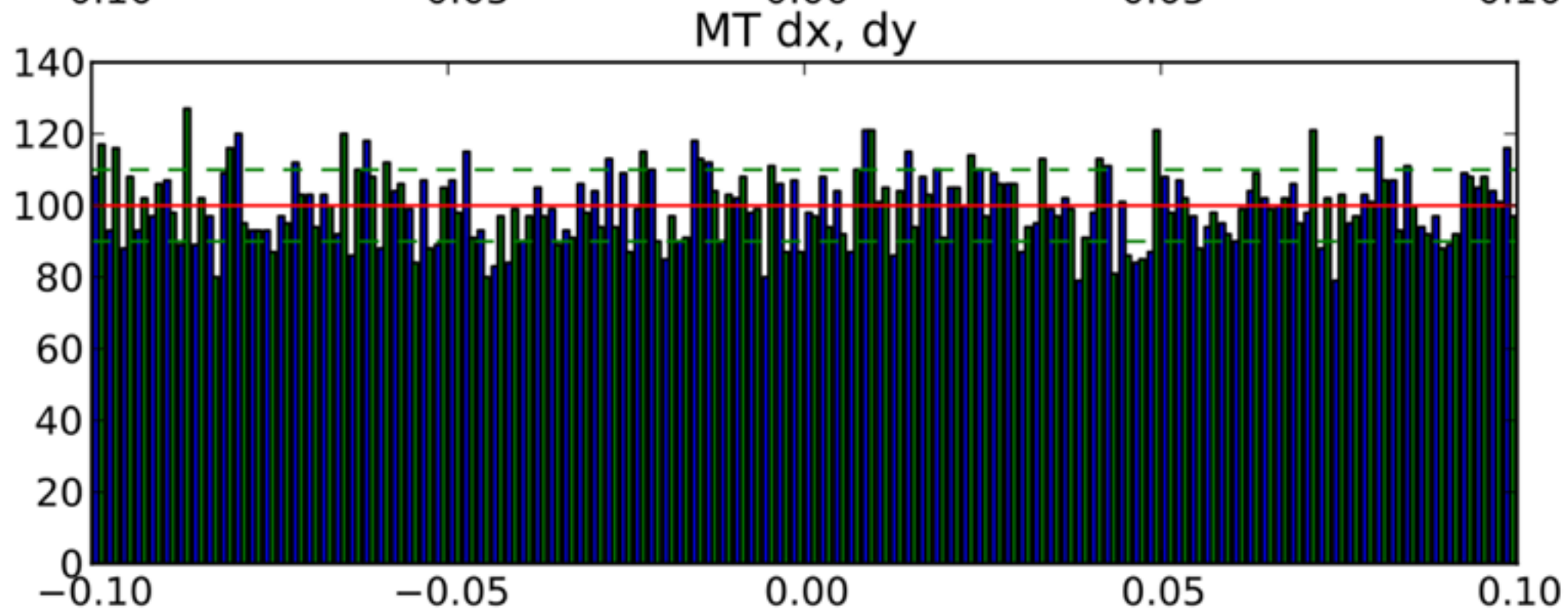
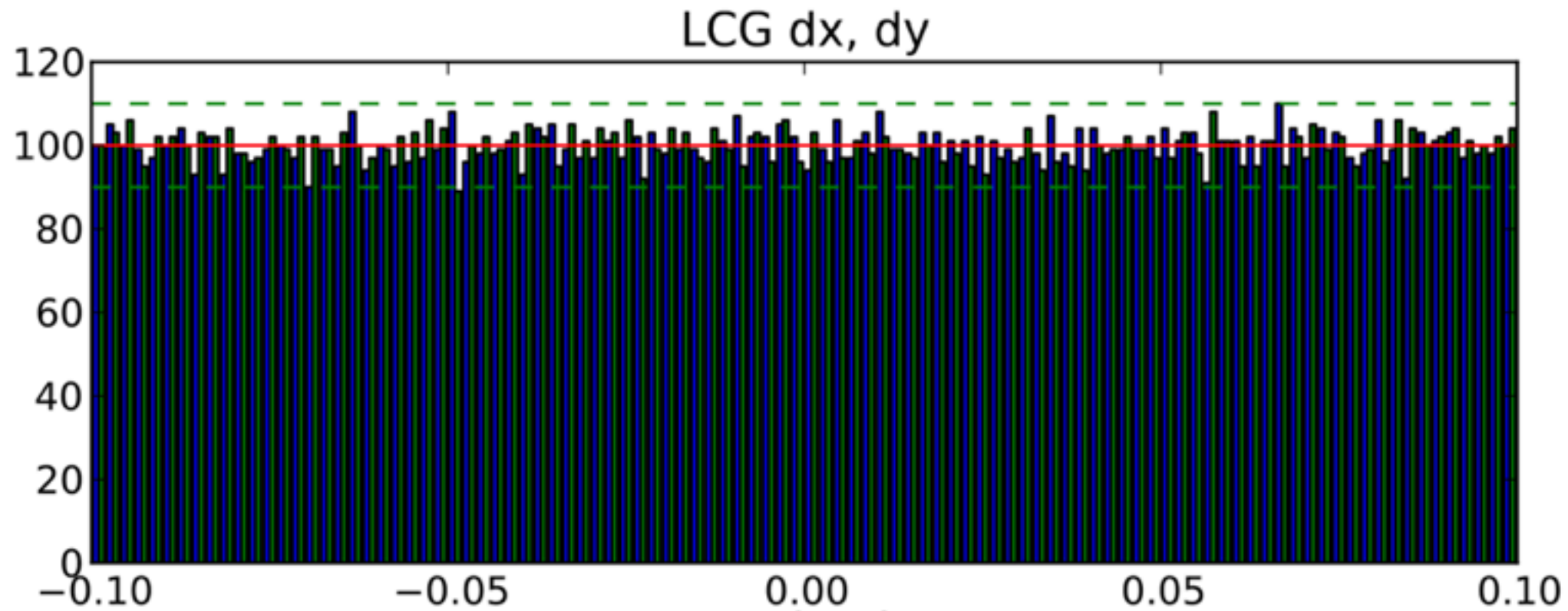
```
double doublesum(float *list, const int n) {  
    double tot = 0.;  
  
    for (int i=0; i<n; i++)  
        tot += list[i];  
  
    return tot;  
}
```

# HWI - Summing

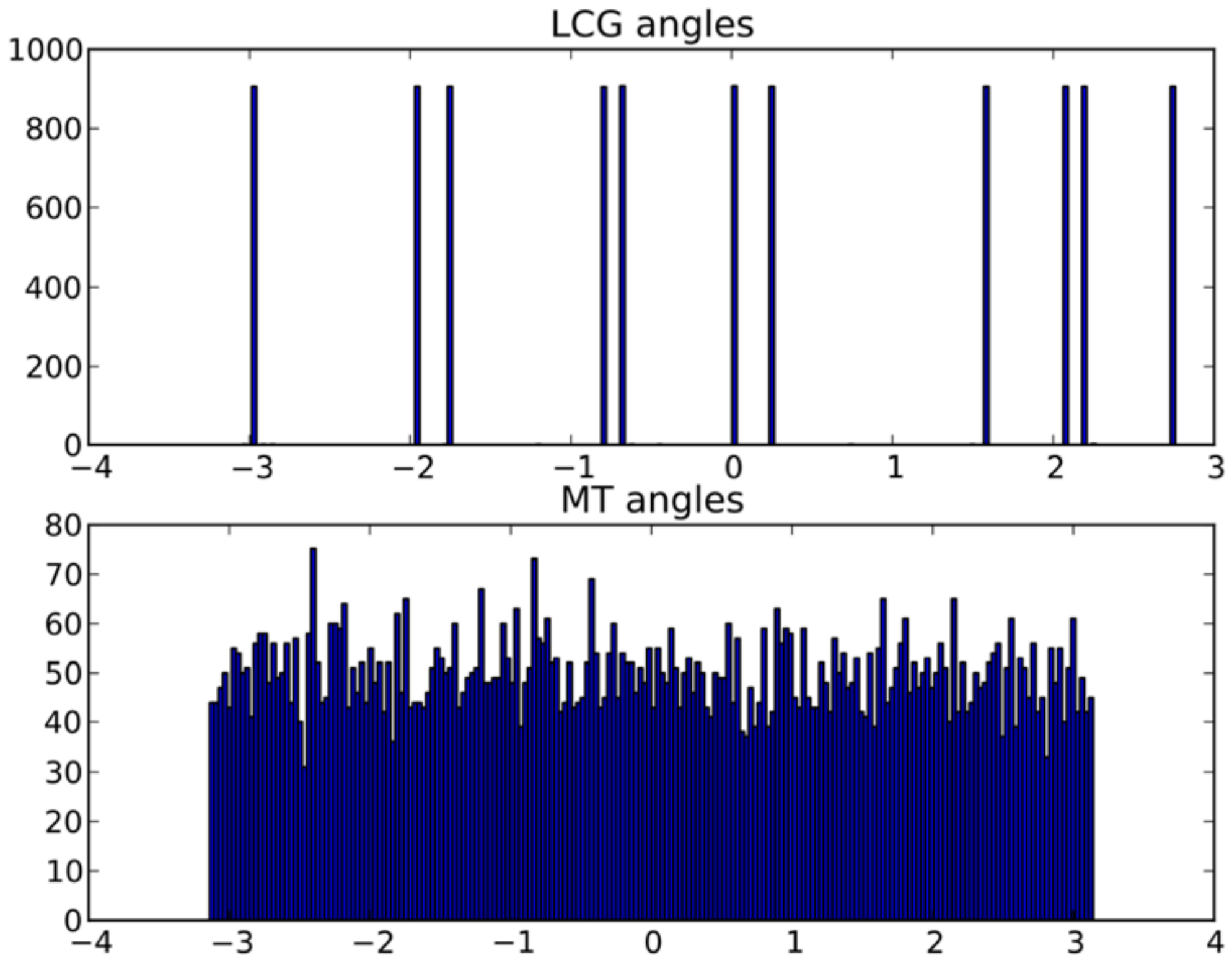
- Lots of ways around this:

```
$ ./part1
Left sum: 1
Right sum: 1.25
Pairwise sum: 2
Kahen sum: 2
Double precision sum: 2
```

# HWI - Random Walks



# HWI - Random Walks



# HWI - Seed

- Some issues with seeding
- General workflow; seed **once**, then generate all the random numbers you need.
- Showing how LCG worked may have confused things; seed was just last (integer) random deviate chosen



# HWI - Seed

- In general, current state of a PRNG can be quite large.
- Generally explicit functions to query state, set state (so can continue exactly where left off)
- Most PRNGs also have a convenience function to set state from small ( $\sim 1$  int) seed; bootstrap state from seed + smaller RNG
- Use once; *don't* keep seeding - don't know how it interacts with the PRNG

# Numerical Linear Algebra

Scientific Computing Course  
Mar 2013

# Conclusion

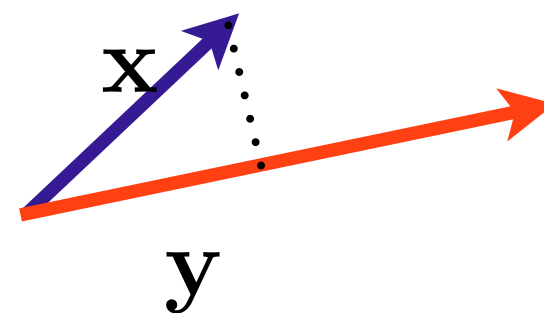
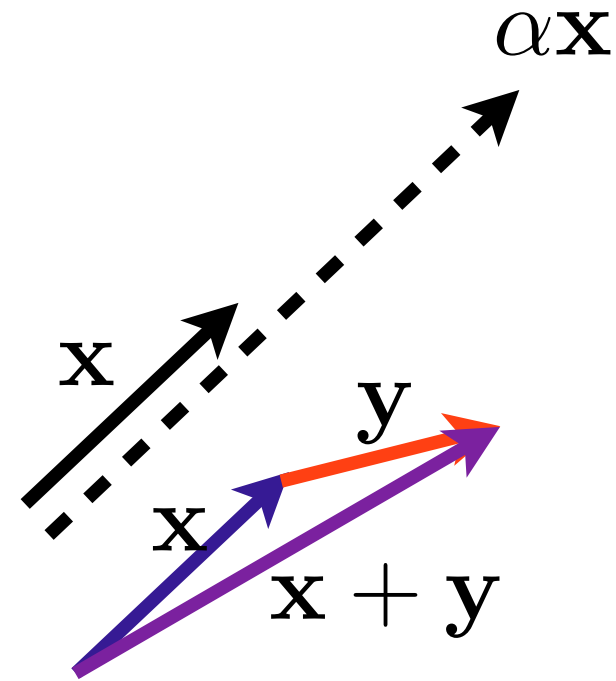
- Linear algebra pops up everywhere, even if you don't notice
- Statistics, data fitting, graph problems, PDE/coupled ODE solves...
- There exist very highly tuned packages for any sort of problem that can be cast into matrices and vectors - use them
- LAPACK, BLAS
- Exploit structure in your matrices
- Don't ever invert a matrix

# Outline

- Reminder of Linear Algebra
- Gaussian Elimination
- BLAS
- Solving  $Ax = b$
- Sparse matrices
- Iterative solvers
- Eigenproblems

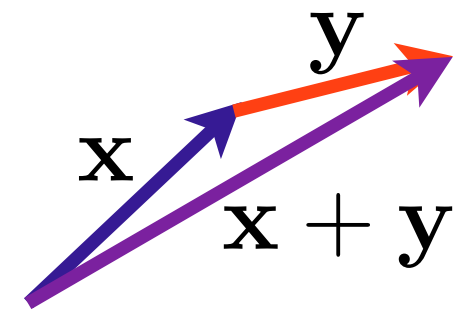
# Vector operations

- Geometric Interpretation
- Scaling a vector, adding two vectors together...
- Dot product (or any inner product)



# Vector spaces

- A set of vectors  $x$  spans a space  $S$  iff every vector in  $S$  can be expressed as a linear combination of  $x_i$



# Vector orthogonality - no overlap

- A set of vectors is said to be orthogonal if

$$x_i \cdot x_j \iff i \neq j$$

and orthonormal if

$$x_i \cdot x_j = \begin{cases} 0 & i \neq j \\ 1 & i = j \end{cases}$$

- A set of  $n$  orthogonal vectors necessarily span a subspace of dimension  $n$

# Matrix · Vector: Change of Basis

$$A\mathbf{x} = \mathbf{b}$$

$$\left[ \begin{array}{c|c|c|c} a_1 & a_2 & \dots & a_n \end{array} \right] \begin{pmatrix} x_1 \\ x_2 \\ \vdots \\ \vdots \\ x_n \end{pmatrix} = x_1 \begin{pmatrix} a_1 \end{pmatrix} + x_2 \begin{pmatrix} a_2 \end{pmatrix} + \dots + x_n \begin{pmatrix} a_n \end{pmatrix}$$



# Matrix · Vector: Change of Basis

- $Ax = b$ :  $x$  is the (unique) vector of coefficients that represents  $b$  in the basis of columns of  $A$
- Basis for  $b$ :  $\{e_1, e_2, \dots, e_n\}$
- Basis for  $x$ :  $\{a_1, a_2, \dots, a_n\}$

# Column spaces

- Column space of  $A$  - the space spanned by the column vectors  $a_i$
- eg, column space is all vectors that can be formed by linear combinations of the  $a_i$

# Matrix Vector: Linear mapping

- $Ax = b$  : Linear transformation of  $x$ .
  - $Ax_1 = b_1$  ;  $Ax_2 = b_2$
  - $A(x_1 + x_2) = (b_1 + b_2)$
  - $A(\alpha x_1) = \alpha b_1$

# Range of A - all possible b

- The range of a matrix A is the space of all possible vectors it can map to:

$$b \in \text{Range}(A) \implies \exists x \mid Ax = b$$

eg, column space.

# Nullspace of A: vectors that map to zero

- The nullspace of a matrix  $A$  is the space of all vectors it maps to zero:

$$\mathbf{x} \in \text{Null}(\mathbf{A}) \implies \mathbf{Ax} = \mathbf{0}, \mathbf{x} \neq \mathbf{0}$$

- For matrices  $A$  with a non-empty nullspace, there may be no solution to  $\mathbf{Ax}=\mathbf{b}$ , or infinitely many solutions.

# Column Rank: Dimension of Range

- The Rank of a matrix  $A$  is the dimension (eg, minimum number of basis vectors) of its column space.
- For square ( $n \times n$ ) matrix, a Full-Rank matrix has rank  $n$ .
- Column rank = Row Rank (not obvious, but true.) So generally just say “Rank”

# Rank + Nullity

- Rank of Matrix
- + Nullity (rank of nullspace) of matrix
- = # of columns of matrix

# Invertability

- Square, full-rank  $n \times n$  matrix  $A$  has an inverse,  $A^{-1}$ , such that  $AA^{-1} = A^{-1}A = I$
- For  $n \times n$  matrix, following statements are equivalent:
  - Has an inverse
  - $\text{rank}(A) = n$
  - $\text{range}(A) = \mathbb{R}^n$
  - $\text{null}(A) = \{\}$
  - No eigenvalues are 0
  - No singular values are 0
  - determinant is non-zero



# Solving Linear Systems

$Ax=b$ , solve for  $x$

# Sets of linear equations: don't invert

- $Ax = b$  implies  $x = A^{-1}b$
- Mathematically true, but numerically, inversion:
  - is slower than other solution methods
  - is numerically much less stable
  - ruins sparsity (**huge** memory disadvantage for, eg, PDEs on meshes)
  - loses any special structure of matrix  $A$

# Easy systems to solve

- We'll talk about methods to solve linear systems of equations
- Will assume nonsingular matrices (so there exists a unique solution)
- But some systems much easier to solve than others. Be aware of “nice” properties of your matrices!

# Diagonal Matrices

- (generally called  $D$ , or  $\Lambda$ )
- Ridiculously easy
- Matrix multiplication - just  $d_i x_i$

$$\begin{pmatrix} d_1 & & & \\ & d_2 & & \\ & & \dots & \\ & & & d_n \end{pmatrix} \begin{pmatrix} x_1 \\ x_2 \\ \vdots \\ x_n \end{pmatrix} = \begin{pmatrix} b_1 \\ b_2 \\ \vdots \\ b_n \end{pmatrix}$$

$$x_i = \frac{b_i}{d_i}$$

# Upper Triangular Matrices

- Generally called U
- “Back Substitution”: solve (easy) last one first
- Use that to solve previous one, etc.
- Lower triangular (L): “Forward substitution”, same deal.

$$\begin{pmatrix} u_{1,1} & u_{1,2} & \cdots & u_{1,n} \\ & u_{2,2} & \cdots & u_{2,n} \\ & & \cdots & \vdots \\ & & & u_{n,n} \end{pmatrix} \begin{pmatrix} x_1 \\ x_2 \\ \vdots \\ x_n \end{pmatrix} = \begin{pmatrix} b_1 \\ b_2 \\ \vdots \\ b_n \end{pmatrix}$$

$$x_n = \frac{b_n}{u_{n,n}}$$
$$x_{n-1} = \frac{b_n - u_{n-1,n}x_n}{u_{n-1,n-1}}$$
$$\vdots$$

# Orthogonal matrices

- Generally called  $Q$
- Columns (rows) are orthogonal unit vectors
- Transpose is inverse!
- *That* inverse I'll let you compute.
- Orthogonal matrices are numerically very nice - all row, col vectors are same "length".

$$Q^T Q = I$$

$$Q\mathbf{x} = \mathbf{b}$$

$$Q^T Q\mathbf{x} = Q^T \mathbf{b}$$

$$\mathbf{x} = Q^T \mathbf{b}$$

# Symmetric Matrices

- No special nomenclature
- Half the work; only have to deal with half the matrix
- (I'm assuming real matrices, here; complex: Hermetian)

$$A^T = A$$

$$a_{i,j} = a_{j,i}$$

# Symmetric Positive Definite

- Very special but common (covariance matrices, some PDEs)
- Always non-singular
- All eigenvalues positive
- Numerically very nice to work with

$$A^T = A$$

$$\mathbf{x}^T A \mathbf{x} > 0$$

$$A = LL^T$$



# Structure matters

- Find structure in your problems
- If writing equations in slightly different way gives you nice structure, do it
- Preserve structure when possible

# Gaussian Elimination

- For general square matrices (can't exploit above properties)
- We all learned this in high school:
- Subtract off multiples of previous rows to zero out below-diagonals
- Back-substitute when done

$$\begin{pmatrix} 10 & -7 & 0 \\ 5 & -1 & 5 \\ -2 & 2 & 6 \end{pmatrix} \begin{pmatrix} x_1 \\ x_2 \\ x_3 \end{pmatrix} = \begin{pmatrix} 7 \\ 6 \\ 4 \end{pmatrix}$$

$$\begin{pmatrix} 10 & -7 & 0 \\ & 2.5 & 5 \\ & 3.4 & 6 \end{pmatrix} \begin{pmatrix} x_1 \\ x_2 \\ x_3 \end{pmatrix} = \begin{pmatrix} 7 \\ -0.5 \\ 2.6 \end{pmatrix}$$

$$\begin{pmatrix} 10 & -7 & 0 \\ & 2.5 & 5 \\ & & -0.8 \end{pmatrix} \begin{pmatrix} x_1 \\ x_2 \\ x_3 \end{pmatrix} = \begin{pmatrix} 7 \\ -0.5 \\ 3.28 \end{pmatrix}$$

# Basic Linear Algebra Subroutines

- Linear algebra fairly simple: matrices and vectors
- Row vector operations, column vector operations, matrix-matrix operations
- BLAS: Basic Linear Algebra Subroutines.
  - Level 1: vector-vector operations
  - Level 2: matrix-vector operations
  - Level 3: matrix-matrix operations

# Basic Linear Algebra Subroutines

- A well defined standard interface for these routines
- Many highly-tuned implementations exist for various platforms. (Atlas, Flame, Goto, PLASMA, cuBLAS...)
- (Interface vs. Implementation! Trick is designing a sufficiently general interface.)
- Higher-order operations (matrix factorizations, like as we'll see, gaussian elimination) defined in LAPACK, on top of BLAS.

# Typical BLAS routines

- Level 1: sdot (dot product, single), zaxpy ( $a\mathbf{x} + \mathbf{y}$ , dbl complex)
- Level 2: dgemv (dbl matrix\*vec), dsymv (dbl symmetric matrix\*vec)
- Level 3: sgemm (general matrix-matrix), ctrmm (triangular matrix-matrix)
- Incredibly cryptic names, interfaces.

## Prefixes:

S: Single C: Complex  
D: Double Z: Double Complex

## Matrix Types:

GE: General	SY: Symmetric
GB: General Banded	SB: Symmetric Banded
HY: Hermetian	HB: Hermetian Banded
TR: Triangular	TB: Triangular Banded
TP: Triangular Packed	

# Why bother?

- Finding, downloading library
- Figuring out how to link
- C/Fortran issues
- Just write it - it's not rocket science.

$$C = AB$$

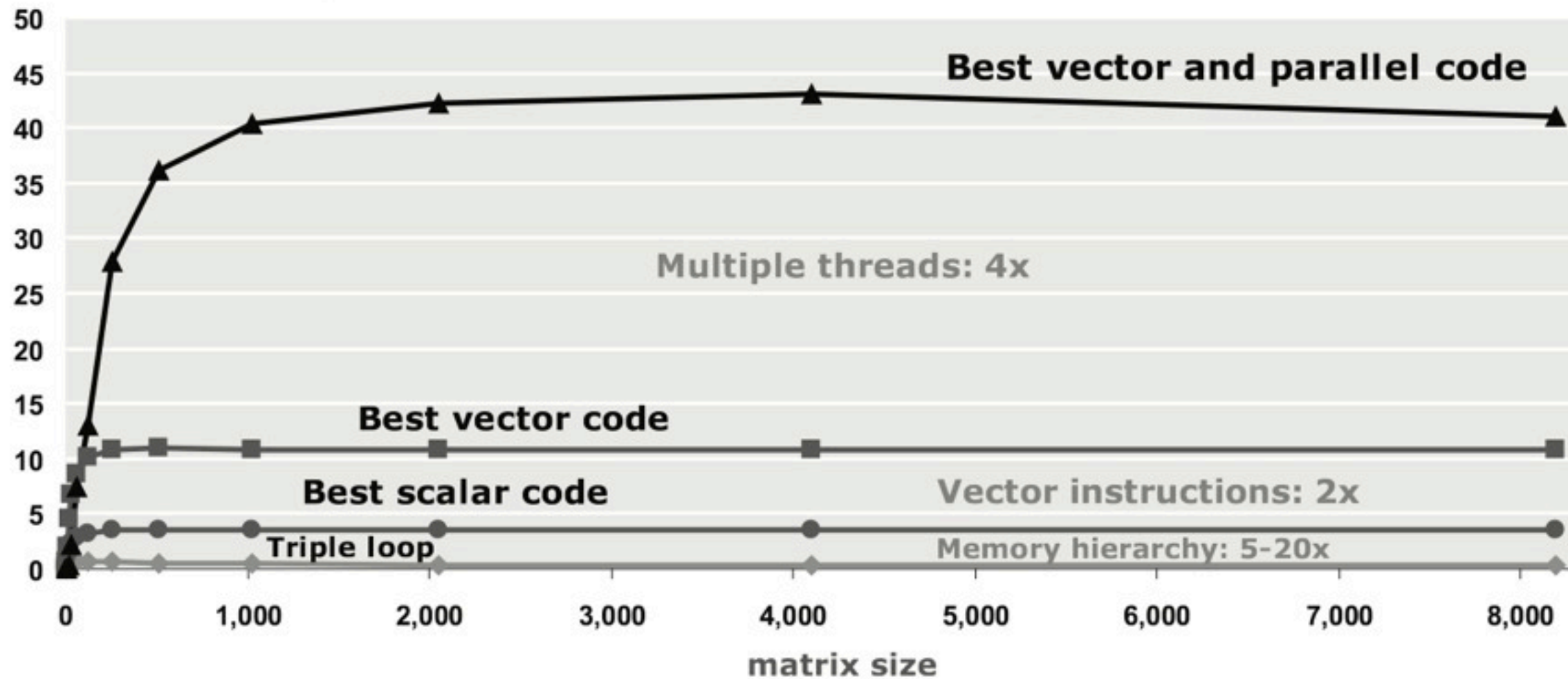
$$c_{i,j} = \sum_k a_{i,k} b_{k,j}$$

```
for (i=0; i<N; i++)  
  for (j=0; j<N; j++)  
    for (k=0; k<N; k++)  
      c[i][j] = a[i][k]*b[k][j];
```

# Never, ever, write your own

## Matrix-Matrix Multiplication (MMM) on 2 x Core 2 Extreme 3 GHz

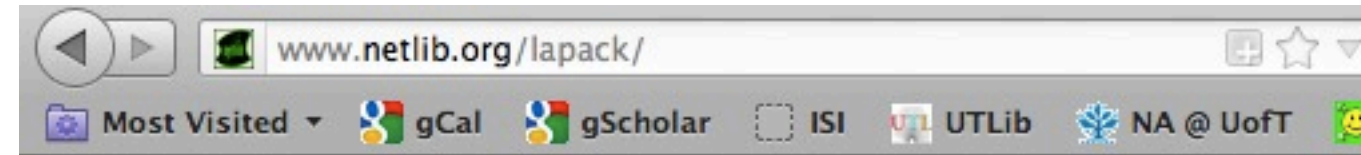
Performance [Gflop/s]



“How to Write Fast Numerical Code: A Small Introduction”, Chellappa et al  
[www.ece.cmu.edu/~franzf/papers/gttse07.pdf](http://www.ece.cmu.edu/~franzf/papers/gttse07.pdf)

# Division of Labour

- Focus on the science you need to do
- Write code for your problem - stuff that you know best
- Let people who enjoy making fast linear algebra software for a living do that.



## LAPACK — Linear Algebra PA

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# Gaussian Elimination = LU Decomposition

- With each stage of the elimination, we were subtracting off some multiple of a previous row
- That means the factored U can have the same multiple of the row added to it to get back to A
- Decomposing to give us  $A = LU$

$$\begin{pmatrix} 10 & -7 & 0 \\ 5 & -1 & 5 \\ -2 & 2 & 6 \end{pmatrix} = \begin{pmatrix} 1 & & \\ & 1 & \\ & & 1 \end{pmatrix} \begin{pmatrix} 10 & -7 & 0 \\ 5 & -1 & 5 \\ -2 & 2 & 6 \end{pmatrix}$$

$$\begin{pmatrix} 10 & -7 & 0 \\ 5 & -1 & 5 \\ -2 & 2 & 6 \end{pmatrix} = \begin{pmatrix} 1 & & \\ +\frac{1}{2} & 1 & \\ -\frac{1}{5} & & 1 \end{pmatrix} \begin{pmatrix} 10 & -7 & 0 \\ & 2.5 & 5 \\ & 0.6 & 6 \end{pmatrix}$$

$$\begin{pmatrix} 10 & -7 & 0 \\ 5 & -1 & 5 \\ -2 & 2 & 6 \end{pmatrix} = \begin{pmatrix} 1 & & \\ -\frac{1}{2} & 1 & \\ +\frac{1}{5} & +\frac{6}{25} & 1 \end{pmatrix} \begin{pmatrix} 10 & -7 & 0 \\ & 2.5 & 5 \\ & & 4.8 \end{pmatrix}$$

$$A = LU$$

# Solving is fast with LU

- Once have  $A = LU$  ( $O(n^3)$  steps) can solve for  $x$  quickly ( $O(n^2)$  steps)
- Can solve for same  $A$  with different  $b$  very cheaply
- Backsubstitute, then forward substitute

$$A\mathbf{x} = \mathbf{b}$$

$$LU\mathbf{x} = \mathbf{b}$$

$$L(\mathbf{y}) = \mathbf{b}$$

$$\mathbf{y} = \text{Backsubst}(L, \mathbf{b})$$

$$U\mathbf{x} = \mathbf{y}$$

$$\mathbf{x} = \text{Forwardsubst}(U, \mathbf{y})$$

# Conditioning

- A problem is said to be inherently ill-conditioned if any small perturbation in the initial conditions generates huge changes in the results

- Say, calculating  $f(x)$ : if

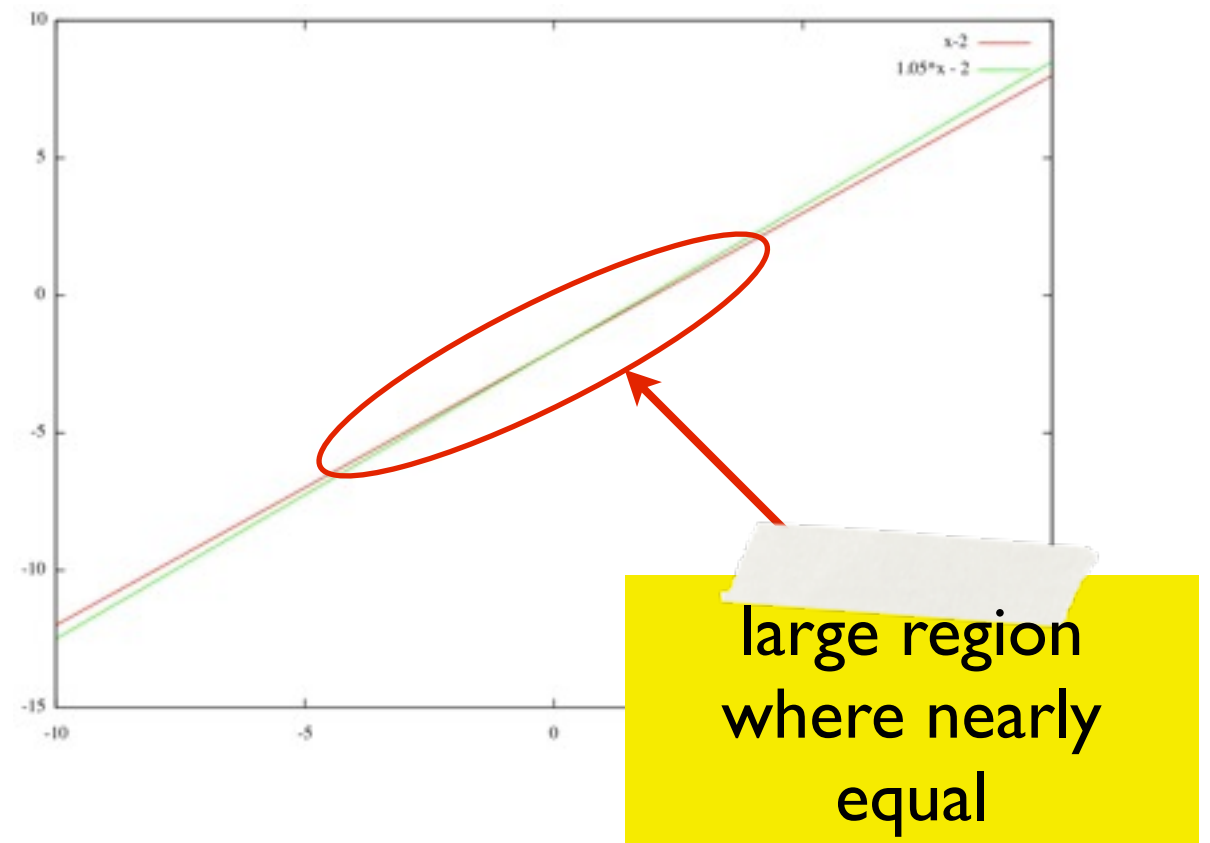
$$\frac{\|f(x + \delta x)\|}{\|f(x)\|} \gg \frac{\|\delta x\|}{\|x\|}$$

then the problem is inherently hard to do numerically  
(or with any sort of approximate method)

# Conditioning

- In matrix problems, this can happen in nearly singular matrices - nearly linearly dependant columns.
- Carve out strongly overlapping subspaces
- Very small changes in b (say) can result in hugely different change in x

$$\begin{pmatrix} 1 & 1 \\ 1 & 1.05 \end{pmatrix} \begin{pmatrix} x \\ y \end{pmatrix} = \begin{pmatrix} 2 \\ 2 \end{pmatrix}$$



# Try it

- Order unity change in answer with 1/2 part in  $10^{-4}$  change in input.
- Would be true with infinite-precision arithmetic.
- Inherently a tough problem.

```
$ ipython --pylab

In [1]: a = numpy.array([[1,1],
                        [1,1.0001]])

In [2]: b = numpy.array([2,2])

In [3]: scipy.linalg.solve(a,b)
Out[3]: array([ 2.,  0.])

In [4]: scipy.linalg.solve(a,
                        b+numpy.array([0,0.0001]))
Out[4]: ??
```

# Condition number

- Condition number can be estimated using “sizes” (matrix norms) of  $A$ , inverse of  $A$ .
- Lapack routines exist:     CON
- Relative error in  $x$  can't be less than condition number \* machine epsilon.

$$\kappa(A) = \|A\| \cdot \|A^{-1}\|$$

$$\frac{\|\delta x\|}{\|x\|} < \kappa(A) \frac{\|\delta b\|}{\|b\|}$$

# Residuals

- Computational scientists have over 20 words for “numerical error”
- Absolute, relative error - error in  $x$ .
- **Residual**: answer in result provided by erroneous  $x$  - error in  $b$ .
- Which is more important is entirely problem dependant

# Pivoting

- The diagonal elements we use to “zero out” lower elements are called pivots.
- May need to change pivots, if for instance zeros appear in wrong place
- Matrix might be singular, or fixed by reordering
- PLU factorization

$$A = \begin{pmatrix} 0 & a & b \\ 0 & 0 & c \\ d & e & f \end{pmatrix}$$



# Pivoting

- Important numerically, too - avoid catastrophic loss of precision.
- Consider 3 digits of decimal precision. Problem nowhere near singular
- What does scipy say?

$$\begin{pmatrix} 10^{-4} & 1 \\ 1 & 1 \end{pmatrix} \begin{pmatrix} x \\ y \end{pmatrix} = \begin{pmatrix} 1 \\ 2 \end{pmatrix}$$
$$\begin{pmatrix} 10^{-4} & 1 \\ 1 + 10^4 & 10^4 \end{pmatrix} \begin{pmatrix} x \\ y \end{pmatrix} = \begin{pmatrix} 1 \\ 2 + 10^4 \end{pmatrix}$$
$$\begin{pmatrix} 10^{-4} & 1 \\ & 10^4 \end{pmatrix} \begin{pmatrix} x \\ y \end{pmatrix} = \begin{pmatrix} 1 \\ 10^4 \end{pmatrix}$$
$$\begin{pmatrix} x \\ y \end{pmatrix} = \begin{pmatrix} 0 \\ 1 \end{pmatrix}$$

# Residuals

- Good linear algebra algorithms (and implementations) should give residuals no more than (some function of size of matrix)  $\times$  (machine epsilon)
- And errors in  $x$  no more than condition number times that.
- An exact solution to a nearby problem
- Bad algorithms/implementations will depend on  $\sqrt{\text{machine epsilon}}$  or worse, and/or will be matrix dependant (eg, LU without pivoting).

# Cholesky Factorization

- For symmetric, positive definite matrices (surprisingly common), use Cholesky factorization instead.
- $A = LL^T$
- No pivoting; more numerically stable; faster.

```
In [10]: a =  
numpy.array([[25,15,-5],  
[15,18,0],[-5,0,11]])
```

```
In [11]:  
scipy.linalg.cholesky(a)  
Out[11]:  
array([[ 5.,  3., -1.],  
       [ 0.,  3.,  1.],  
       [ 0.,  0.,  3.]])
```

# $Ax \sim b$ : QR factorizations

- Not all  $Ax=b$  s can be solved; consider an overdetermined system (data fitting).
- LU won't even work on non-square systems.
- What to do?

$$\begin{pmatrix} x_0^3 & x_0^2 & x_0 & 1 \\ x_1^3 & x_1^2 & x_1 & 1 \\ \dots & & & \\ x_n^3 & x_n^2 & x_n & 1 \end{pmatrix} \begin{pmatrix} a \\ b \\ c \\ d \end{pmatrix} = \begin{pmatrix} y_0 \\ y_1 \\ \dots \\ y_n \end{pmatrix}$$

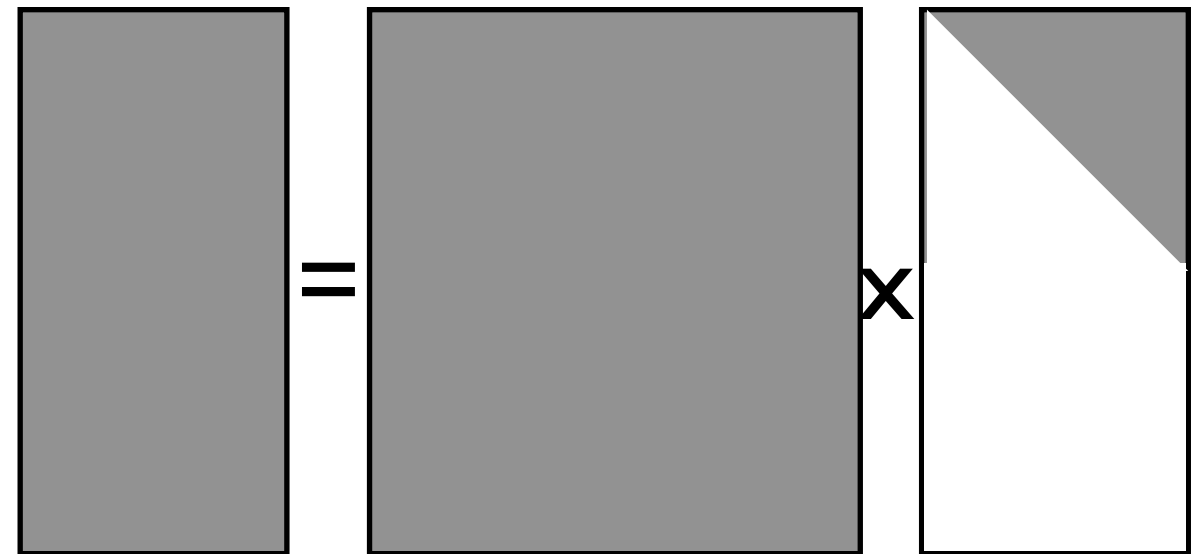
# Minimize residual: Residual not in Range(A)

- Want to project out residual somehow
- Normal equations
- Much of linear algebra is decompositions into useful forms

$$\begin{aligned}r^2 &= \|\mathbf{b} - A\mathbf{x}\|_2^2 \\&= (\mathbf{b} - A\mathbf{x})^T (\mathbf{b} - A\mathbf{x}) \\&= \mathbf{b} \cdot \mathbf{b} - 2\mathbf{b}^T A\mathbf{x} + \mathbf{x}^T A^T A\mathbf{x} \\0 &= -2\mathbf{b}^T A + 2\mathbf{x}^T A^T A \\(A^T A)\mathbf{x} &= A^T \mathbf{b}\end{aligned}$$

# QR decomposition

- All matrices can be decomposed into QR, even  $m \times n$ ,  $m > n$
- Bottom half of R is necessarily empty (below diagonal)
- All columns in Q are orthogonal bases of  $m$ -d space, and R is the combination of them that makes up A



# Orthogonalizing columns of A

- Let's take these n column vectors of length m and make an orthonormal basis.
- Divide  $a_1$  by its norm; I done. What about rest?

$$\left[ \begin{array}{c|c|c|c} a_1 & a_2 & \dots & a_n \end{array} \right]$$

$$\left[ \begin{array}{c} q_1 \end{array} \right] = \frac{\mathbf{a}_1}{\|\mathbf{a}_1\|}$$

# Gram-Schmidt (don't use this)

- Easiest to follow at first isn't numerically stable (should use Householder transforms).
- Subtract off  $q_1$  component from  $a_2$ , take unit vector of that -  $q_2$ .
- And so on.
- Bit like LU, but instead of making zeros, you're making orthogonality

$$\left[ \begin{array}{c|c|c|c} a_1 & a_2 & \dots & a_n \end{array} \right]$$

$$\left[ \begin{array}{c} q_2 \end{array} \right] = \frac{\mathbf{a}_2 - (\mathbf{a}_2 \cdot \mathbf{q}_1) \mathbf{q}_1}{\|\mathbf{a}_2 - \mathbf{a}_2 \cdot \mathbf{q}_1\|}$$



# Gram-Schmidt (don't use this)

- Gram-Schmidt handy for generating orthogonal series of basis functions from (say) polynomials, as well.
- Same procedure, just different definition of inner product, norm.

$$\left[ \begin{array}{c|c|c|c} a_1 & a_2 & \dots & a_n \end{array} \right]$$

$$\left[ \begin{array}{c} q_2 \end{array} \right] = \frac{\mathbf{a}_2 - (\mathbf{a}_2 \cdot \mathbf{q}_1) \mathbf{q}_1}{\|\mathbf{a}_2 - \mathbf{a}_2 \cdot \mathbf{q}_1\|}$$

# QR Factor a random matrix

```
In [13]: r = numpy.random.random((50,50))
```

```
In [14]: for i in xrange(50):
.....:     for j in xrange(i):
.....:         r[i,j] = 0.
.....:
```

```
In [15]: print r[0:3,0:3]
[[ 0.4147775  0.64843642  0.41133882]
 [ 0.         0.88592831  0.54711704]
 [ 0.         0.         0.23438925]]
```

```
In [16]: q,x = scipy.linalg.qr(numpy.random.random((50,50)))
```

```
In [17]: a = numpy.dot(q,r)
```

```
In [18]: q2,r2 = scipy.linalg.qr(a)
```

```
In [19]: a2 = numpy.dot(q2,r2)
```

```
In [20]: print scipy.linalg.norm(a2-a)/scipy.linalg.norm(a)
6.60894445883e-16
```

# Errors and residuals

- Generate random matrices Q,R; calculate A
- QR factorization of A
- Errors in Q2, R2  $\sim \sqrt{\text{machine epsilon}}$
- (Random matrix tends to be ill-conditioned)
- Residual in A: (machine epsilon). Would be  $\sqrt{\text{machine epsilon}}$  with classical G-S

```
In [18]: q2,r2 = scipy.linalg.qr(a)
In [19]: a2 = numpy.dot(q2,r2)
In [20]: print scipy.linalg.norm(a2-a)/
          scipy.linalg.norm(a)
6.60894445883e-16
In [21]: print scipy.linalg.norm(q2-q)/
          scipy.linalg.norm(q)
3.67030163525e-07
In [22]: print scipy.linalg.norm(r2-r)/
          scipy.linalg.norm(r)
6.36755093518e-08
```

# Normal equations with QR are easy

- Now this is fairly straightforward
- End up with  $(R\mathbf{x})$  -- forward solve -- equal to matrix-vector product.
- Done!

$$(A^T A)\mathbf{x} = A^T \mathbf{b}$$

$$R^T Q^T Q R \mathbf{x} = R^T Q^T \mathbf{b}$$

$$R^T R \mathbf{x} = R^T Q^T \mathbf{b}$$

$$R \mathbf{x} = Q^T \mathbf{b}$$

# Eigenproblems

- Tells a great deal about the structure of a matrix
- How it will act on a vector: project onto its eigenvectors, multiply by eigenvalues.
- Goal is a complete decomposition:

$$A\mathbf{x} = \lambda\mathbf{x}$$

$$A \begin{bmatrix} | & | & \dots & | \\ x_1 & x_2 & \dots & x_n \\ | & | & \dots & | \end{bmatrix} = \begin{bmatrix} | & | & \dots & | \\ x_1 & x_2 & \dots & x_n \\ | & | & \dots & | \end{bmatrix} \begin{bmatrix} \lambda & & & \\ & \lambda & & \\ & & \ddots & \\ & & & \lambda \end{bmatrix}$$

# Eigenvalue Decomposition

- For square matrix
- “Similarity Transform”
- No restrictions on the structure of  $X$
- Can only happen if there are a full set of eigenvectors.
- Diagonalizability:  $N$  non-null eigenvectors;
- Invertability:  $N$  non-zero eigenvalues

$$A \begin{bmatrix} | & | & & | \\ x_1 & x_2 & \dots & x_n \\ | & | & & | \end{bmatrix} = \begin{bmatrix} | & | & & | \\ x_1 & x_2 & \dots & x_n \\ | & | & & | \end{bmatrix} \begin{bmatrix} \lambda & & & \\ & \lambda & & \\ & & \dots & \\ & & & \lambda \end{bmatrix}$$

$$AX = X\Lambda$$

$$A = X\Lambda X^{-1}$$

# Defective Matrices

- Both these matrices have eigenvalue 2, with multiplicity 3
- But A has full set of eigenvectors ( $e_1, e_2, e_3$ )
- B has only one eigenvector;  $e_1$
- Not diagonalizable

$$A = \begin{pmatrix} 2 & & \\ & 2 & \\ & & 2 \end{pmatrix}$$
$$B = \begin{pmatrix} 2 & 1 & \\ & 2 & 1 \\ & & 2 \end{pmatrix}$$

# Iterative Methods

- So far, have dealt solely with direct methods.
- Solution takes one (long) step, then answer is complete, as exact as matrix/method allows.
- Other approach; take successive approximations, get closer.
- Typically converge to machine accuracy in much less time than direct, esp for large matrices



# Krylov Subspaces

- Krylov subspace:  
repeated action on  $\mathbf{b}$  by  
 $A$ .

$$A\mathbf{x} = \mathbf{b}$$

- For sufficiently large  $n$ ,  
final term should  
converge to eigenvector  
with largest eigenvalue

$$\mathcal{K} = [\mathbf{b}, A\mathbf{b}, A^2\mathbf{b}, \dots, A^{n-1}\mathbf{b}]$$

- But slow, and only one  
eigenvalue?

# Krylov Subspaces

- Can orthogonalize (Gram Schmidt, Householder) to project out other components
- Should give approximations to eigenvectors (random  $\mathbf{b}$ )
- But not numerically stable

$$A\mathbf{x} = \mathbf{b}$$

$$\mathcal{K} = [\mathbf{b}, A\mathbf{b}, A^2\mathbf{b}, \dots, A^{n-1}\mathbf{b}]$$

# Arnoldi Iteration

- Stabilized orthogonalization
- Becomes Lanczos iteration for symmetric  $A$
- Orthogonal projection of  $A$  onto the Krylov subspace,  $H$
- $H$  is of modest size, can have eigenvalues calculated
- Note: Only requires matrix-vector, vector-vector products
- GMRES: Arnoldi iteration for solving  $Ax=b$

$$q_1 \leftarrow e_1$$

for  $j \in [1, k-1]$  :

$$h_{j,k-1} \leftarrow q_j^T q_k$$

$$q_k \leftarrow q_k - h_{j,k-1} q_j$$

$$h_{k,k-1} \leftarrow \|q_k\|$$

$$q_k \leftarrow \frac{q_k}{h_{k,k-1}}$$

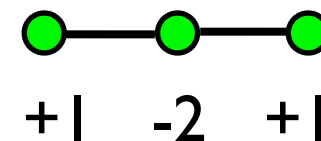
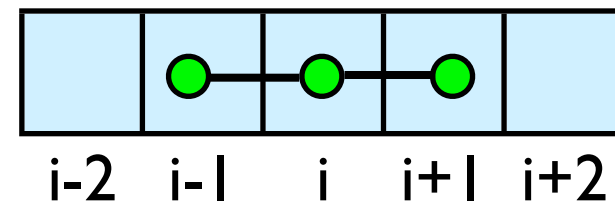
# Sparse Matrices

- So far, we've been assuming our matrices are dense; there are numbers stored for every entry in matrix.
- This is indeed often the case, but it's also often that huge numbers of the entries are zero: some roughly constant number of entries per row, much less than  $n$ .
- Difference between  $n^2$  and  $n$  can be huge if  $n \sim 10^6$ ; difference between doing and not doing the problem.
- Happens particularly often in discretizing PDEs.

# Discretizing Derivatives

- Done by finite differencing the discretized values
- Implicitly or explicitly involves interpolating data and taking derivative of the interpolant

$$\left. \frac{d^2 q}{dx^2} \right|_i \approx \frac{q_{i+1} - 2q_i + q_{i-1}}{\Delta x^2}$$



$$\frac{d\mathbf{q}}{dt} = \sigma \begin{pmatrix} -2 & 1 & & & \\ & 1 & -2 & 1 & \\ & & \dots & & \\ & & & 1 & -2 & 1 \\ & & & & 1 & -2 \end{pmatrix} \mathbf{q}$$

$$\frac{d\mathbf{q}}{dt} \approx \frac{\mathbf{q}^{n+1} - \mathbf{q}^n}{\Delta t}$$

$$\mathbf{q}^{n+1} = \mathbf{q}^n + \sigma \Delta t \mathbf{A} \mathbf{q}^n$$

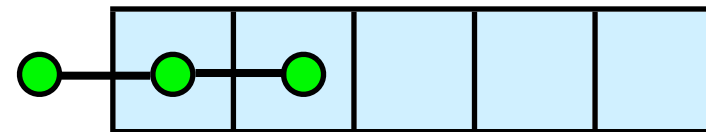
$$\mathbf{q}^{n+1} = (\mathbf{I} + \sigma \Delta t \mathbf{A}) \mathbf{q}^n$$

$$\mathbf{q}^{n+1} = \sigma \Delta t \begin{pmatrix} \frac{1}{\sigma \Delta t} - 2 & 1 & & & \\ & 1 & \frac{1}{\sigma \Delta t} - 2 & 1 & \\ & & \dots & & \\ & & & 1 & \frac{1}{\sigma \Delta t} - 2 & 1 \\ & & & & 1 & \frac{1}{\sigma \Delta t} - 2 \end{pmatrix} \mathbf{q}^n$$

# Boundary Conditions

- What happens when stencil goes off of the end of the box?
- Depends on how you want to handle boundary conditions.
- Typically easiest to have extra points on end, set values to enforce desired BCs.

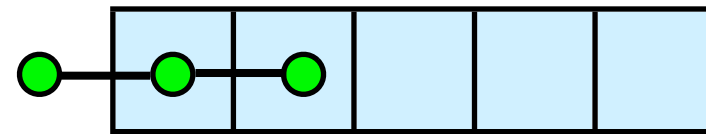
$$\left. \frac{d^2 q}{dx^2} \right|_i \approx \frac{q_{i+1} - 2q_i + q_{i-1}}{\Delta x^2}$$



# Boundary Conditions

- Dirichlet (fixed value) boundary conditions: just have 1 on diagonal, 0 elsewhere, keeps value there constant.
- Neumann (derivative) bcs: requires more manipulation of the equations.

$$\left. \frac{d^2 q}{dx^2} \right|_i \approx \frac{q_{i+1} - 2q_i + q_{i-1}}{\Delta x^2}$$





# Inverses destroy sparsity

- For sparse matrices like above, LU decompositions may maintain much sparsity (particularly if banded, etc)

$$\begin{pmatrix} 1 & -1 & & & \\ -1 & 2 & -1 & & \\ & -1 & 2 & -1 & \\ & & -1 & 2 & -1 \\ & & & -1 & 2 \end{pmatrix} = \begin{pmatrix} 1 & & & & \\ -1 & 1 & & & \\ & -1 & 1 & & \\ & & -1 & 1 & \\ & & & -1 & 1 \end{pmatrix} \begin{pmatrix} 1 & -1 & & & \\ & 1 & -1 & & \\ & & 1 & -1 & \\ & & & 1 & -1 \\ & & & & 1 \end{pmatrix}$$

$$\begin{pmatrix} 1 & -1 & & & \\ -1 & 2 & -1 & & \\ & -1 & 2 & -1 & \\ & & -1 & 2 & -1 \\ & & & -1 & 2 \end{pmatrix}^{-1} = \begin{pmatrix} 5 & 4 & 3 & 2 & 1. \\ 4 & 4 & 3 & 2 & 1. \\ 3 & 3 & 3 & 2 & 1. \\ 2 & 2 & 2 & 2 & 1. \\ 1 & 1 & 1 & 1 & 1. \end{pmatrix}$$

- Inverses in general are full
- For large n, difference between cn and n<sup>2</sup> huge.

# Sparse (banded) LU

- If entries only exist within a narrow band around diagonal, then row, column operations fast.

$$\begin{pmatrix} 1 & -1 & & & \\ -1 & 2 & -1 & & \\ & -1 & 2 & -1 & \\ & & -1 & 2 & -1 \\ & & & -1 & 2 \end{pmatrix} =$$

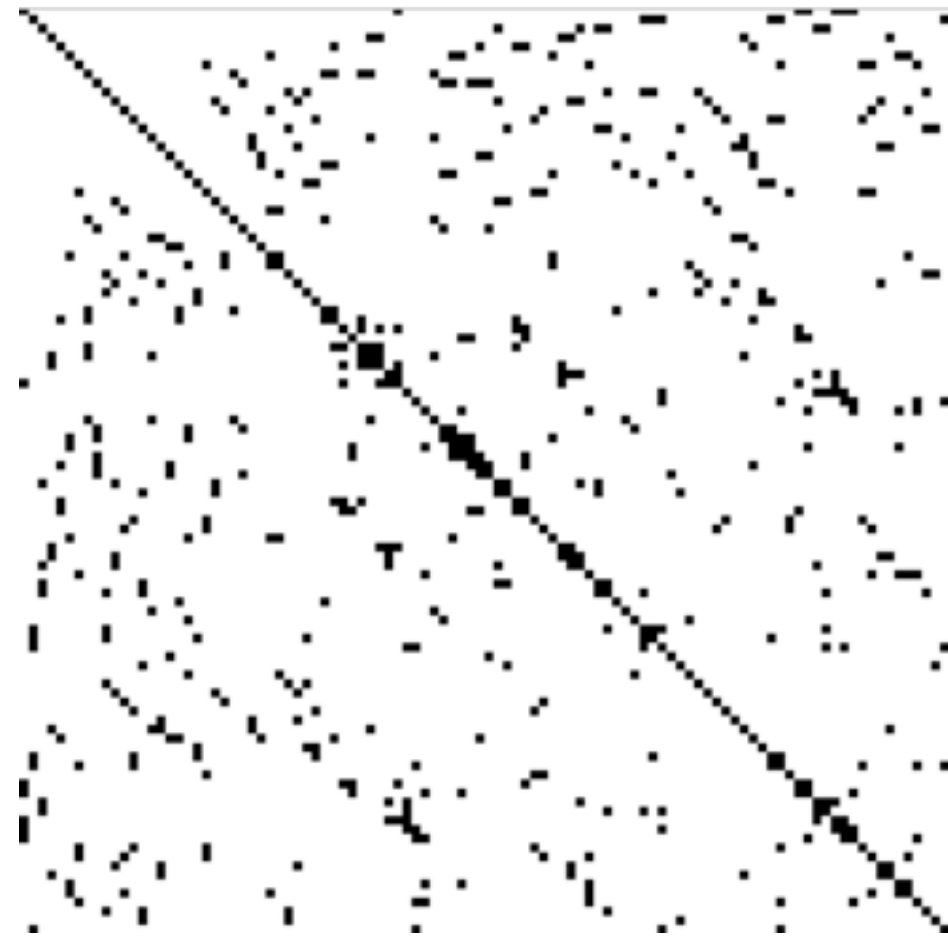
- May get significant “fill in” depending on exact structure of matrix

$$\begin{pmatrix} 1 & & & & \\ -1 & 1 & & & \\ & -1 & 1 & & \\ & & -1 & 1 & \\ & & & -1 & 1 \end{pmatrix} \begin{pmatrix} 1 & -1 & -1 & & \\ & 1 & 1 & -1 & \\ & & 1 & 1 & -1 \\ & & & 1 & -1 \\ & & & & 1 \end{pmatrix}$$

- (This is artificially good example)

# Sparsity patterns

- Sparse matrices can have arbitrary sparsity patterns
- Typically need at less than 10% nonzeros to make dealing with sparse matrices worth it.
- Half zeros - typically just store full matrix.



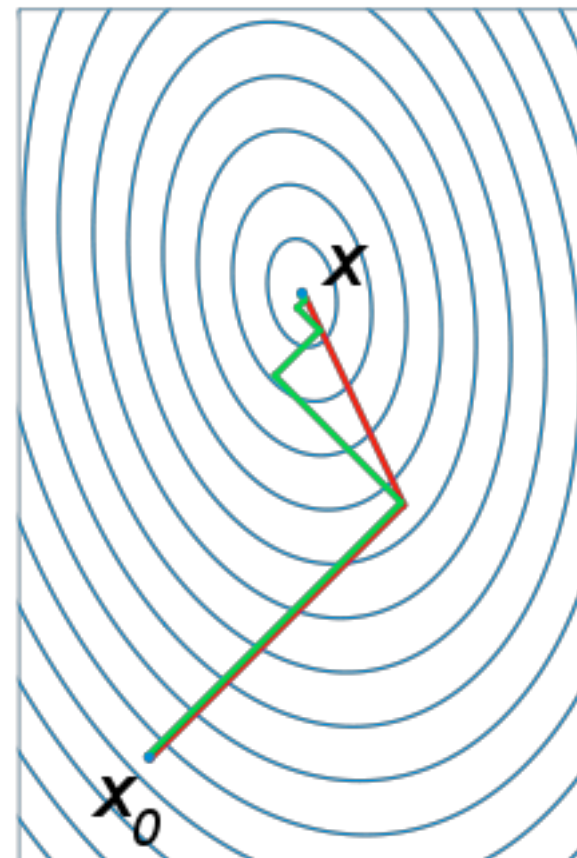
[http://en.wikipedia.org/wiki/File:Finite\\_element\\_sparse\\_matrix.png](http://en.wikipedia.org/wiki/File:Finite_element_sparse_matrix.png)

# Common Sparse Matrix Formats:

- CSR (Compressed Sparse Row): Just join all the nonzeros in rows together, with pointers to where each starts, and (similar sized) array of column for each value
- CSC (Compressed Sparse Column): Same, but flip row/column
- Banded: just store diagonals +/- some bandwidth
- Many many more.

# Iterative $Ax=b$ solvers: Conjugate Gradient

- SPD matrices, works particularly well on sparse systems
- “Steepest Descent”, but only on conjugate (w/r  $A$ ) directions: no “doubling back”



[http://en.wikipedia.org/wiki/Conjugate\\_gradient\\_method](http://en.wikipedia.org/wiki/Conjugate_gradient_method)

# Conjugate Gradient Method

$$\mathbf{r}_0 := \mathbf{b} - \mathbf{A}\mathbf{x}_0$$

$$\mathbf{p}_0 := \mathbf{r}_0$$

$$k := 0$$

repeat

$$\alpha_k := \frac{\mathbf{r}_k^T \mathbf{r}_k}{\mathbf{p}_k^T \mathbf{A} \mathbf{p}_k}$$

$$\mathbf{x}_{k+1} := \mathbf{x}_k + \alpha_k \mathbf{p}_k$$

$$\mathbf{r}_{k+1} := \mathbf{r}_k - \alpha_k \mathbf{A} \mathbf{p}_k$$

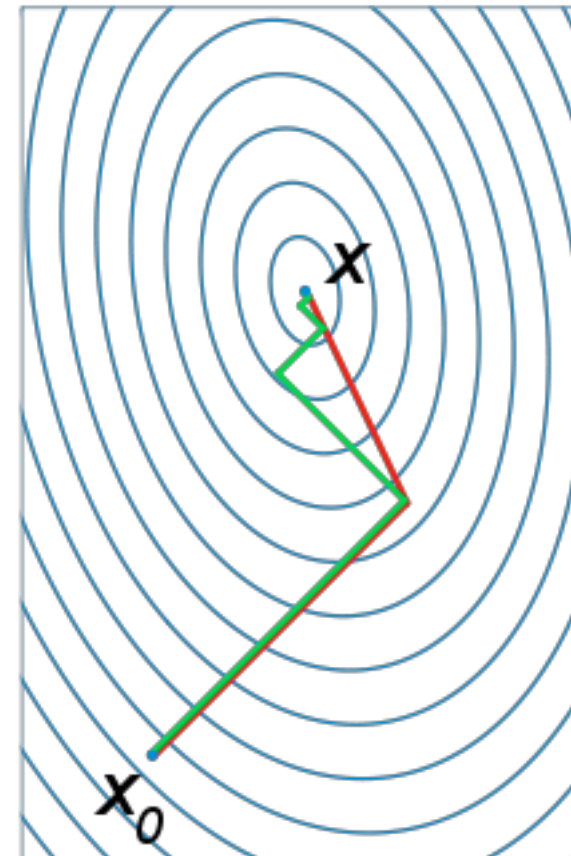
if  $\mathbf{r}_{k+1}$  is sufficiently small then exit loop end if

$$\beta_k := \frac{\mathbf{r}_{k+1}^T \mathbf{r}_{k+1}}{\mathbf{r}_k^T \mathbf{r}_k}$$

$$\mathbf{p}_{k+1} := \mathbf{r}_{k+1} + \beta_k \mathbf{p}_k$$

$$k := k + 1$$

end repeat



[http://en.wikipedia.org/wiki/Conjugate\\_gradient\\_method](http://en.wikipedia.org/wiki/Conjugate_gradient_method)

# Resources

- Trefethen & Bau, “Numerical Linear Algebra”  
<http://people.maths.ox.ac.uk/trefethen/text.html>
- Strang on iTunes U: “Mathematical Methods for Engineers” or “Linear Algebra” - excellent lectures by a master.

# Homework

Educational *and* fun.



# Homework: Part I

- The time-explicit formulation of the 1d heat diffusion equation has a term that looks like this (ignoring boundary conditions)

$$\frac{D\Delta t}{\Delta x^2} \begin{pmatrix} -2 & 1 & & & & & \\ 1 & -2 & 1 & & & & \\ & 1 & -2 & 1 & & & \\ & & & \dots & & & \\ & & & 1 & -2 & 1 & \\ & & & & 1 & -2 & \end{pmatrix} x^n$$

# Homework: Part I

- Ignoring the constants, what are the eigenvalues for this problem - what might we expect to get amplified/damped by this operator? (use 100 points; D\_\_EV)
- Plot the eigenmode with the largest and smallest absolute eigenvalues, and explain them.
- Use the largest abs. eigenvalue to put a constraint on  $dt$  given  $dx$ ,  $D$ . This is a stability constraint on the numerical method; for larger timesteps, method blows up.

# Lapack Hints

- If you are using an  $n \times n$  array, the “leading dimension” of the array is  $n$ . (This argument is so that you could work on sub-matrices if you wanted)
- Have to make sure the 2d array is contiguous block of memory
- C vs FORTRAN array orderings
- C bindings for LAPACK - lapacke

# Homework: Part 2

- For a 1d grid of size 100 (eg, a 100x100 matrix A), using lapack, evolve this PDE. Plot and explain results.
- Have an initial condition where  $x = 1$  at the first zone, and zero everywhere else (hot plate “turns on” in a cold domain).
- You’ll want to use driver routines for linear solves ( <http://www.netlib.org/lapack/lug/node26.html> ). Do solve in double precision (D\_\_SV). Which solver should you use?
- Using a small enough timestep, timestep the temperature evolution by finding dT. Do solution in double precision (D\_\_SV) .