```
vimrc
"Windows: :e $HOME/_vimrc
"Linux: :e $HOME/.vimrc
set nocompatible
set number
syntax on
filetype plugin indent on
set bs=indent,eol,start
set et
set sw=4
set ts=4
set hls
nnoremap j gj
nnoremap k gk
nnoremap tn :tabnew<Space>
nnoremap <C-1> gt
nnoremap <C-h> gT
nnoremap <C-m> :make<CR>
"set backspace=indent,eol,start
"set expandtab
"set shiftwidth=4
"set tabstop=4
"set hlsearch
    Algebra.cc
typedef signed long long int T;
```

return a:

```
// Throughout all following code, it's assumed that inputs are nonnegative.
// However, a signed type is used for two purposes:
// 1. -1 is used as an error code sometimes.
// 2. Some of these (egcd) actually have negative return values.
typedef vector<T> VT;
typedef vector<VT> VVT;
// basic acd
T gcd( T a, T b ) {
   if( a < 0 ) return gcd(-a,b);</pre>
    if( b < 0 ) return gcd(a,-b);</pre>
   while( b ) { c = a \% b; a = b; b = c; }
// basic lcm
T 1cm(Ta, Tb) {
    if (a < 0) return lcm(-a,b);
   if( b < 0 ) return lcm(a,-b);
    return a/gcd(a,b)*b; // avoids overflow
// returns qcd(a,b), and additionally finds x,y such that qcd(a,b) = ax + by
T egcd( T a, T b, T &x, T &y ) {
    if(a < 0) {
       T r = egcd(-a,b,x,y);
        x *= -1;
        return r;
    if( b < 0 ) {
        T r = egcd(a, -b, x, y);
        y *= -1;
       return r:
    T u = v = 0. v = x = 1:
    while(b) {
        T q = a/b,
                      r = a % b:
         a = b,
                      b = r;
        T m = u,
                      n = v;
         u = x - q*u, v = y - q*v;
         x = m
                      y = n;
```

```
// Compute b so that ab = 1 \pmod{n}.
// Returns n if gcd(a,n) != 1, since no such b exists.
T modinv( T a, T n ) {
    T x, y, g = egcd(a, n, x, y);
    if( g != 1 ) return -1;
    x %= n;
    if( x < 0 ) x += n;
    return x;
// Find all solutions to ax = b (mod n),
// and push them onto S.
// Returns the number of solutions.
// Solutions exist iff gcd(a,n) divides b.
// If solutions exist, then there are exactly gcd(a,n) of them.
size_t modsolve( T a, T b, T n, VT &S ) {
    T_1,_2, g = egcd(a,n,_1,_2); // modinv uses egcd already
    if((b\%g) == 0) {
       T x = modinv(a/g, n/g);
        x = (x * b/g) \% (n/g);
        for( T k = 0; k < g; k++)
            S.push_back( (x + k*(n/g)) \% n);
        return (size t)g:
    return 0;
// Chinese remainder theorem, simple version.
// Given a, b, n, m, find z which simultaneously satisfies
    z = a \pmod{m} and z = b \pmod{n}.
// This z, when it exists, is unique mod lcm(n,m).
// If such z does not exist, then return -1.
// z exists iff a == b (mod gcd(m,n))
T CRT( T a, T m, T b, T n ) {
    T s, t, g = egcd(m, n, s, t);
    T 1 = m/g*n, r = a \% g;
    if( (b \% g) != r ) return -1;
    if( g == 1 ) {
       s = s \% 1; if( s < 0 ) s += 1;
        t = t \% 1: if( t < 0 ) t += 1:
       T r1 = (s * b) % 1, r2 = (t * a) % 1;
          r1 = (r1 * m) % 1, r2 = (r2 * n) % 1;
        return (r1 + r2) % 1;
    else f
        return g*CRT(a/g, m/g, b/g, n/g) + r;
// Chinese remainder theorem, extended version.
// Given a[K] and n[K], find z so that, for every i,
     z = a[i] \pmod{n[i]}
// The solution is unique mod lcm(n[i]) when it exists.
// The existence criteria is just the extended version of what it is above.
T CRT_ext( const VT &a, const VT &n ) {
    T \text{ ret } = a[0], 1 = n[0];
    FOR(i 1 a size()) {
        ret = CRT( ret, 1, a[i], n[i]);
        1 = 1 cm(1 n[i])
        if( ret == -1 ) return -1:
    return ret;
// Compute x and y so that ax + by = c.
// The solution, when it exists, is unique up to the transformation
    x \rightarrow x + kb/q
// y \rightarrow y - ka/q
// for integers k, where g = gcd(a,b).
// The solution exists iff gcd(a,b) divides c.
// The return value is true iff the solution exists.
bool linear_diophantine( T a, T b, T c, T &x, T &y ) {
    T s,t, g = egcd(a,b,s,t);
    if( (c % g) != 0 )
       return false;
    x = c/g*s; y = c/g*t;
    return true;
```

```
// Given an integer n-by-n matrix A and (positive) integer m,
// compute its determinant mod m.
T integer_det( VVT A, const T M ) {
   const size_t n = A.size();
   FOR(i,0,n) FOR(j,0,n) A[i][j] %= M;
   T det = 1 % M;
   FOR(i,0,n) {
       FOR(j,i+1,n) {
           while( A[j][i] != 0 ) {
               T t = A[i][i] / A[j][i];
               FOR(k,i,n) A[i][k] = (A[i][k] - t*A[j][k]) % M;
               swap( A[i], A[j] );
               det *= -1;
        if( A[i][i] == 0 ) return 0;
       det = (det * A[i][i]) % M;
   if( det < 0 ) det += M;
   return det;
T mult_mod(T a, T b, T m) {
   Τq;
   Tr:
   asm(
            "mulq %3;"
            "divq %4;"
           : "=a"(q), "=d"(r)
           : "a"(a), "rm"(b), "rm"(m));
   return r:
/* Computes a^b mod m. Assumes 1 <= m <= 2^62-1 and 0^0=1.
* The return value will always be in [0, m) regardless of the sign of a.
T pow_mod(T a, T b, T m) {
   if (b == 0) return 1 % m;
   if (b == 1) return a < 0 ? a % m + m : a % m;
   T t = pow_mod(a, b / 2, m);
   t = mult mod(t, t, m);
   if (b % 2) t = mult_mod(t, a, m);
   return t \ge 0 ? t : t + m;
/* A deterministic implementation of Miller-Rabin primality test.
* This implementation is guaranteed to give the correct result for n < 2^64
* by using a 7-number magic base.
* Alternatively, the base can be replaced with the first 12 prime numbers
* (prime numbers <= 37) and still work correctly.
bool is_prime(T n) {
   T small_primes[] = {2, 3, 5, 7, 11, 13, 17, 19, 23, 29, 31, 37};
   for (int i = 0; i < 12; ++i)
        if (n > small_primes[i] && n % small_primes[i] == 0)
           return false:
   T base[] = {2, 325, 9375, 28178, 450775, 9780504, 1795265022};
   T d = n - 1:
   int s = 0:
   for (; d \% 2 == 0; d /= 2, ++s);
   for (int i = 0; i < 7; ++i) {
       T a = base[i] % n:
       if (a == 0) continue;
       T t = pow_mod(a, d, n);
       if (t == 1 \mid | t == n - 1) continue:
       bool found = false:
       for (int r = 1; r < s; ++r) {
           t = pow_mod(t, 2, n);
           if (t == n - 1) {
               found = true;
               break;
        if (!found)
           return false:
```

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return true:
    FFT.cc
// Based on implementation at: http://e-maxx.ru/algo/fft_multiply,

→ http://serjudging.vanb.org/wp-content/uploads/kinversions_hcheng.cc

// Solves UVa Online Judge "Golf Bot" in 2.35 seconds
// f[0...N-1] and g[0..N-1] are numbers, N is a power of 2.
// Want to compute the convolution h, defined by
// h[n] = sum \ of \ f[k]q[n-k] \ (k = 0, ..., N-1).
// Here, the index is cyclic; f[-1] = f[N-1], f[-2] = f[N-2], etc.
// Let F[0...N-1] be FFT(f), and similarly, define G and H.
// The convolution theorem says H[n] = F[n]G[n] (element-wise product).
// To compute h[] in O(N \log N) time, do the following:
   1. Compute F and G (pass invert=false as the argument).
   2. Get H by element-wise multiplying F and G
// 3. Get h by taking the inverse FFT (pass invert=true)
#define MAXN (1 << 3) // must be power of 2
typedef complex<double> T;
typedef vector<T> VT;
const double PI = 4*atan(1);
void FFT(VT &a. bool invert) {
   size_t n = a.size();
   for (int i=1, j=0; i<n; ++i) {
       int bit = n >> 1;
       for (; j>=bit; bit>>=1)
           j -= bit;
       i += bit:
       if (i < j)
           swap(a[i], a[j]);
   }
   for (int len=2; len<=n; len<<=1) {
       double ang = 2*PI/len * (invert ? -1 : 1);
       T wlen (cos(ang), sin(ang));
       for (int i=0; i<n; i+=len) {
           for (int j=0; j<len/2; ++j) {
               T u = a[i+j], v = a[i+j+len/2] * w;
               a[i+i] = u + v:
               a[i+j+len/2] = u - v;
               w *= wlen;
       }
   }
   if(invert)
       for (int i=0; i<n; ++i)
           a[i] /= n;
    Linear Algebra.cc
// Useful linear algebra routines.
#define FOR(v,l,u) for (size_t v = l; v < u; ++v)
typedef double
                      T; // the code below only supports fields
typedef vector<T>
                      VT.
typedef vector<VT>
                      VVT:
typedef vector<size_t> VI;
typedef vector<bool> VB;
typedef vector<long long> VN;
typedef vector<VN> VVN;
typedef long long 11;
// Given an m-by-n matrix A, compute its reduced row echelon form,
// returning a value like the determinant.
// If m = n, the returned value *is* the determinant of A.
```

// If m != n, the returned value is nonzero iff A has full row rank.

// To compute rank(A), get its RREF, and count the nonzero rows.

```
T GaussJordan ( VVT &A ) {
    const size_t m = A.size(), n = A[0].size();
    T det = 1:
    size_t pj = 0;
                          // walking pointer for the pivot column
    FOR(k,0,m) {
        size_t pi = k;
        while(pj < n) { // find the best row below k to pivot
            FOR(i,k,m) if( fabs(A[i][pj]) > fabs(A[pi][pj]) ) pi = i;
            if( !feq(0.0, A[pi][pj]) ) { // we have our new pivot
                if( pi != k ) {
                    swap( A[pi], A[k] );
                   pi = k:
                    det *= -1:
                break;
           FOR(i,k,m) A[i][pj] = 0; // This column is zeros below row k
                                     // So move on to the next column
        if( pj == n ) { det = 0; break; } // we're done early
        T s = 1.0/A[pi][pj]; // scale the pivot row
        FOR(j,pj,n) A[pi][j] *= s;
        FOR(i,0,m) if( i != pi ) { // subtract\ pivot\ from\ other\ rows
            T a = A[i][pj];
                                    // multiple of pivot row to subtract
            FOR(j,pj,n) A[i][j] = a*A[pi][j];
        ++pj;
    }
    return det;
// In-place invert A.
void InvertMatrix( VVT &A ) {
    const size t n = A.size():
    FOR(i,0,n) FOR(j,0,n) A[i].push_back((i==j) ? 1 : 0); // augment
    Gauss.Jordan( A ):
                                               // compute RREF
    FOR(i,0,n) FOR(j,0,n) A[i][j] = A[i][j+n]; // copy A inverse over
    FOR(i,0,n) A[i].resize(n);
                                               // get rid of cruft
// Given m-by-n A and m-by-q b, compute a matrix x with Ax = b.
// This solves q separate systems of equations simultaneously.
// Fix k in [0,q).
// x[*][k] indicates a candidate solution to the jth equation.
// has_sol[k] indicates whether a solution is actually solution.
// The return value is the dimension of the kernel of A.
// Note that this is the dimension of the space of solutions when
size_t SolveLinearSystems( const VVT &A, const VVT &b, VVT &x, VB &has_sol )
    const size_t m = A.size(), n = A[0].size(), q = b[0].size();
    FOR(i,0,m) \ FOR(j,0,q) \ M[i].push\_back(b[i][j]); \ // \ augment
    Gauss.Jordan( M ):
                                                   // RREF
    x = VVT(n, VT(q, 0));
    size_t i = 0, jz = 0;
    while( i < m ) {
        while( jz < n \&\& feq(M[i][jz],0) ) ++jz;
        if( jz == n ) break; // all zero means we're starting the kernel
        FOR(k,0,q) x[jz][k] = M[i][n+k]; // first nonzero is always 1
    size_t kerd = n - i; // i = row rank = column rank
    has_sol = VB(q,true);
    while( i < m ) {
        FOR(k,0,q) if( !feq(M[i][n+k],0) ) has_sol[k] = false;
    }
    return kerd:
// Given m-by-n A, compute a basis for the kernel of A.
// The return value is in K, which is interpreted as a length-d array of
// n-dimensional vectors. (So K.size() == dim(Ker(A)))
// The return value is K.size().
size_t KernelSpan( const VVT &A, VVT &K ) {
    const size_t m = A.size(), n = A[0].size();
    VVT M = A:
    Gauss Jordan (M) :
    K = VVT():
```

```
VB all zero(n.true):
    FOR(i,0,m) {
        size t iz = 0:
        while( jz < n && feq(M[i][jz],0) ) ++jz;
        if( jz == n ) break; // skip to the easy part of the kernel
        all_zero[jz] = false;
        FOR(j,jz+1,n) if( !feq(M[i][j],0) ) {
            all_zero[j] = false;
            K.push_back( VT(n,0) );
           K.back()[jz] = -1 * M[i][j];
           K.back()[j] = 1;
       }
   FOR(j,0,n) if( all_zero[j] ) {
       K.push_back( VT(n,0) );
       K.back()[i] = 1;
   return K.size();
// code for fast linear recurrence evaluation. Based on blog post at:
// http://fusharblog.com/solving-linear-recurrence-for-programming-contest/
// used for AC answer at: http://codeforces.com/contest/678/problem/D
// compute AB with entries mod M
VVN matrix_mult(VVN A, VVN B, 11 M) {
   VVN result(A.size(), VN(B[0].size(), 0));
   for (int i = 0; i < result.size(); ++i) {</pre>
       for (int j = 0; j < result[i].size(); ++j) {
           for (int k = 0; k < A[0].size(); ++k) {
               result[i][j] = (result[i][j] + A[i][k] * B[k][j]) % M;
   }
   return result;
// compute A^n with entries mod M
// if A is m x m, takes O(m^3 \log n) time
VVN matrix_pow(VVN A, 11 n, 11 M) {
   if (n == 1) {
       return A;
   else if (n \% 2 == 0) {
        VVN smaller = matrix_pow(A, n/2, M);
        return matrix_mult(smaller, smaller, M);
   else {
        return matrix_mult(matrix_pow(A, n-1, M), A, M);
// computes nth term of f(n) = rec[0]*f(n-1) + rec[1]*f(n-2) + ... +
\hookrightarrow rec[k-1]*f(n-k) + c
// given that f(1) = init[0], f(2) = init[1], ..., f(k) = init[k-1]
// in O(k^3 log n) time
11 eval_rec(VN rec, VN init, T n, T M, T c = 0) {
   size t k = rec.size():
   VVN mat(k+1, VN(k+1, 0));
   for (int i = 0; i < k; ++i) {
       mat[i][i+1] = 1;
   for (int i = 0; i < k; ++i) {
       mat[k-1][i] = rec[k-1-i];
   mat[k][k] = 1:
       return init[0];
   VVN mat_new = matrix_pow(mat, n-1, M);
    VVN vect(k+1, VN(1));
   for (int i = 0; i < k; ++i)
       vect[i][0] = init[i];
    vect[k][0] = c;
```

Algebra.cc LinearAlgebra.cc

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VVN vect new = matrix mult(mat new, vect, M);
   return vect_new[0][0];
    Simplex.cc
// Ripped from http://web.stanford.edu/~liszt90/acm/notebook.html#file17
#include <iostream>
#include <iomanin>
#include <vector>
#include <cmath>
#include inits>
using namespace std;
// BEGIN CUT
#define ACM_assert(x) \{if(!(x))*((long *)0)=666;\}
//#define TEST LEAD OR GOLD
#define TEST_HAPPINESS
typedef long double DOUBLE;
typedef vector<DOUBLE> VD;
typedef vector<VD> VVD;
typedef vector<int> VI;
const DOUBLE EPS = 1e-9;
struct LPSolver {
   int m, n;
   VI B, N;
   VVD D:
   LPSolver(const VVD &A, const VD &b, const VD &c) :
       m(b.size()), n(c.size()), N(n+1), B(m), D(m+2, VD(n+2)) {
       for (int i = 0; i < m; i++) for (int j = 0; j < n; j++) D[i][j] =
        for (int i = 0; i < m; i++) { B[i] = m+i; D[i][n] = -1; D[i][n+1] = -1
        \hookrightarrow b[i]; }
       for (int j = 0; j < n; j++) { N[j] = j; D[m][j] = -c[j]; }
       N[n] = -1; D[m+1][n] = 1;
   void Pivot(int r, int s) {
       for (int i = 0; i < m+2; i++) if (i != r)
            for (int j = 0; j < n+2; j++) if (j != s)
               D[i][j] -= D[r][j] * D[i][s] / D[r][s];
       for (int j = 0; j < n+2; j++) if (j != s) D[r][j] /= D[r][s];
       for (int i = 0; i < m+2; i++) if (i != r) D[i][s] /= -D[r][s];
       D[r][s] = 1.0 / D[r][s];
       swap(B[r], N[s]);
   bool Simplex(int phase) {
       int x = phase == 1 ? m+1 : m;
       while (true) {
            for (int j = 0; j \le n; j++) {
                if (phase == 2 && N[j] == -1) continue;
                \hookrightarrow N[j] < N[s]) s = j;
            if (D[x][s] >= -EPS) return true;
            int r = -1;
            for (int i = 0: i < m: i++) {
               if (D[i][s] <= 0) continue;</pre>
                if (r == -1 \mid \mid D[i][n+1] / D[i][s] < D[r][n+1] / D[r][s] \mid \mid
                        \label{eq:defD} D[i]\,[n+1] \ / \ D[i]\,[s] \ == \ D[r]\,[n+1] \ / \ D[r]\,[s] \ \&\& \ B[i] \ <
                         \hookrightarrow B[r]) r = i;
            if (r == -1) return false;
            Pivot(r, s);
   DOUBLE Solve(VD &x) {
       int r = 0;
       for (int i = 1; i < m; i++) if (D[i][n+1] < D[r][n+1]) r = i;
       if (D[r][n+1] \leftarrow -EPS) {
            if (!Simplex(1) || D[m+1][n+1] < -EPS) return
```

-numeric limits<DOUBLE>::infinity():

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for (int i = 0; i < m; i++) if (B[i] == -1) {
                for (int j = 0; j \le n; j++)
                    if (s == -1 || D[i][j] < D[i][s] || D[i][j] == D[i][s] &&
                    \hookrightarrow N[j] < N[s]) s = j;
                Pivot(i, s);
        if (!Simplex(2)) return numeric_limits<DOUBLE>::infinity();
        for (int i = 0; i < m; i++) if (B[i] < n) x[B[i]] = D[i][n+1];
        return D[m][n+1];
    // BEGIN CUT
    void Print() {
        cout << "N = "; for (int i = 0; i < N.size(); i++) printf("%8d",</pre>
        cout << "B = "; for (int i = 0; i < B.size(); i++) printf("%8d",</pre>

    B[i]); cout << endl;
</pre>
        cout << endl:
        for (int i = 0; i < D.size(); i++) {
           for (int j = 0; j < D[i].size(); j++) {
               printf("%8.2f", double(D[i][j]));
           printf("\n");
       printf("\n");
    // END CUT
// BEGIN CUT
#ifdef TEST_HAPPINESS
int main() {
   int n. m:
    while (cin >> n >> m) {
        ACM_assert(3 <= n && n <= 20);
        ACM_assert(3 <= m && m <= 20);
        VVD A(m, VD(n));
        VD b(m), c(n);
        for (int i = 0; i < n; i++) {
           cin >> c[i]:
            ACM_assert(c[i] >= 0);
            ACM_assert(c[i] <= 10);</pre>
        for (int i = 0; i < m; i++) {
            for (int j = 0; j < n; j++)
               cin >> A[i][j];
            cin >> b[i]:
            ACM_assert(b[i] >= 0);
            ACM assert(b[i] <= 1000):
        LPSolver solver(A, b, c);
        VD sol:
        DOUBLE primal_answer = m * solver.Solve(sol);
        VVD AT(A[0].size(), VD(A.size()));
        for (int i = 0; i < A.size(); i++)
            for (int j = 0; j < A[0].size(); j++)
                AT[j][i] = -A[i][j];
        for (int i = 0; i < c.size(); i++)
           c[i] = -c[i];
        for (int i = 0; i < b.size(); i++)
            b[i] = -b[i];
        LPSolver solver2(AT, c, b);
        DOUBLE dual_answer = -m * solver2.Solve(sol);
        ACM_assert(fabs(primal_answer - dual_answer) < 1e-10);
        int primal rounded answer = (int) ceil(primal answer);
        int dual_rounded_answer = (int) ceil(dual_answer);
        // The following assert fails b/c of the input data.
        // ACM_assert(primal_rounded_answer == dual_rounded_answer);
        cout << "Nasa can spend " << primal_rounded_answer << " taka." <</pre>
         \hookrightarrow endl;
   }
#ifdef TEST_LEAD_OR_GOLD
int main() {
    int n:
```

```
int ct = 0:
    while (cin >> n) {
        if (n == 0) break:
        VVD A(6, VD(n));
        VD b(6), c(n, -1);
        for (int i = 0; i < n; i++) {
           for (int j = 0; j < 3; j++) {
               cin >> A[j][i]; A[j+3][i] = -A[j][i];
        for (int i = 0; i < 3; i++) {
           cin >> b[i]; b[i+3] = -b[i];
       if (ct > 0) cout << endl;
        cout << "Mixture " << ++ct << endl;</pre>
       LPSolver solver(A, b, c);
       VD x:
        double obj = solver.Solve(x);
       if (isfinite(obj)) {
           cout << "Possible" << endl:
       } else {
           cout << "Impossible" << endl;</pre>
   }
   return 0;
#else
int main() {
   const int m = 4;
   const int n = 3;
   DOUBLE A[m][n] = {
       { 6, -1, 0 },
       \{-1, -5, 0\},
       { 1, 5, 1 },
       { -1, -5, -1 }
   DOUBLE _b[m] = \{ 10, -4, 5, -5 \};
   DOUBLE _c[n] = \{ 1, -1, 0 \};
   VVD A(m):
   VD b(_b, _b + m);
   VD c(_c, _c + n);
   for (int i = 0; i < m; i++) A[i] = VD(_A[i], _A[i] + n);
   LPSolver solver(A, b, c);
   DOUBLE value = solver.Solve(x);
   cerr << "VALUE: "<< value << endl;
   cerr << "SOLUTION:":
   for (size_t i = 0; i < x.size(); i++) cerr << " " << x[i];
   cerr << endl:
   return 0;
// BEGIN CUT
#endif
#endif
// END CUT
    Rational.cc
```

```
// Rational struct. Uses lcm to keep in simplified form.
// Simply replace "double" (or "int") with "rat" and use
// contructor to initialize constants. Rat class handles
// everything else.
//
// Written in such a way as to avoid overflow if possible.
struct rat {
    T n, d;
    rat(T n, T d) {
        T k = gcd(n, d);
        this->n = n/k;
}
```

Linear Algebra.cc Rational.cc

```
this -> d = d/k:
    rat(T n) : n(n), d(1) {}
rat operator + (const rat &a, const rat &b) {
   T new_d = lcm(a.d, b.d); // if overflow occurs, this may be 0
                             // causing floating point exception
   T a scale = new d / a.d:
   T b_scale = new_d / b.d;
    return rat(a.n*a_scale + b.n*b_scale, new_d);
rat operator * (const T s, const rat &a) {
    return rat(a.n * s. a.d):
rat operator * (const rat &a, const T s) {
rat operator - (const rat &a, const rat &b) {
    return a + (-1 * b);
rat operator * (const rat &a, const rat &b) {
   return rat(a.n*b.n, a.d*b.d);
rat operator / (const rat &a, const rat &b) {
    return a * rat(b.d, b.n);
    SegmentTree.cc
// Segment Tree with lazy propagation. This solves
// AhoyPirates on UVa Online Judge in 1.172 seconds.
// To use, you only need to change SegmentNode data,
// the merge function, update, and initialization;
// the functions are written in such a way as to handle everything else.
// If range updates are not necessary,
// ignore the updateVal and rangeUpdate function.
// The following implementation is an example for RMQ.
#define MAXN 1000 // the maximum input length of the sequnece, good idea to
 \hookrightarrow add 10 or so to this
typedef signed long long int T; // the type of the underlying sequence
typedef vector<size_t> VI;
typedef vector<T> VT;
struct SegmentNode {
    // segment data
   T maxVal:
    // update data
   T updateVal;
SegmentNode st[MAXN*4];
T A [MAXN]:
size_t left(size_t cur) { return cur << 1; }</pre>
size_t right(size_t cur) { return (cur << 1) + 1; }</pre>
// create merging function here
void merge(SegmentNode &left, SegmentNode &right, SegmentNode &result) {
    result.maxVal = max(left.maxVal, right.maxVal);
// only use if range update is needed
void updateChildren(size_t cur, size_t L, size_t R);
// merge handles all querying, no changes needed here
// note that when calling any segment functions in main,
```

```
// the first three parameters will be 1, 0, length-1
SegmentNode query(size_t cur, size_t L, size_t R, size_t LQ, size_t RQ) {
    if (L >= LO && R <= RO)
        return st[cur];
    updateChildren(cur, L, R);
    size_t M = (L+R)/2;
       return query(right(cur), M+1, R, LQ, RQ);
    if (M+1 > RQ)
        return query(left(cur), L, M, LQ, RQ);
    SegmentNode leftResult = query(left(cur), L, M, LQ, RQ);
    SegmentNode rightResult = query(right(cur), M+1, R, LQ, RQ);
    SegmentNode result:
    merge(leftResult, rightResult, result);
    return result;
// only implement if necessary
void update(size_t cur, size_t L, size_t R, size_t idx, T val) {
    if (L == idx && R == idx) {
       // write update of single value here
        st[cur].maxVal = val;
    else if (L <= idx && R >= idx) {
       size_t M = (L+R)/2;
        update(left(cur), L, M, idx, val);
        update(right(cur), M+1, R, idx, val);
        merge(st[left(cur)], st[right(cur)], st[cur]);
   }
// only implement if necessary
void rangeUpdate(size_t cur, size_t L, size_t R, size_t Lbound, size_t

→ Rbound, T val) {
    if (L >= Lbound && R <= Rbound) {
        // implement range update here
       st[cur].maxVal += val;
        // set undate vals here
        st[cur].updateVal += val;
    else if (L <= Rbound && R >= Lbound) {
        updateChildren(cur, L, R);
        size_t M = (L+R)/2;
        rangeUpdate(left(cur), L, M, Lbound, Rbound, val);
        rangeUpdate(right(cur), M+1, R, Lbound, Rbound, val);
        merge(st[left(cur)], st[right(cur)], st[cur]);
void updateChildren(size_t cur, size_t L, size_t R) {
    rangeUpdate(left(cur), L, R, L, R, st[cur].updateVal);
    rangeUpdate(right(cur), L, R, L, R, st[cur].updateVal);
    // reset update vals
    st[cur].updateVal = 0;
void build(size_t cur, size_t L, size_t R) {
    // initialize update vals
    st[cur].updateVal = 0;
        // initialize single value here
        st[cur].maxVal = A[L];
        size t M = (L+R)/2:
        build(left(cur), L, M);
```

```
build(right(cur), M+1, R);
       merge(st[left(cur)], st[right(cur)], st[cur]);
}
    BIT.cc
// T is a type with +/- operations and identity element '0'.
// Least significant bit of a. Used throughout.
int LSB( int a ) { return a ^ (a & (a-1)); }
// To use it, instantiate it as 'BIT(n)' where n is the size of the

    underlying

// array. The BIT then assumes a value of 0 for every element. Update each
// index individually (with 'add') to use a different set of values.
// Note that it is assumed that the underlying array has size a power of 2!
// This mostly just simplifies the implementation without any loss in speed.
// Just use the closest power of 2 larger than the max input size. Even if

→ some test

// cases do not test this high, initialization is extremely quick.
// The comments below make reference to an array 'arry'. This is the
// array. (A is the data stored in the actual tree.)
struct BIT {
   VT A:
   BIT( int n ): N(n), A(N+1,0) {} // n must be a power of 2
    // add v to arry[idx]
   void add( int idx, T v ) {
       for( int i = idx+1; i <= N; i += LSB(i) ) A[i] += v;
   // get sum( arry[0..idx] )
   T sum( int idx ) {
       for( int i = idx+1; i > 0; i -= LSB(i) ) ret += A[i];
        return ret;
    // get sum( arry[l..r] )
   T sum_range( int 1, int r ) { return sum(r) - sum(1-1); }
   // Find largest r so that sum( arry[0..r] ) <= thresh
   // This assumes arry[i] >= 0 for all i > 0, for monotonicity.
   // This takes advantage of the specific structure of LSB() to simplify
    \hookrightarrow the
   // binary search.
   int largest_at_most( T thresh ) {
       int r = 0, del = N;
       while( del && r <= N ) {
           int q = r + del;
           if( A[q] <= thresh ) {
               thresh -= A[q];
           del /= 2;
        return r-1;
   }
};
// A 'range-add'/'index query' BIT
struct BIT_flip {
   BIT_flip( int n ) : A(n) {}
   // add v to arry[l,r]
   void add( int 1, int r, T v ) {
       A.add(1.v):
       A.add(r+1,-v);
    // get arry[idx]
   T query( int idx ) {
       return A.sum(idx);
```

Rational.cc BIT.cc

```
// A 'range-add'/'range-query' data structure that uses BITs.
struct BIT_super {
    int N;
    BIT_flip m, b; // linear coefficient, constant coefficient
   BIT_super( int n ) : N(n), m(n), b(n) {}
    // add v to arry[l..r]
    void add( int 1, int r, T v ) {
        m.add(1,r,v);
                             // add slope on active interval
        b.add(1,N,1*(-v)); // subtract contribution from pre-interval
        b.add(r+1,N,(r+1)*v);
                                    // add total contribution to
         \hookrightarrow after-interval
    // get sum( arry[0..r] )
   T query( int r ) {
        return m.query(r)*r + b.query(r);
    // get sum( arry[l..r] )
   T query_range( int 1, int r ) {
       return query(r) - query(1-1);
// A 2-dimensional specialization of BITd. (see below)
// What took 'nlogn' before now takes 'nlog^2(n)'.
struct BIT2 {
   int N2;
   vector<BIT> A:
   BIT2( int n1, int n2 ) : N1(n1), N2(n2) {
       A.resize(N1+1, BIT(n2));
    // add v to arry[x][y]
    void add( int x, int y, T v ) {
       for( int i = x+1; i <= N1; i += LSB(i) ) A[i].add(y,v);
    // get sum( arry[0..x][0..y] ).
   T sum( int x, int y ) {
       T ret = 0;
       for( int i = x+1; i > 0; i -= LSB(i) ) ret += A[i].sum(y);
        return ret;
    // get sum( arry[xL..xH)[yL..yH) ).
   T sum_range( int xL, int yL, int xH, int yH ) {
        return sum(xH,yH) + sum(xL-1,yL-1) - sum(xH,yL-1) - sum(xL-1,yH);
// A d-dimensional binary indexed tree
// What took 'nlogn' before now takes 'nlog^d(n)'.
// To construct it, set dims to be the vector of dimensions, and pass
// d <- dims.size().
typedef vector<int> VI;
struct BITd {
   int N:
   int D:
    vector<BITd> A;
   T V·
   BITd( const VI &dims, int d ) : N(dims[d-1]), D(d) {
       if( D == 0 ) V = 0;
                    A.resize( N+1, BITd( dims, D-1 ) );
   void add( const VI &idx, T v ) {
        if( D == 0 ) V += v;
        for (int i = idx[D-1]+1; i \le N; i += LSB(i)) A[i].add(idx.v);
   T sum( const VI &idx ) {
       if( D == 0 ) return V;
        T ret = 0:
        for( int i = idx[D-1]+1; i > 0; i -= LSB(i) ) ret += A[i].sum(idx);
   T sum_range( VI lo, VI hi ) {
        FOR(i,0,D) --lo[i];
        // In higher dimensions, we have to use inclusion-exclusion
        int BD = ((int)1) << D;</pre>
```

```
T ret = 0:
        FOR(S,0,BD) {
            int sign = 1:
            VI q(lo);
           FOR(b,0,BD) if((S >> b) & 1) {
               q[b] = hi[b];
               sign *= -1;
           ret += sign * sum(q);
        return ret;
};
    KMP.cc
// An implemention of Knuth-Morris-Pratt substring-finding.
// The table constructed with KMP_table may have other uses.
typedef vector<size_t> VI;
// In the KMP table, T[i] is the *length* of the longest *prefix*
// which is also a *proper suffix* of the first i characters of w.
void KMP_table( string &w, VI &T ) {
   T = VI(w.size()+1);
    size_t i = 2, j = 0;
    T[1] = 0; // T[0] is undefined
    while( i <= w.size() ) {
       if( w[i-1] == w[j] ) { T[i] = j+1; ++i; ++j; } // extend previous
        else if(j > 0)
                              {j = T[j];}
                                                       // fall back
        else
                              \{T[i] = 0; ++i; \}
                                                       // give up
   }
// Search for first occurrence of q in s in O(|q|+|s|) time.
size_t KMP( string &s, string &q ) {
    size_t m, z; m = z = 0; // m is the start, z is the length so far
    VI T; KMP_table(q, T);
                             // init the table
    while( m+z < s.size() ) { // while we're not running off the edge...
        if( q[z] == s[m+z] ) { // next character matches
            if( z == q.size() ) return m; // we're done
       else if( z > 0 ) { // fall back to the next best match
           m += z - T[z]; z = T[z];
        else {
                               // go back to start
           m += 1:
       }
    return s.size():
    AhoCorasick.cc
// An implementation of Aho-Corasick dictionary matching algorithm
// Taken directly from the paper
typedef vector<size t> VI:
typedef vector<string> VS;
struct node {
    unordered_map<char, node*> g;
    node* f;
    node* output = NULL;
    bool isWord = false:
    size_t num;
    node(size_t num) : num(num) {}
    node() {}
void enter(string S, node *root, size_t &n) {
    node *state = root;
    size t i = 0:
    while (state->g.count(S[j]) != 0) {
        state = state->g[S[j]];
        ++j;
```

```
for (; j < S.size(); ++j) {
        state->g[S[j]] = new node(n++);
       state = state->g[S[j]];
   state->isWord = true:
void construct_f(node *root) {
   queue<node*> q;
   for (auto it = root->g.begin(); it != root->g.end(); ++it) {
       q.push(it->second);
       it->second->f = root;
   while (!q.empty()) {
       node *r = q.front(); q.pop();
        for (auto it = r->g.begin(); it != r->g.end(); ++it) {
           q.push(it->second);
           node *state = r->f;
           while (state->g.count(it->first) == 0 && state->num != 0)
               state = state->f:
           if (state->g.count(it->first) != 0)
               it->second->f = state->g[it->first];
               it->second->f = root:
           if (it->second->f->isWord)
               it->second->output = it->second->f;
           else // may assign NULL pointer
               it->second->output = it->second->f->output;
       7-
   }
node* ConstructAutomaton(VS &dictionary) {
   size t n = 0:
    node *root = new node(n++);
   for (size_t i = 0; i < dictionary.size(); ++i) {</pre>
        enter(dictionary[i], root, n);
   construct_f(root);
    return root:
void PrintMatches(node *root, string x) {
   node *state = root:
   for (size_t i = 0; i < x.size(); ++i) {
       while (state->g.count(x[i]) == 0 \&\& state->num != 0) state =

    state->f;

       if (state->g.count(x[i]) != 0)
           state = state->g[x[i]];
       node *out = state;
       while (true) {
           if (out->isWord) cout << i << " " << out->num << " " << endl;
           if (out->output == NULL) break;
           out = out->output;
   }
    Suffix Array.cc
// A prefix-doubling suffix array construction implementation.
#define FOR(v,l,u) for (size_t v = l; v < u; ++v)
typedef vector<size_t> VI;
typedef pair<int, int> II;
#define MAX_N 100010
int RA[MAX_N], tempRA[MAX_N];
int SA[MAX_N], tempSA[MAX_N];
int c[MAX_N];
// uses Radix Sort as a subroutine to sort in O(n)
void CountingSort(int n, int k) {
   int i, sum, maxi = max(300, n);
   memset(c. 0. sizeof c):
   for (i = 0; i < n; i++)
```

```
c[i+k < n ? RA[i+k] : 0]++;
    for (i = sum = 0; i < maxi; i++) {
        int t = c[i]; c[i] = sum; sum += t;
    for (i = 0: i < n: i++)
        tempSA[c[SA[i]+k < n ? RA[SA[i]+k] : 0]++] = SA[i];
    for (i = 0; i < n; i++)
        SA[i] = tempSA[i];
// Construct SA in O(n log n) time. Solves UVa Online Judge "Glass Beads" in

    ∴ 5 seconds

void ConstructSA(string T) {
    int i, k, r, n = T.size();
    for (i = 0; i < n; i++) RA[i] = T[i];
    for (i = 0; i < n; i++) SA[i] = i;
   for (k = 1; k < n; k <<= 1) {
        CountingSort(n, k);
        CountingSort(n, 0);
        tempRA[SA[O]] = r = 0;
        for (i = 1; i < n; i++)
           tempRA[SA[i]] =
               (RA[SA[i]] == RA[SA[i-1]] &\& RA[SA[i]+k] == RA[SA[i-1]+k]) ?
                for (i = 0; i < n; i++)
            RA[i] = tempRA[i];
        if (RA[SA[n-1]] == n-1) break;
// Given a "string" w, and suffix array SA, compute the array LCP for which
// the suffix starting at SA[i] matches SA[i+1] for exactly LCP[i]
 // It is assumed that the last character of w is the unique smallest-rank
// character in w. Runs in O(n).
void LongestCommonPrefix( const string &w, VI &LCP ) {
    const size_t N = w.size(); VI rk(N);
   FOR(i,0,N) rk[SA[i]] = i;
   LCP = VI(N-1); size_t k = 0;
   FOR(i,0,N) {
        if( rk[i] == N-1 ) continue;
        size_t j = SA[ rk[i]+1 ];
        while(w[i+k] == w[j+k]) ++k;
        LCP[rk[i]] = k;
        if(k > 0) --k;
   }
// Finds the smallest and largest i such that the prefix of suffix SA[i]
// the pattern string P. Returns (-1, -1) if P is not found in T. Runs in O(m
 \hookrightarrow log n).
II StringMatching(const string &T, const string P) {
    int n = T.size(), m = P.size();
   int lo = 0, hi = n-1, mid = lo;
   while (lo < hi) {
        mid = (lo + hi) / 2;
        int res = T.compare(SA[mid], m, P);
        if (res >= 0) hi = mid;
                     lo = mid+1;
   if (T.compare(SA[lo], m, P) != 0) return II(-1, -1);
   II ans; ans.first = lo;
   lo = 0; hi = n-1; mid = lo;
    while (lo < hi) {
       mid = (lo + hi) / 2;
        int res = T.compare(SA[mid], m, P);
        if (res > 0) hi = mid;
        else
                    lo = mid+1:
   if (T.compare(SA[hi], m, P)) hi--;
    ans.second = hi;
   return ans;
}
    ArticulationPoint.cc
// This is code for computing articulation points of graphs,
// ie points whose removal increases the number of components in the graph.
```

```
// This works when the given graph is not necessarily connected, too.
typedef vector<size_t> VI;
typedef vector<VI> VVI;
typedef vector<bool> VB;
struct artpt_graph {
   // basic graph stuff
    VI parent, n_children, rank; // dfs tree
    VB is_art; VI reach;
                                 // articulation points
    artpt_graph( size_t N ) : N(N), adj(N), is_art(N) {}
    void add_edge( size_t s, size_t t ) {
        adj[s].push_back(t);
        adj[t].push_back(s);
    size_t dfs_artpts( size_t rt, VB &visited, size_t R ) {
        visited[rt] = true;
        rank[rt] = R++:
        reach[rt] = rank[rt]; // reach[rt] <= rank[rt] always.
        FOR(i,0,adj[rt].size()) {
           size_t v = adj[rt][i];
            if( v == parent[rt] ) continue;
            if ( visited[v] )
               reach[rt] = min(reach[rt], rank[v]);
            else {
               ++n_children[rt];
               parent[v] = rt;
               R = dfs_artpts( v, visited, R );
               reach[rt] = min(reach[rt], reach[v]);
        if( reach[rt] < rank[rt] || n_children[rt] == 0 )</pre>
           is_art[rt] = false;
        return R;
    void comp_articulation_points() {
        is_art = VB(N, true); reach = VI(N);
        parent = VI(N,N);
                             rank = VI(N);
                                                n children = VI(N.0):
        VB visited(N,false); size_t R = 0;
        FOR(i,0,N) {
            if( visited[i] ) continue;
           R = dfs_artpts(i, visited, R); // this is not right on i
           is_art[i] = (n_children[i] >= 2); // but we can fix it!
   }
};
    BellmanFord.cc
// A Bellman-Ford implementation.
// bellmanford(S) computes the shortest paths from S to all other nodes.
// It returns true if there are no negative cycles in the graph,
// and false otherwise.
// D[v] is set to the shortest path from S to v (when it exists).
// P[v] is set to the parent of v in the shortest-paths tree,
// or N (for which there is no index) if v is not reachable from S.
#define FOR(v,l,u) for (size_t v = l; v < u; ++v)
typedef signed long long int T;
typedef vector<T>
typedef vector<VT>
                      VVT:
typedef vector<bool>
typedef vector<VB>
                     VVB;
typedef vector<size_t> VI;
typedef vector<VI> VVI;
const T UNBOUNDED = numeric_limits<T>::min(); // -infinity for doubles
const T INFINITY = numeric_limits<T>::max(); // infinity for doubles
struct bellmanford_graph {
    size_t N; // number of nodes
    VVI A; // adjacency list
          W; // weight of edges
          D; // shortest distance
          P; // parent in the shortest path tree
    bellmanford_graph( size_t N ) : N(N), A(N), W(N) {}
    void add_edge( size_t s, size_t t, T w ) {
        A[s].push_back(t);
        W[s].push_back(w);
```

```
bool bellmanford( size t S ) {
       D = VT(N, INFINITY); D[S] = 0; P = VI(N,N);
       FOR(k.O.N)
       FOR(s,0,N)
       FOR(i,0,A[s].size()) {
           size_t t = A[s][i];
           if( D[s] == INFINITY ) continue;
           if( D[t] > D[s] + W[s][i] ) {
               if(k == N-1) {
                   D[t] = UNBOUNDED;
               else {
                   D[t] = D[s] + W[s][i];
                   P[t] = s;
               }
       FOR(v,0,N) if( D[v] == UNBOUNDED ) return false;
       return true;
};
```

FloydWarshall.cc

```
// Floyd-Warshall implementation with negative cycle detection.
// This will modify the graph, computing its transitive closure.
// If there is an upper bound for any simple path length,
// then create a constant INF equal to that,
// and set W[i][j] = INF when there is no edge i \rightarrow j.
// You can then remove all reference to A.
// Notable generalizations:
// - Finding paths with maximum minimum-capacity-along-path
// - Transitive closure (done with A below)
#define FOR(v,l,u) for (size_t v = l; v < u; ++v)
typedef signed long long int T; // anything with <, +, and 0
typedef vector<T>
                      VT;
typedef vector<VT>
                      VVT:
typedef vector<size_t> VI; // only if you want the actual paths
typedef vector<VI>
                      VVI; // ^^^^
typedef vector <bool> VB; // only if you don't have an upper bound
typedef vector<VB>
                      VVB; // ------
struct floydwarshall_graph {
   size_t N; // Number of nodes
   VVB A; // [i][j] is true iff there exists an edge i -> j
          W; //[i][j] is the weight of the edge i \rightarrow j.
   VVI P; // [i][j] is the next node in shortest path i -> j
   floydwarshall_graph( size_t n ) :
       N(n), A(n,VB(n,false)), W(n,VT(n,0)), P(n,VI(n,n)) {}
    void add_edge( size_t s, size_t t, T w ) {
        A[s][t] = true; W[s][t] = w; P[s][t] = t;
   bool floydwarshall() {
       FOR(k,0,N) // We've computed paths using only \{0, 1, ..., k-1\}
       FOR(i,0,N) // Now compute the shortest path from i \rightarrow j
       FOR(j,0,N) { // when considering a path using k.
            if( !A[i][k] || !A[k][j] ) continue;
                                                    // skip invalid
           if( !A[i][j] ) {
                                                    // first time
               A[i][i] = true:
               W[i][j] = W[i][k] + W[k][j];
               P[i][j] = P[i][k];
           if( W[i][k] + W[k][j] < W[i][j] ) {
                                                    // future times
               P[i][j] = P[i][k];
               W[i][j] = W[i][k] + W[k][j];
        FOR(i,0,N) if(W[i][i] < 0) return false; // negative cycle.
        return true; // no negative cycle.
};
```

${\bf Max Card Bipartite Matching.cc}$

```
// This code performs maximum (cardinality) bipartite matching.
// Does not support weighted edges.
// Running time: O(|E| |V|) -- often much faster in practice
11
     INPUT: adj_list[i][j] = edge between row node i and column node
            mr[i] = vector of size #rows, initialized to -1
            mc[j] = vector of size #columns, initialized to -1
    OUTPUT: mr[i] = assignment for row node i, -1 if unassigned
            mc[j] = assignment for column node j, -1 if unassigned
             function returns number of matches made
typedef vector<int> VI;
typedef vector<VI> VVI;
typedef vector<bool> VB;
bool FindMatch(int i, const VVI &adj_list, VI &mr, VI &mc, VB &seen) {
 for (int j = 0; j < adj_list[i].size(); j++) {</pre>
   int item = adj_list[i][j];
   if (!seen[item]) {
      seen[item] = true;
     if (mc[item] < 0 || FindMatch(mc[item], adj_list, mr, mc, seen)) {</pre>
       mr[i] = item;
       mc[item] = i;
       return true:
 return false;
// mr should be a vector of size number of row items, initialized to -1
// mc should be a vector of size number of column items, initialized to -1
int BipartiteMatching(const VVI &adj_list, VI &mr, VI &mc) {
 for (int i = 0; i < adj_list.size(); i++) {
   VB seen(mc.size(), false);
   if (FindMatch(i, adj_list, mr, mc, seen)) ct++;
 return ct;
```

MaximumFlow-Dinic.cc

```
// Adjacency list implementation of Dinic's blocking flow algorithm.
// This is very fast in practice, and only loses to push-relabel flow.
// Running time:
     0(|V|^2 |E|)
11
// INPUT:
     - graph, constructed using AddEdge()
     - source
     - sink
// OUTPUT:
    - maximum flow value
     - To obtain the actual flow values, look at all edges with
       capacity > 0 (zero capacity edges are residual edges).
// Taken from Stanford ACM:
const int INF = 2000000000;
struct Edge {
   int from, to, cap, flow, index;
   Edge(int from, int to, int cap, int flow, int index) :
   from(from), to(to), cap(cap), flow(flow), index(index) {}
struct Dinic {
   int N:
   vector<vector<Edge> > G;
   vector<Edge *> dad;
   vector<int> Q;
```

```
Dinic(int N) : N(N), G(N), dad(N), Q(N) {}
    void AddEdge(int from, int to, int cap) {
       G[from].push_back(Edge(from, to, cap, 0, G[to].size()));
        if (from == to) G[from].back().index++;
        G[to].push_back(Edge(to, from, 0, 0, G[from].size() - 1));
    long long BlockingFlow(int s, int t) {
        fill(dad.begin(), dad.end(), (Edge *) NULL);
        dad[s] = &G[0][0] - 1;
        int head = 0. tail = 0:
       Q[tail++] = s;
        while (head < tail) {
           int x = O[head++];
           for (int i = 0; i < G[x].size(); i++) {
               Edge &e = G[x][i];
               if (!dad[e.to] && e.cap - e.flow > 0) {
                   dad[e.to] = &G[x][i];
                   Q[tail++] = e.to;
        if (!dad[t]) return 0;
        long long totflow = 0;
        for (int i = 0; i < G[t].size(); i++) {
           Edge *start = \&G[G[t][i].to][G[t][i].index];
            int amt = INF;
           for (Edge *e = start; amt \&\& e != dad[s]; e = dad[e->from]) {
               if (!e) { amt = 0; break; }
               amt = min(amt, e->cap - e->flow);
            if (amt == 0) continue:
            for (Edge *e = start; amt && e != dad[s]; e = dad[e->from]) {
               e->flow += amt;
               G[e->to][e->index].flow -= amt;
           totflow += amt:
        return totflow:
    long long GetMaxFlow(int s, int t) {
        long long totflow = 0;
        while (long long flow = BlockingFlow(s, t))
            totflow += flow;
        return totflow;
};
    MaximumFlow-EdmondsKarp.cc
typedef double T; // also works for doubles, but use feq, etc
const T INFTY = numeric_limits<T>::max(); // ::infinity() for doubles
typedef vector<size_t> VI;
typedef vector<VI>
typedef vector<T>
                       VT:
typedef vector<VT>
                      VVT;
// Edmonds-Karp algorithm for max-flow
// Runs in O(VE^2). Alternate complexity: O(f(V+E)),
// where f is the value of the maximum flow
struct edmondskarp_graph {
    size t N:
    VVI A;
    WVT C, F; // references to F can be removed if you don't want flows
    edmondskarp_graph( size_t _N ) : N(_N), A(N), C(N,VT(N,0)), F(N,VT(N,0))
    void add_capacity( size_t s, size_t t, T cap ) {
        if( cap == 0 ) return;
```

if(C[s][t] == 0 && C[t][s] == 0) {

A[s].push_back(t);

A[t].push_back(s);

```
C[s][t] += cap;
       // If you subtract capacities, and want to remove edges with cap 0,
       // do so here, or afterward.
   T Augment( const VI &P ) {
       T amt = INFTY;
       FOR(i,0,P.size()-1) amt = min(amt, C[ P[i] ][ P[i+1] ]);
       FOR(i,0,P.size()-1) {
           size_t u = P[i], v = P[i+1];
           C[u][v] -= amt;
           C[v][u] += amt;
           if( F[v][u] > amt ) {
               F[v][u] -= amt;
           else {
               F[u][v] += amt - F[v][u];
               F[v][u] = 0;
       return amt:
   bool bfs( size_t s, size_t t, VI &P ) {
       P = VI(N,N):
       VI Q(N); size_t qh=0, qt=0;
       P[Q[qt++] = s] = s;
       while( qh < qt \&\& P[t] == N ) {
           size_t c = Q[qh++];
           FOR(i,0,A[c].size()) {
               size_t u = A[c][i];
               if( C[c][u] == 0 ) continue;
               if( P[u] != N ) continue;
               P[Q[qt++] = u] = c;
       7
       return P[t] != N;
   T ComputeMaxFlow( size_t s, size_t t ) {
       T flow = 0;
       while(bfs(s,t,P)) {
           VI path(1,t);
           size t z = t:
           while( z != P[z] ) path.push_back( z = P[z] );
           path = VI(path.rbegin(), path.rend());
           flow += Augment(path);
       return flow;
   }
};
```

MinCostBipartiteMatching.cc

```
/// Min cost bipartite matching VIa shortest augmenting paths
// This is an O(n^3) implementation of a shortest augmenting path
// algorithm for finding min cost perfect matchings in dense
// graphs. In practice, it solves 1000x1000 problems in around 1
// second.
//
// cost[i][j] = cost for pairing left node i with right node j
// Lmate[i] = index of right node that left node i pairs with
// Rmate[j] = index of left node that right node j pairs with
// The values in cost[i][j] may be positive or negative. To perform
// maximization, simply negate the cost[][] matrix.
///
// typedef vector<double> VD;
typedef vector<int> VID;
typedef vector<int> VI;
double MinCostMatching(const VVD &cost, VI &Lmate, VI &Rmate) {
    int n = int(cost.size());
```

```
// construct dual feasible solution
VD u(n):
VD v(n);
for (int i = 0; i < n; i++) {
    u[i] = cost[i][0];
    for (int j = 1; j < n; j++) u[i] = min(u[i], cost[i][j]);</pre>
for (int j = 0; j < n; j++) {
    v[j] = cost[0][j] - u[0];
    for (int i = 1; i < n; i++) v[j] = min(v[j], cost[i][j] - u[i]);
// construct primal solution satisfying complementary slackness
Lmate = VI(n, -1);
Rmate = VI(n, -1);
int mated = 0;
for (int i = 0; i < n; i++) {
    for (int j = 0; j < n; j++) {
        if (Rmate[j] != -1) continue;
        if (fabs(cost[i][j] - u[i] - v[j]) < 1e-10) {
            Lmate[i] = j;
            Rmate[j] = i;
            mated++;
            break:
        }
    }
VD dist(n):
VI dad(n):
VI seen(n):
// repeat until primal solution is feasible
while (mated < n) {
    // find an unmatched left node
    int s = 0:
    while (Lmate[s] != -1) s++;
    // initialize Dijkstra
    fill(dad.begin(), dad.end(), -1);
    fill(seen.begin(), seen.end(), 0);
    for (int k = 0: k < n: k++)
    dist[k] = cost[s][k] - u[s] - v[k];
    int i = 0:
    while (true) {
        // find closest
        i = -1:
        for (int k = 0; k < n; k++) {
            if (seen[k]) continue;
            if (j == -1 || dist[k] < dist[j]) j = k;
        seen[i] = 1;
        // termination condition
        if (Rmate[j] == -1) break;
        // relax neighbors
        const int i = Rmate[j];
        for (int k = 0; k < n; k++) {
            if (seen[k]) continue;
            const double new_dist = dist[j] + cost[i][k] - u[i] - v[k];
            if (dist[k] > new_dist) {
                dist[k] = new_dist;
                dad[k] = j;
           }
       }
    // update dual variables
    for (int k = 0; k < n; k++) {
        if (k == j || !seen[k]) continue;
        const int i = Rmate[k];
        v[k] += dist[k] - dist[j];
        u[i] -= dist[k] - dist[j];
```

```
u[s] += dist[i]:
       // augment along path
       while (dad[j] >= 0) {
           const int d = dad[j];
           Rmate[j] = Rmate[d];
           Lmate[Rmate[j]] = j;
           j = d;
       Rmate[j] = s;
       Lmate[s] = j;
       mated++;
    double value = 0;
   for (int i = 0; i < n; i++)
   value += cost[i][Lmate[i]];
   return value:
    MinCostMaxFlow.cc
// Min-cost Maximum Flow. The implementation has a few stages, some of which
// can be outright ignored if the graph is guaranteed to satisfy certain
// conditions. These are documented below.
#define FOR(v,l,u) for (size_t v = l; v < u; ++v)
typedef signed long long int T; // the basic type of costs and flow.
typedef vector<T>
                            VT:
struct edge {
   size_t s, t;
   T cap, flow, cost; // Note: cap is *residual* capacity.
   size_t di; // index of dual in t's edgelist.
   edge *dual; // the actual dual (see "compile_edges")
typedef vector<edge> VE;
typedef vector<VE>
                      VVE;
typedef vector<size_t> VI;
typedef pair<T,size_t> DijkP; // Dijkstra PQ element.
typedef priority_queue<DijkP, vector<DijkP>, std::greater<DijkP> > DijkPQ;
struct mcmf_graph {
   size_t N; VVE adj; VT pot;
   mcmf_graph( size_t N ) : N(N), adj(N), pot(N,0) {}
   void add_edge( size_t s, size_t t, T cap, T cost ) {
       edge f. r:
       f.s = s; f.t = t; f.cap = cap; f.flow = 0; f.cost = cost;
       r.s = t; r.t = s; r.cap = 0; r.flow = 0; r.cost = -cost;
       f.di = adj[t].size(); r.di = adj[s].size();
       adj[s].push_back(f); adj[t].push_back(r);
    void compile edges() {
       FOR(v,0,N) FOR(i,0,adj[v].size()) { // This has to be done after all
           edge &e = adj[v][i]; // edges are added because vectors can
           e.dual = &adj[e.t][e.di]; // resize and move their contents.
   T Augment( const VE &path ) {
       T push = path[0].cap;
       FOR(i,0,path.size()) push = min(push, path[i].cap);
       FOR(i,0,path.size()) {
           edge &e = *(path[i].dual->dual); // the actual edge, not a copy
           e.cap -= push; e.dual->cap += push;
           if( e.dual->flow >= push ) e.dual->flow -= push;
           else { e.flow += push - e.dual->flow; e.dual->flow = 0; }
       return push;
   void ApplyPotential( const VT &delta ) {
       FOR(v.O.N) {
           FOR(i,0,adj[v].size()) {
               adi[v][i].cost += delta[v];
                adj[v][i].dual->cost -= delta[v];
```

pot[v] += delta[v];

```
/* The following, down to "CancelNegativeCycles", are unnecessary if the
 * graph is guaranteed to have no negative cycles.
 * Alternatively, if you compute any maxflow, you can include these, and
 * run CancelNegativeCycles to find a cost-optimal maxflow. */
bool dfs_negcycle_r( const size_t rt, VI &par, VE &cycle ) {
    FOR(i,0,adj[rt].size()) {
        edge &e = adj[rt][i];
        if( e.cap == 0 || e.cost >= 0 ) continue;
        size_t v = e.t;
        if( par[v] < N ) { // found a negative cycle!
            size_t fr = 0; while( cycle[fr].s != v ) ++fr;
            cycle = VE( cycle.begin()+fr, cycle.end() );
            cycle.push_back(e);
            return true;
        else if( par[v] == N ) { // unvisited node
            par[v] = rt; cycle.push_back(e);
            if ( dfs_negcycle_r(v,par,cycle) ) return true;
            par[v] = N+1; cycle.pop_back();
    return false;
bool dfs_negcycle( VE &cycle ) {
    cycle.clear(); VI par(N,N);
    FOR(v,0,N) if( par[v] == N \&\& dfs_negcycle_r(v,par,cycle)) return
     \hookrightarrow true;
    return false:
void CancelNegativeCycles() { // only if the graph has negative cycles
    while( dfs_negcycle(cycle) )
       Augment(cycle);
/* The following is unnecessary if the graph is guaranteed to have no
 * negative-cost edges with positive capacity before MCMF is run. */
void FixNegativeEdges( size_t SRC ) {
    VT W(N); VI P(N,N); P[SRC] = 0;
    FOR(kk,0,N-1) {
       FOR(v,0,N) FOR(i,0,adj[v].size()) {
            if( adj[v][i].cap == 0 ) continue;
            size_t u = adj[v][i].t; T w = adj[v][i].cost;
           if( P[u] == N | | W[v]+w < W[u] ) {
               W[u] = W[v] + w;
               P[u] = v;
       }
    ApplyPotential(W);
/* The following form the crux of min-cost max-flow, unless you go with
 * pure cycle-canceling approach by precomputing a maxflow. */
void shortest_paths( size_t S, VE &P, VT &W ) {
    DijkPQ Q; P = VE(N); W = VT(N,0); // DO init everything to 0!
    FOR(i,0,N) P[i].s = N; edge x; x.s = x.t = S;
    Q.push(DijkP(0,0)); VE trv; trv.push_back(x);
    while( !Q.empty() ) {
       T wt = Q.top().first; edge e = trv[Q.top().second];
        size_t v = e.t;
                               Q.pop();
        if( P[v].s != N ) continue;
        W[v] = wt; P[v] = e;
        FOR(i,0,adj[v].size()) {
            edge &f = adj[v][i];
            if(f.cap == 0) continue;
            Q.push(DijkP(W[v]+f.cost, trv.size()));
            trv.push_back(f);
// Note that this returns the total *maximum flow*, not its cost. Use
// "Cost()" after calling this for that.
T ComputeMinCostMaxFlow( size_t SRC, size_t DST ) {
    compile_edges(); // we have to do this after all edges are added
    CancelNegativeCycles(); // Only if necessary!
    FixNegativeEdges( SRC ); // Ditto!
```

T flow = 0; VE P; VT W; shortest_paths(SRC, P, W);

```
while( P[DST] .s != N ) { // while there is a path S->T
            for( size_t v = DST; v != SRC; v = P[v].s ) ap.push_back(P[v]);
            ap = VE( ap.rbegin(), ap.rend() ); // I love C++ sometimes
            flow += Augment( ap );
            ApplyPotential( W ); // This eliminates negative cycles from ^
            shortest_paths( SRC, P, W );
        return flow:
   T Cost() {
        T c = 0:
        FOR(v,0,N) FOR(i,0,adj[v].size()) {
            edge &e = adj[v][i];
            c += e.flow * (e.cost - pot[e.s] + pot[e.t]);
        return c;
}:
    SCC.cc
// An implementation of Kosaraju's algorithm for strongly-connected
// This includes code which constructs a "meta" graph with one node per SCC.
#define FOR(v,l,u) for (size_t v = l; v < u; ++v)
typedef vector<size_t> VI;
typedef vector<VI>
                      VVT:
typedef vector<bool> VB:
struct graph {
   size_t N;
    VVI A; // Adjacency lists.
          B; // Reversed adjacency lists.
   VI scc; // scc[i] is the component to which i belongs
    size_t n_sccs; // the number of components
   graph( size_t n ) : N(n), A(n), B(n), scc(n) {}
    void add_edge( size_t s, size_t t ) {
        A[s].push_back(t);
        B[t].push_back(s);
   bool has_edge( size_t s, size_t t ) { // only for compute_scc_graph
        FOR(i,0,A[s].size()) if( A[s][i] == t ) return true;
    void dfs_order( size_t rt, VB &Vis, VI &order ) {
        Vis[rt] = true:
        FOR(i,0,A[rt].size()) {
            size_t v = A[rt][i];
            if ( Vis[v] ) continue:
            dfs_order( v, Vis, order );
        order.push_back(rt);
    void dfs_label( size_t rt, VB &Vis, size_t lbl, VI &out ) {
        Vis[rt] = true;
        out[rt] = lbl:
        FOR(i,0,A[rt].size()) {
            size_t v = A[rt][i];
            if( Vis[v] ) continue;
            dfs_label( v, Vis, lbl, out );
       }
    void compute sccs() {
        VB visited(N,false); VI order;
        FOR(v,0,N) if( !visited[v] ) dfs_order(v, visited, order);
        swap(A,B);
        visited = VB(N,false); n_sccs = 0;
        FOR(i,0,N) {
            size t v = order[N-1-i]:
            if( !visited[v] ) dfs_label(v, visited, n_sccs++, scc);
        swap(A,B);
    void compute_scc_graph( graph &H ) {
        H = graph(n_sccs);
        FOR(v,0,N) {
```

```
FOR(i,0,A[v].size()) {
                size_t u = A[v][i];
               size_t vv = scc[v], uu = scc[u];
               if( vv != uu && !H.has_edge(vv,uu) )
                   H.add_edge(vv,uu);
       }
   }
    UnionFind.cc
// UnionFind data structure that implements
// path compression, stolen from stanfordacm
// should not stackoverflow in correcty configured environment
// (ulimit -s BIGNUMBER)
typedef vector<size_t> VI;
// Union find is now a vector of integers, C[i] = parent(i).
// Initialize with C[i] = i
int find(VI &C, size_t x) { return (C[x] == x) ? x : C[x] = find(C, C[x]); }
void merge(VI &C, size_t x, size_t y) { C[find(C, x)] = find(C, y); }
    Kruskal.cc
// Kruskal's algorithm to return MST using
// Union-Find data structure.
typedef pair<size_t, size_t> ii;
typedef pair < double, ii > dii;
typedef vector<dii> vdii;
// edges is a list of all edges in the graph, n is number
// of vertices in the graph
double kruskal(vdii &edges, size_t n) {
    sort(edges.begin(), edges.end());
    for (size_t i = 0; i < n; ++i)
       uf[i] = i:
    double cost = 0;
    for (size_t i = 0; i < edges.size(); ++i) {</pre>
        size_t u = edges[i].second.first;
        size_t v = edges[i].second.second;
        if (find(uf, u) != find(uf, v)) {
           merge(uf, u, v);
            cost += edges[i].first;
            // if MST edges used are needed, add them here
   }
    return cost;
    LCA.cc
// Lowest Common Ancestor on a weighted, rooted tree.
// Solves UVa Online Judge "AntsColony" in 2.04 seconds.
// O(n log n) time and space precomputation, O(log n) LCA query.
typedef vector<int> VI;
typedef vector<VI> VVI;
typedef pair<int, int> II;
typedef vector<II> VII;
typedef vector<VII> VVII;
// computes an integer upper bound of the binary logarithm of n
int log2(int n) {
   int j;
    for (j = 0; (1 << j) < n; ++j);
```

```
// computes the distance from the root to every node (not necessary for LCA)
// and the level of each node in the tree (necessary for LCA)
// expects an adjacency list of the tree, where each entry is (idx, weight)
void get_dist(VVII &tree, VI &dist, VI &L) {
   stack<int> s;
   s.push(0);
   dist[0] = 0;
   L[0] = 0;
   while (!s.empty()) {
       int cur = s.top(); s.pop();
for (int i = 0; i < tree[cur].size(); ++i) {</pre>
            dist[tree[cur][i].first] = dist[cur] + tree[cur][i].second;
            s.push(tree[cur][i].first);
            L[tree[cur][i].first] = L[cur]+1;
   }
// uses dynamic programming to preprocess the P table
void preprocess(VVI &P, int N) {
   for (int j = 1; (1 << j) < N; ++j) {
       for (int i = 0; i < N; ++i) {
           if (P[i][j-1] != -1) {
                P[i][j] = P[P[i][j-1]][j-1];
   }
// comptues the LCA of p and q, given P, L, and N
int LCA(int p, int q, VVI &P, VI &L, int N) {
   if (L[p] < L[q])
       swap(p, q);
   for (int i = log2(N)-1; i >= 0; --i) {
       if (P[p][i] != -1 && L[P[p][i]] >= L[q])
            p = P[p][i];
   if (p == q)
       return p:
   for (int i = log2(N)-1; i >= 0; --i) {
       if (P[p][i] != P[q][i]) {
           p = P[p][i], q = P[q][i];
   }
   return P[p][0];
    FloatCompare.cc
// Short function for comparing floating point numbers.
const double EPS_ABS = 1e-10; // for values near 0.0. Keep small.
const double EPS_REL = 1e-8; // for values NOT near 0.0. Balance.
bool feq( double a, double b ) {
   double d = fabs(b-a);
   if( d <= EPS_ABS ) return true;</pre>
   if( d <= max(fabs(a),fabs(b))*EPS_REL ) return true;</pre>
   return false;
bool flt( double a, double b ) {
   return !feq(a,b) && a < b;
```

Vector.cc

MinCostMaxFlow.cc

```
// A simple library used elsewhere in the notebook.
                                                                                    return fabs(a*x+b*y+c*z-d)/sqrt(a*a+b*b+c*c);
// Provides basic vector/point operations.
typedef double T:
                                                                                // Decide if lines ab and cd are parallel.
                                                                                // If a=b or c=d, then this will return true.
struct Pt {
                                                                                bool LinesParallel( Pt a, Pt b, Pt c, Pt d ) {
   Тх, у;
                                                                                   return feq( cross(b-a,c-d), 0 );
   Pt( T x, T y ) : x(x), y(y) {}
                                                                                // Decide if lines ab and cd are the same line
   Pt( const Pt &h ) : x(h.x), y(h.y) {}
                                                                                // If a=b and c=d, then this will return true.
                                                                                // If a=b xor c=d, (wlog a=b), then this is true iff a is on cd.
                                                                                bool LinesColinear( Pt a, Pt b, Pt c, Pt d ) {
Pt operator + ( const Pt &a, const Pt &b ) { return Pt(a.x+b.x, a.y+b.y); }
                                                                                   return LinesParallel(a,b, c,d)
Pt operator - ( const Pt &a, const Pt &b ) { return Pt(a.x-b.x, a.y-b.y); }
                                                                                        && isLeft(a,b, c) == 0
Pt operator * ( const T s, const Pt &a ) { return Pt(s*a.x, s*a.y); }
                                                                                        && isLeft(c,d, a) == 0; // to make a=b, c=d cases symmetric
Pt operator * ( const Pt &a, const T s ) { return s*a; }
Pt operator / ( const Pt &a, const T s ) { return Pt(a.x/s,a.y/s); }
                                                                                // Determine if the segment ab intersects with segment cd
// Note the kind of division that occurs when using integer types.
                                                                                // Use line-line intersection (below) to find it.
                                                                                // This *will* do the right thing if a=b, c=d, or both!
// Use rationals if you want this to work right.
bool operator == ( const Pt &a, const Pt &b ) {
                                                                                bool SegmentsIntersect( Pt a, Pt b, Pt c, Pt d ) {
                                                                                    if( LinesColinear(a,b, c,d) ) {
   return feq(a.x,b.x) && feq(a.y,b.y);
                                                                                        if( a==c || a==d || b==c || b==d ) return true;
                                                                                        if( dot(a-c,b-c) > 0 && dot(a-d,b-d) > 0 && dot(c-b,d-b) > 0 )
bool operator != ( const Pt &a, const Pt &b ) { return !(a == b); }
                                                                                            return false:
T dot( const Pt &a, const Pt &b ) { return a.x*b.x + a.v*b.v; }
                                                                                        return true:
T cross( const Pt &a, const Pt &b ) { return a.x*b.y - a.y*b.x; }
T norm2( const Pt &a )
                                  { return a.x*a.x + a.y*a.y; } //
                                                                                    if( isLeft(a,b, d) * isLeft(a,b, c) > 0 ) return false;
 \hookrightarrow dot(a, a)
                                                                                    if( isLeft(c,d, a) * isLeft(c,d, b) > 0 ) return false;
T norm( const Pt &a )
                                   { return sqrt(a.x*a.x + a.y*a.y); }
T dist2( const Pt &a, const Pt &b ) { // dot(a-b,a-b)
   T dx = a.x - b.x, dy = a.y - b.y;
                                                                                // Determine if c is on the segment ab
   return dx*dx + dy*dy;
                                                                                bool PointOnSegment( Pt a, Pt b, Pt c )
                                                                                    { return SegmentsIntersect(a,b,c,c); }
T dist( const Pt &a, const Pt &b ) { // sqrt(dot(a-b,a-b))
                                                                                // Compute the intersection of lines ab and cd.
                                                                                // ab and cd are assumed to be *NOT* parallel
   T dx = a.x - b.x, dy = a.y - b.y;
   return sqrt(dx*dx + dy*dy);
                                                                               Pt ComputeLineIntersection( Pt a, Pt b, Pt c, Pt d ) {
                                                                                    b=b-a; d=d-c; c=c-a; // translate to a, set b,d to directions
                                                                                    return a + b*cross(c,d)/cross(b,d); // solve s*b = c + t*d by Cramer
bool lex_cmp_xy( const Pt &lhs, const Pt &rhs ) {
   if( !feq(lhs.x,rhs.x) ) return lhs.x < rhs.x;</pre>
                                                                                // Compute the center of the circle uniquely containing three points.
   if( !feq(lhs.y,rhs.y) ) return lhs.y < rhs.y;</pre>
                                                                                // It's assume the points are *NOT* colinear, so check that first.
                                                                                Pt ComputeCircleCenter(Pt a, Pt b, Pt c) {
   return false:
                                                                                    b=(b-a)/2; c=(c-a)/2; // translate to a=origin, shrink to midpoints
                                                                                    return a+ComputeLineIntersection(b,b+RotateCW90(b),c,c+RotateCW90(c));
    PlaneGeometry.cc
                                                                                // Compute intersection of line ab with circle at c with radius r.
                                                                                // This assumes a!=b.
                                                                               VP CircleLineIntersection( Pt a, Pt b, Pt c, T r ) {
// Some routines for basic plane geometry.
                                                                                    VP ret;
// Depends on Vector.cc.
                                                                                    b = b-a; a = a-c; // translate c to origin, make b the direction
typedef vector<Pt> VP;
                                                                                                         // Let P(t) = a + t*b, and Px, Py projections
                                                                                    T A = dot(b,b):
                                                                                                         // Solve Px(t)^2 + Py(t)^2 = r^2
                                                                                   T B = dot(a,b):
int isLeft( Pt a, Pt b, Pt c ) {
                                                                                    T C = dot(a,a) - r*r; // Get A*t^2 + 2B*t + C = 0
   T z = cross(b-a,c-a);
                                                                                    T D = B*B - A*C; //4*D is the discriminant
   if (feq(z,0)) return 0; //c is on the line ab
                                                                                    if( flt(D,0) ) return ret;
   else if( z > 0 ) return 1; // c is left of the line ab
                                                                                    D = sqrt(max((T)0,D));
                       return -1: // c is right of the line ab
                                                                                    ret.push_back( c+a + b*(-B + D)/A );
                                                                                    if( feq(D,0) ) return ret;
Pt RotateCCW90( Pt p ) { return Pt(-p.y,p.x); }
                                                                                    ret.push_back( c+a + b*(-B - D)/A );
Pt RotateCW90(Pt p) { return Pt(p.y,-p.x); }
Pt RotateCCW( Pt p, T t ) { // This only makes sense for T=double
   return Pt(p.x*cos(t)-p.y*sin(t), p.x*sin(t)+p.y*cos(t));
                                                                                // Compute intersection of circle at a with radius r
                                                                                // with circle at b with radius s.
// Project the point c onto line ab
                                                                                // This assumes the circles are distinct, ie (a,r)!=(b,s)
// This assumes a != b, so check that first.
                                                                                VP CircleCircleIntersection( Pt a, T r, Pt b, T s ) {
Pt ProjectPointLine( const Pt &a, const Pt &b, const Pt &c ) {
   return a + (b-a) * dot(b-a,c-a)/norm2(b-a);
                                                                                    T d = dist(a, b);
                                                                                    if( d > r+s || d+min(r.s) < max(r.s) ) return ret: // emptu
// "Project" the point c onto segment ab
                                                                                    T x = (d*d-s*s+r*r)/(2*d); // The rest of this is magic.
// Nicely paired with dist: dist(c, ProjectPointSegment(a,b,c))
                                                                                    T y = sqrt(r*r-x*x);
                                                                                                                // (It's actually basic geometry.)
Pt ProjectPointSegment( Pt a, Pt b, Pt c ) {
                                                                                    Pt. v = (b-a)/d
   T r = dist2(a,b);
                                                                                    ret.push_back(a+v*x + RotateCCW90(v)*y);
   if( feq(r,0) ) return a;
                                                                                    if( !feq(y,0) )
   r = dot(c-a, b-a)/r;
                                                                                        ret.push_back(a+v*x - RotateCCW90(v)*y);
   if(r < 0) return a;
                                                                                    return ret;
```

if(r > 1) return b:

// Compute the distance between point (x,y,z) and plane ax+by+cz=d

T DistancePointPlane(T x, T y, T z, T a, T b, T c, T d) {

return a + (b-a)*r:

```
// Basic routines for polygon-related stuff.
// Uses Vector.cc and PlaneGeometry.cc.
// Polygons are just vector < Pt>'s.
#define FOR(v,l,u) for (size_t v = l; v < u; ++v)
typedef vector<Pt> VP;
// These generalize to higher-dimensional polyhedra, provided you represent
// them as a collection of facets
// Just replace "cross" with the suitable determinant, and adjust any
// factors.
T ComputeSignedArea( const VP &p ) {
   T area = 0:
   for( size_t i = 0; i < p.size(); i++ ) {
       size_t z = (i + 1) % p.size();
        area += cross( p[i], p[z] );
   return area / 2.0;
T ComputeArea( const VP &p ) {
   return fabs(ComputeSignedArea(p));
T ComputePerimeter( const VP &p ) {
   T perim = 0.0:
   for( size_t i = 0; i < p.size(); ++i )</pre>
       perim += dist(p[i], p[(i+1) % p.size()]);
    return perim:
Pt ComputeCentroid( const VP &p ) {
   Pt c(0,0); T scale = 6.0 * ComputeSignedArea(p);
   for( size_t i = 0; i < p.size(); i++ ) {
       size_t j = (i + 1) % p.size();
       c = c + cross(p[i],p[j]) * (p[i]+p[j]);
   return c / scale;
bool IsSimple( const VP &p ) {
   for( size_t i = 0; i < p.size(); ++i )</pre>
   for( size_t k = i+1; k < p.size(); ++k ) {</pre>
       size_t j = (i + 1) % p.size();
       size_t 1 = (k + 1) % p.size();
        if( i == 1 \mid \mid j == k ) continue;
        if( SegmentsIntersect(p[i],p[j], p[k],p[l]) )
           return false:
   return true:
// Determine the winding number of a point. This is the number of
// times the polygon goes around the given point.
// It is 0 exactly when the point is outside.
// A signed type is used intermediately so that we don't have to
// detect CW versus CCW, but the absolute value is taken in the end.
// If q is *on* the polygon, then the results are not well-defined,
// since it depends on whether q is on an "up" or "down" edge.
size_t WindingNumber( const VP &p, Pt q) {
   int wn = 0; vector<int> state(p.size()); // state decides up/down
   FOR(i,0,p.size())
       if( feq(p[i].y, q.y) ) state[i] = 0; // break ties later
        else if(p[i].y < q.y) state[i] = -1; // we'll use nearest
                               state[i] = 1; // neighbor (either)
   FOR(i,1,p.size()) if( state[i] == 0 ) state[i] = state[i-1];
   if( state[0] == 0 ) state[0] = state.back();
   FOR(i,1,p.size()) if( state[i] == 0 ) state[i] = state[i-1];
   FOR(i,0,p.size()) {
       size_t z = (i + 1) % p.size();
        if( state[z] == state[i] ) continue; // only interested in changes
        else if( state[z] == 1 && isLeft(p[i],p[z],q) > 0 ) ++wn;
        else if( state[i] == 1 && isLeft(p[i],p[z],q) < 0 ) --wn;
   return (size_t)(wn < 0 ? -wn : wn);
// A complement to the above.
bool PointOnPolygon( const VP &p, Pt q ) {
   for( size_t i = 0; i < p.size(); i++ ) {
       size_t z = (i + 1) % p.size();
        if( PointOnSegment(p[i],p[z],q) )
```

Vector.cc Polygon.cc

Polygon.cc

```
return true:
   return false:
// Convex hull.
// This *will* modify the given VP. To save your points, do
// { VP hull(p.begin(),p.end()); ConvexHull(hull); }
// This *will* keep redundant points on the polygon border.
// To ignore those, change the isLeft's < and > to <= and >=.
void ConvexHull( VP &Z ) {
    sort( Z.begin(), Z.end(), lex_cmp_xy);
   Z.resize( unique(Z.begin(),Z.end()) - Z.begin() );
   if(7.size() < 2) return:
    for( size_t i = 0; i < Z.size(); i++ ) {</pre>
        while(up.size() > 1 && isLeft(up[up.size()-2],up.back(),Z[i]) > 0)
            up.pop back():
        while(dn.size() > 1 \& isLeft(dn[dn.size()-2],dn.back(),Z[i]) < 0)
            dn.pop_back();
        up.push_back(Z[i]);
        dn.push_back(Z[i]);
   Z = dn;
   for( size t i = up.size() - 2; i >= 1; i-- ) Z.push back(up[i]);
// Implementation of Sutherland-Hodgman algorithm:
// https://en.wikipedia.org/wiki/Sutherland-Hodgman_algorithm
// Computes the intersection of polygon subject and polygon clip.
// Polygons points must be given in clockwise order. Clip must be convex.
// May return repeated points, especially if intersection is a single point

    or line segment.

// If no intersection occurs, will return an empty vector.
// Undefined behavior if intersection consists of multiple polygons.
VP ConvexClipPolygon( const VP &subject, const VP &clip ) {
    VP output = subject;
    for (size_t i = 0; i < clip.size(); ++i) {
        size_t ip1 = (i+1)%clip.size();
        Pt EdgeStart = clip[i];
        Pt EdgeEnd = clip[ip1];
        VP input = output;
        output.clear():
        Pt S = input.back();
        for (size_t j = 0; j < input.size(); ++j) {</pre>
            Pt E = input[i]:
            if (isLeft(EdgeStart, EdgeEnd, E) <= 0) {
                if (isLeft(EdgeStart, EdgeEnd, S) > 0) {
                    output.push_back(ComputeLineIntersection(EdgeStart,
                     }
               output.push_back(E);
            else if (isLeft(EdgeStart, EdgeEnd, S) <= 0) {</pre>
                output.push_back(ComputeLineIntersection(EdgeStart, EdgeEnd,
                \hookrightarrow S, E));
           S = E:
    return output;
    KDtree.cc
// Fully dynamic n-dimensional sledgehammer kd-tree.
// Constructs 100,000 point kd tree in about 2 seconds
// Handles 100.000 2D NN searches on 100.000 points in about 2 seconds
// Handles 1,000 2D range queries of ranges "O(sqrt(n)) points
// on 50,000 points in about 2 seconds
// May degenerate when points are not uniformly distributed
// I hope you don't implement the whole thing.
// use to change data structure of pts in kdtree
typedef int T:
typedef vector<T> VT;
```

typedef vector<VT*> VVT;
typedef vector<VVT> VVVT;

```
struct kdnode {
    size_t d;
    kdnode *left:
    kdnode *right;
    // number of alive nodes in subtree rooted at this node,
    // including this node if alive
    size_t nAlive;
    // number of flagged dead nodes in subtree rooted at this node.
    // including this node if dead
    size_t nDead;
    // is flagged or not
    hool isAlive:
    VT *pt:
    // computes distance in n-dimensional space
    double dist(VT &pt1, VT &pt2) {
       double retVal = 0:
        for (size_t i = 0; i < pt1.size(); ++i) {
           retVal += (pt1[i]-pt2[i]) * (pt1[i]-pt2[i]);
        return sqrt(retVal);
    // returns closer point to apt
    VT * minPt(VT &qpt, VT *pt1, VT *pt2) {
        if (pt1 == NULL) return pt2;
        if (pt2 == NULL) return pt1;
        return dist(*pt1, qpt) < dist(*pt2, qpt) ? pt1 : pt2;
    // find median based on chosen algorithm
    T findMed(VVT &pts, size_t d) {
        VT arr(pts.size());
        for (size_t i = 0; i < pts.size(); ++i)</pre>
           arr[i] = (*pts[i])[d];
        sort(arr.begin(), arr.end());
        return arr[arr.size()/2]:
    void printPt(VT &pt) {
        for (size_t i = 0; i < pt.size(); ++i)</pre>
           cout << pt[i] << " ";
    // intersects orthogonal region with left or right
    // of orthogonal halfspace on dth dimension
    VT region_intersect(VT region, T line, size_t d, bool goLeft) {
        if (goLeft) {
           region[d*2+1] = line;
        else {
            region[d*2] = line;
        return region:
    // returns true iff the entire region is contained in the given range
    bool region_contained(VT &region, VT &range) {
        for (size_t i = 0; i < region.size(); ++i) {</pre>
           if (i % 2 == 0) {
               if (region[i] < range[i])
                    return false:
           }
           else {
                if (region[i] > range[i])
                    return false;
        return true:
```

```
}
// returns true if point is in range
bool pt_contained(VT *pt, VT &range) {
    for (size_t i = 0; i < pt->size(); ++i) {
        if ((*pt)[i] < range[i*2] || (*pt)[i] > range[i*2+1])
            return false;
    }
    return true:
// creates "infinite" region, unbounded on all dimensions
VT infRegion(size_t d) {
    VT region(d*2);
    for (size_t i = 0; i < region.size(); ++i) {</pre>
        if (i % 2 == 0)
            region[i] = -INF;
            region[i] = INF;
    return region:
}
void build_tree(VVT &pts, size_t d, size_t num_d) {
    this->d = d;
    nAlive = pts.size();
    nDead = 0:
    isAlive = true;
    VVT leftV, rightV;
    T med = findMed(pts, d);
    for (size_t i = 0; i < pts.size(); ++i) {</pre>
        if ((*pts[i])[d] == med && pt == NULL)
            pt = pts[i];
         else if ((*pts[i])[d] <= med)
            leftV.push_back(pts[i]);
            rightV.push_back(pts[i]);
    left = leftV.empty() ? NULL : new kdnode(leftV, (d+1)%num_d, num_d);
    right = rightV.empty() ? NULL : new kdnode(rightV, (d+1)%num_d,
     \hookrightarrow num_d);
// constructs kd tree
kdnode(VVT %pts, size_t d, size_t num_d) {
    build_tree(pts, d, num_d);
// adds pt to tree
void addPt(VT *newPt) {
    ++nAlive:
    bool goLeft = (*newPt)[d] <= (*pt)[d];</pre>
    kdnode *child = goLeft ? left : right;
    size_t childCt = (child == NULL ? 0 : child->nAlive) + 1;
     // rebuild
    if (childCt > (1+ALPHA)/2 * nAlive) {
        VVT allPts:
        addPtToResult(allPts):
        allPts.push_back(newPt);
        delete left; delete right;
        build_tree(allPts, d, pt->size());
    else if (child == NULL) {
        // add node
        VVT ptV(1, newPt);
```

```
if (goLeft)
            left = new kdnode(ptV, (d+1)%pt->size(), pt->size());
        else
            right = new kdnode(ptV, (d+1)%pt->size(), pt->size());
    else {
        // recurse
        child->addPt(newPt);
// deletes existing point from kd-tree, rebalancing if necessary
// returns the number of dead nodes removed from this subtree,
// and bool for if pt found both are necessary in this implementation
// to retain proper balancing invariants
pair<size_t, bool> deletePt(VT *oldPt) {
    ++nDead:
    --nAlive;
    // need to reconstruct - last part is to avoid
    // an empty tree construction. Will get picked up by parent later
    if (nAlive < (1.0-ALPHA) * (nAlive + nDead) && nAlive > 0) {
        VVT allPts;
        addPtToResult(allPts):
        bool found = false;
        for (size_t i = 0; i < allPts.size(); ++i) {</pre>
            if (*allPts[i] == *oldPt) {
                found = true:
                allPts.erase(allPts.begin() + i);
                break;
           }
        delete left; delete right;
        size_t deadRemoved = nDead;
        build_tree(allPts, d, pt->size());
        return make_pair(deadRemoved, found);
    else if (*pt == *oldPt) { // base case, point found
        isAlive = false:
        return make_pair(0, true);
    else {
        bool goLeft = (*oldPt)[d] <= (*pt)[d];</pre>
        kdnode *child = goLeft ? left : right;
        size_t deadRemoved = 0;
        bool found = false;
        if (child != NULL) {
            // recurse
            pair<size_t, bool> result = child->deletePt(oldPt);
            deadRemoved = result.first;
            found = result second:
        // point may not have been found
        if (!found)
            ++nAlive:
```

```
nDead -= deadRemoved;
        return make_pair(deadRemoved, found);
}
// returns points in orthogonal range in O(n^{-((d-1)/d)} + k)
// where k is the number of points returned
VVT range_query(VT &range) {
    VVT result:
    int dummy = 0:
    VT region = infRegion(pt->size());
    range_query(range, region, result, dummy, false);
    return result:
}
// counts number of queries in range, runs in O(n^{(d-1)/d})
int count_query(VT &range) {
    int numPts = 0;
    VT region = infRegion(pt->size());
    range_query(range, region, dummy, numPts, true);
    return numPts;
}
void range_query(VT &range, VT &region, VVT &result, int &numPts, bool
 \hookrightarrow \quad \texttt{count)} \ \{
    if (region_contained(region, range)) {
        if (count)
            //by only adding size, we get rid of parameter
            //k in output sensitive O(n^{(d-1)/d} + k) analysis
            numPts += nAlive:
            addPtToResult(result);
        return;
    else if (isAlive && pt_contained(pt, range)) {
        if (count)
            ++numPts:
            result.push_back(pt);
    // are parts of the range to the right of splitting line?
    if ((*pt)[d] <= range[d*2+1] && right != NULL) {
        VT newRegion = region_intersect(region, (*pt)[d], d, false);
        right->range_query(range, newRegion, result, numPts, count);
    // are parts of the range to the left of splitting line?
    if ((*pt)[d] >= range[d*2] && left != NULL) {
        VT newRegion = region_intersect(region, (*pt)[d], d, true);
        left->range_query(range, newRegion, result, numPts, count);
}
// adds point to vector result and recursively calls addPt on children
void addPtToResult(VVT &result) {
```

```
if (isAlive)
            result.push_back(pt);
       if (left != NULL)
            left->addPtToResult(result);
        if (right != NULL)
            right->addPtToResult(result);
    // overloaded for first call with no current best
    VT * NN(VT &qpt) {
        VT *result = NN(qpt, NULL);
        return result;
    // performs NN query
    VT * NN(VT &qpt, VT *curBest) {
        bool goLeft = qpt[d] <= (*pt)[d];</pre>
        kdnode *child = goLeft ? left : right;
        if (isAlive)
            curBest = minPt(qpt, pt, curBest);
       if (child != NULL)
            curBest = child->NN(qpt, curBest);
        double curDist = curBest == NULL ? INF : dist(*curBest, qpt);
        // need to check other subtree
        if (curDist + EP > abs((*pt)[d] - qpt[d])) {
            kdnode *oppChild = goLeft ? right : left;
            if (oppChild != NULL) {
                curBest = oppChild->NN(qpt, curBest);
        return curBest:
   }
    // prints tree somewhat nicely
    void print_tree() {
       printf("((%c%d", isAlive ? 'A' : 'D', (*pt)[0]);
       for (size_t i = 1; i < pt->size(); ++i) {
            printf(" %d", (*pt)[i]);
        cout << endl;</pre>
       if (left != NULL) left->print_tree();
        printf(", ");
        if (right != NULL) right->print_tree();
        printf(")");
    ~kdnode() {
        delete left;
        delete right:
};
```

KDtree.cc KDtree.cc