UC San Diego

University of California San Diego

MnM

Mohit Gurumukhani, Nalin Bhardwaj, Maxwell Morehead

Contest (1)

template.cpp

```
#include <bits/stdc++.h>
using namespace std:
#define rep(i, a, b) for(int i = a; i < (b); ++i)</pre>
#define trav(a, x) for(auto& a : x)
#define all(x) begin(x), end(x)
#define sz(x) (int)(x).size()
typedef long long ll;
typedef pair <int, int> pii;
typedef vector <int> vi;
int main() {
  cin.sync with stdio(0); cin.tie(0);
  cin.exceptions (cin.failbit);
```

.bashrc

function mccne() { V="\$1"; shift; g++ -std=gnu++17 "\$@" -o "← \$V" "\$V.cpp": } function mcc() { mccne "\$@" -Wall -Werror -Wextra; } setxkbmap -option caps:escape # reset keyboard mappings #setxkbmap -layout us -option

.vimrc

```
set nu noeb sm sc ts=4 sts=4 sw=4 tm=250 ru bs=indent.eol.
sy enable | ino jk <esc> | vn fd <Esc>
filet indent plugin on
colo desert
" Select region and then type : Hash to hash your selection.
" Useful for verifying that there aren't mistypes.
ca Hash w !cpp -dD -P -fpreprocessed \| tr -d '[:space:]' \
\| md5sum \| cut -c-6
```

hash.sh

Hashes a file, ignoring all whitespace and comments. Use \leftarrow # verifying that code was correctly typed. cpp -dD -P -fpreprocessed | tr -d '[:space:]' | md5sum | cut -←

Mathematics (2)

2.1 Equations

$$ax + by = e$$

$$cx + dy = f$$

$$\Rightarrow x = \frac{ed - bf}{ad - bc}$$

$$y = \frac{af - ec}{ad - bc}$$

In general, given an equation Ax = b, the solution to a variable x_i is given by

$$x_i = \frac{\det A_i'}{\det A}$$

where A_i' is A with the i'th column replaced by b.

2.2 Recurrences

If $a_n = c_1 a_{n-1} + \cdots + c_k a_{n-k}$, and r_1, \ldots, r_k are distinct roots of $x^k + c_1 x^{k-1} + \cdots + c_k$, there are d_1, \ldots, d_k s.t.

$$a_n = d_1 r_1^n + \dots + d_k r_k^n.$$

Non-distinct roots r become polynomial factors, e.g. $a_n = (d_1 n + d_2)r^n.$

Trigonometry

$$\sin(v+w) = \sin v \cos w + \cos v \sin w$$
$$\cos(v+w) = \cos v \cos w - \sin v \sin w$$

$$\tan(v+w) = \frac{\tan v + \tan w}{1 - \tan v \tan w}$$
$$\sin v + \sin w = 2\sin\frac{v+w}{2}\cos\frac{v-w}{2}$$
$$\cos v + \cos w = 2\cos\frac{v+w}{2}\cos\frac{v-w}{2}$$

$$(V+W)\tan(v-w)/2 = (V-W)\tan(v+w)/2$$

where V, W are lengths of sides opposite angles v, w.

$$a\cos x + b\sin x = r\cos(x - \phi)$$
$$a\sin x + b\cos x = r\sin(x + \phi)$$

where $r = \sqrt{a^2 + b^2}$, $\phi = \operatorname{atan2}(b, a)$.

Geometry - 1

2.4.1 Triangles

Side lengths: a, b, c

Semiperimeter: $p = \frac{a+b+c}{2}$

Area: $A = \sqrt{p(p-a)(p-b)(p-c)}$

Circumradius: $R = \frac{abc}{4A}$

Inradius: $r = \frac{A}{}$

Length of median (divides triangle into two equal-area

triangles): $m_a = \frac{1}{2}\sqrt{2b^2 + 2c^2 - a^2}$

Length of bisector (divides angles in two):

$$s_a = \sqrt{bc \left[1 - \left(\frac{a}{b+c} \right)^2 \right]}$$

Law of sines:
$$\frac{\sin \alpha}{a} = \frac{\sin \beta}{b} = \frac{\sin \gamma}{c} = \frac{1}{2R}$$

Law of cosines: $a^2 = b^2 + c^2 - 2bc \cos \alpha$

Law of tangents:
$$\frac{a+b}{a-b} = \frac{\tan \frac{\alpha+\beta}{2}}{\tan \frac{\alpha-\beta}{2}}$$

2.4.2 Quadrilaterals

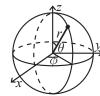
With side lengths a, b, c, d, diagonals e, f, diagonals angle θ , area A and magic flux $F = b^2 + d^2 - a^2 - c^2$:

$$4A = 2ef \cdot \sin \theta = F \tan \theta = \sqrt{4e^2f^2 - F^2}$$

For cyclic quadrilaterals the sum of opposite angles is 180°, ef = ac + bd, and $A = \sqrt{(p-a)(p-b)(p-c)(p-d)}$.

2.5Geometry - 2

2.5.1 Spherical coordinates



$$x = r \sin \theta \cos \phi \qquad r = \sqrt{x^2 + y^2 + z^2}$$

$$y = r \sin \theta \sin \phi \qquad \theta = a\cos(z/\sqrt{x^2 + y^2 + z^2})$$

$$z = r \cos \theta \qquad \phi = a\tan(2(y, x))$$

2.6Sums

$$c^{a} + c^{a+1} + \dots + c^{b} = \frac{c^{b+1} - c^{a}}{c-1}, c \neq 1$$

$$1 + 2 + 3 + \dots + n = \frac{n(n+1)}{2}$$

$$1^{2} + 2^{2} + 3^{2} + \dots + n^{2} = \frac{n(2n+1)(n+1)}{6}$$

$$1^{3} + 2^{3} + 3^{3} + \dots + n^{3} = \frac{n^{2}(n+1)^{2}}{4}$$

$$1^{4} + 2^{4} + 3^{4} + \dots + n^{4} = \frac{n(n+1)(2n+1)(3n^{2} + 3n - 1)}{30}$$

2.6.1 Discrete distributions

Binomial distribution

The number of successes in n independent yes/no experiments, each which yields success with probability p is $Bin(n, p), n = 1, 2, ..., 0 \le p \le 1.$

$$p(k) = \binom{n}{k} p^k (1-p)^{n-k}$$

$$\mu = np, \ \sigma^2 = np(1-p)$$

Bin(n, p) is approximately Po(np) for small p.

First success distribution

The number of trials needed to get the first success in independent yes/no experiments, each wich yields success with probability p is Fs(p), $0 \le p \le 1$.

$$p(k) = p(1-p)^{k-1}, k = 1, 2, \dots$$

$$\mu = \frac{1}{p}, \, \sigma^2 = \frac{1-p}{p^2}$$

Poisson distribution

The number of events occurring in a fixed period of time t if these events occur with a known average rate κ and independently of the time since the last event is $Po(\lambda), \lambda = t\kappa.$

$$p(k) = e^{-\lambda} \frac{\lambda^k}{k!}, k = 0, 1, 2, \dots$$
$$\mu = \lambda, \sigma^2 = \lambda$$

Continuous distributions

Uniform distribution

If the probability density function is constant between a and b and 0 elsewhere it is U(a, b), a < b.

$$f(x) = \begin{cases} \frac{1}{b-a} & a < x < b \\ 0 & \text{otherwise} \end{cases}$$

$$\mu = \frac{a+b}{2}, \, \sigma^2 = \frac{(b-a)^2}{12}$$

Exponential distribution

The time between events in a Poisson process is $\operatorname{Exp}(\lambda), \lambda > 0.$

$$f(x) = \begin{cases} \lambda e^{-\lambda x} & x \ge 0\\ 0 & x < 0 \end{cases}$$
$$\mu = \frac{1}{\lambda}, \, \sigma^2 = \frac{1}{\lambda^2}$$

Normal distribution

Most real random values with mean μ and variance σ^2 are well described by $\mathcal{N}(\mu, \sigma^2)$, $\sigma > 0$.

$$f(x) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(x-\mu)^2}{2\sigma^2}}$$

If $X_1 \sim \mathcal{N}(\mu_1, \sigma_1^2)$ and $X_2 \sim \mathcal{N}(\mu_2, \sigma_2^2)$ then $aX_1 + bX_2 + c \sim \mathcal{N}(\mu_1 + \mu_2 + c, a^2\sigma_1^2 + b^2\sigma_2^2)$

2.7Markov chains

A Markov chain is a discrete random process with the property that the next state depends only on the current state. Let X_1, X_2, \ldots be a sequence of random variables generated by the Markov process. Then there is a transition matrix $\mathbf{P} = (p_{ij})$, with $p_{ij} = \Pr(X_n = i | X_{n-1} = j)$, and $\mathbf{p}^{(n)} = \mathbf{P}^n \mathbf{p}^{(0)}$ is the probability distribution for X_n (i.e., $p_i^{(n)} = \Pr(X_n = i)$, where $\mathbf{p}^{(0)}$ is the initial distribution.

 π is a stationary distribution if $\pi = \pi \mathbf{P}$. If the Markov chain is *irreducible* (it is possible to get to any state from any state), then $\pi_i = \frac{1}{\mathbb{E}(T_i)}$ where $\mathbb{E}(T_i)$ is the expected time between two visits in state i. π_i/π_i is the expected number of visits in state i between two visits in state i.

For a connected, undirected and non-bipartite graph, where the transition probability is uniform among all neighbors, π_i is proportional to node i's degree.

A Markov chain is *ergodic* if the asymptotic distribution is independent of the initial distribution. A finite Markov chain is ergodic iff it is irreducible and aperiodic (i.e., the gcd of cycle lengths is 1). $\lim_{k\to\infty} \mathbf{P}^k = \mathbf{1}\pi$.

A Markov chain is an A-chain if the states can be partitioned into two sets A and G, such that all states in A are absorbing $(p_{ii} = 1)$, and all states in **G** leads to an absorbing state in **A**. The probability for absorption in state $i \in \mathbf{A}$, when the initial state is j, is $a_{ij} = p_{ij} + \sum_{k \in \mathbf{G}} a_{ik} p_{kj}$. The expected time until absorption, when the initial state is i, is $t_i = 1 + \sum_{k \in \mathbf{G}} p_{ki} t_k$.

2.7.1 Pick's theorem

B = number of lattice points on the boundary of the

I = number of lattice points in the interior of thepolygon.

 $Area = \frac{B}{2} + I - 1$

Data structures (3)

OrderStatisticTree.h

Description: A set (not multiset!) with support for finding the n'th element, and finding the index of an element. To get a map, change null_type. Time: $\mathcal{O}(\log N)$

```
#include <bits/extc++.h>
using namespace gnu pbds;
template<class T>
using Tree = tree <T, null type, less <T>, rb tree tag,
    tree order statistics node update>;
void example() {
  Tree<int> t, t2; t.insert(8);
  auto it = t.insert(10).first;
  assert(it == t.lower bound(9));
  assert(t.order of key(10) == 1);
  assert(t.order_of_key(11) == 2);
assert(*t.find_by_order(0) == 8);
  t.join(t2); // assuming T < T2 or T > T2, merge t2 into t
```

HashMap.h

Description: Hash map with the same API as unordered map, but ~3x faster. Initial capacity must be a power of 2 (if provided). 1443bc, 7 lines

```
#include <bits/extc++.h>
// To use most bits rather than just the lowest ones:
struct chash {
  const uint64 t C = ll(2e18 * M PI) + 71; // large odd ←
  ll operator()(ll x) const { return __builtin_bswap64(x*C); ←
};
_gnu_pbds::gp_hash_table<ll,int,chash> h←
     (\{\},\{\},\{\},\{\},\{1<<16\});
```

Matrix.h

Description: Basic operations on square matrices. Usage: Matrix<int, 3> A; A.d = $\{\{\{1,2,3\}\}, \{\{4,5,6\}\}, \{\{7,8,9\}\}\}\}\};$ vector<int> vec = $\{1,2,3\}$; $vec = (A^N) * vec;$ c43c7d, 26 lines template<class T, int N> struct Matrix { typedef Matrix M; array <array <T, N>, N> d{};

```
M operator*(const M& m) const {
  rep(i,0,N) rep(j,0,N)
    rep(k,0,N) a.d[i][j] += d[i][k]*m.d[k][j];
  return a;
vector <T> operator*(const vector <T>& vec) const {
  vector <T> ret(N);
  rep(i,0,N) rep(j,0,N) ret[i] += d[i][j] * vec[j];
```

```
return ret;
  M operator^(ll p) const {
     assert (p \geq 0);
    M a, b(*this);
     rep(i,0,N) a.d[i][i] = 1;
     while (p) {
       if (p&1) a = a*b;
      b = b*b;
      p >>= 1;
     return a;
};
Treap.h
Description: Treap with lazy propogation and climbing for reverse lookup.
Time: Split and merge are \mathcal{O}(\log N)
struct pnode {
  int sz, prior;
  int v, lz, vval = 0;
  pnode *L, *R, *P;
  pnode(pnode* l = NULL, pnode* r = NULL, int val = 0) {
    L = l, R = r, P = NULL, v = val, sz = 1, lz = 0, prior =\leftarrow
};
typedef pnode* node;
int get sz(node t) { return t?t->sz:0; }
int get v(node t) { return t?t->v:0; }
int get lz(node t) { return t?t->lz:0; }
void upd P(node& t) {
  if(t) {
     if(t\rightarrow L) t\rightarrow L\rightarrow P = t;
     if(t\rightarrow R) t\rightarrow R\rightarrow P = t;
void upd sz(node& t) {
  if(t) t\rightarrow sz = get sz(t\rightarrow L)+get sz(t\rightarrow R)+1;
void upd lz(node& t) {
  if(t && t->lz) {
    t\rightarrow vval += t\rightarrow lz;
    t \rightarrow v += get sz(t) + t \rightarrow lz;
    if(t\rightarrow L) t\rightarrow L\rightarrow lz += t\rightarrow lz;
    if(t->R) t->R->lz += t->lz;
     t\rightarrow lz = 0;
void upd v(node& t) {
  if(t) {
     upd lz(t->L); upd lz(t->R);
    t \rightarrow v = get v(t \rightarrow L) + get v(t \rightarrow R) + t \rightarrow vval;
void split(node t, node& l, node& r, int key, int add) {
  upd lz(t); upd lz(l); upd lz(r);
  upd P(t); upd P(1); upd P(r);
  if(!t) {
    l = r = NULL;
```

```
return;
  int cur key = add+get sz(t->L);
 if(cur_key < key) {</pre>
    split(t->R, t->R, r, key, cur key+1);
  else {
    split(t->L, l, t->L, key, add);
    r = t;
  upd sz(t); upd v(t);
  upd sz(1); upd v(1);
  upd sz(r); upd v(r);
  upd_P(t); upd_P(l); upd_P(r);
void merge(node& t, node l, node r) {
  upd lz(1); upd lz(r); upd lz(t);
  upd P(1); upd P(r); upd P(t);
  if(!| || !r) {
    t = 1?1:r:
    return;
 if(l->prior > r->prior) {
    merge(1->R, 1->R, r);
    t = 1;
  else {
    merge(r\rightarrow L, l, r\rightarrow L);
    t = r;
 upd sz(1); upd v(1);
  upd_sz(r); upd_v(r);
  upd sz(t); upd v(t);
 upd P(1); upd P(r); upd P(t);
int climber (node t, bool add) {
  int res = (add?get sz(t\rightarrow L)+1:0);
 if(t->P) {
    if(t->P->R) {
      if(t\rightarrow P\rightarrow R\rightarrow prior == t\rightarrow prior) res += climber(t\rightarrow P,1);
      else res += climber(t \rightarrow P, 0);
    else res += climber(t \rightarrow P, 0);
  return res;
void printer(node t, lli add = 0) {
 if(t) {
     printer(t->L, add);
    cerr << t->v << " " << t->tr << " " << t->vval << " " << \leftarrow
           t \rightarrow prior \ll " " \ll add+get sz(t \rightarrow L) \ll "\n";
    printer(t->R, add+get sz(t->L)+1);
void fix(node& t) {
    if(t\rightarrow P) t\rightarrow P = NULL;
// Initialise
srand (time (NULL));
```

```
// Call after each qry/upd, root is overall root of running \
fix (root);
FenwickTree2d.h
Description: Computes sums a[i,j] for all i<I, j<J, and increases single ele-
ments a[i,j]. Requires that the elements to be updated are known in advance
(call fakeUpdate() before init()).
Time: \mathcal{O}(\log^2 N). (Use persistent segment trees for \mathcal{O}(\log N).)
"FenwickTree.h"
                                                         b28c27, 22 lines
struct FT2 {
  vector <vi>vs; vector <FT> ft;
  FT2(int limx): ys(limx) {}
  void fakeUpdate(int x, int y) {
    for (; x < sz(ys); x = x + 1) ys[x].push back(y);
  void init() {
    trav(v, ys) sort(all(v)), ft.emplace back(sz(v));
  int ind(int x, int y) {
    return (int)(lower bound(all(ys[x]), y) - ys[x].begin())←
  void update(int x, int y, ll dif) {
    for (; x < sz(ys); x = x + 1)
      ft[x].update(ind(x, y), dif);
  11 query(int x, int y) {
    11 sum = 0;
    for (; x; x &= x - 1)
      sum += ft[x-1].query(ind(x-1, y));
    return sum;
};
CHT.h
Description: Convex hull trick for finding max f(x) given a number of lines
f(x) = mx + c. During inserting, m should be in increasing order.
Time: Query is \mathcal{O}(\log N). Insert is \mathcal{O}(1) amortized
                                                         efe2ae, 36 lines
typedef long long int lli;
const lli maxn = lli(1e5)+5;
struct line {
  lli m, c;
  line(lli m = 0, lli c = 0)
   m = _m, c = _c;
};
struct cht {
  lli sz = 0;
  line st[maxn];
  double intersect(line a, line b) { return double(a.c-b.c) ←
       /(b.m-a.m); }
  void insert(line a) {
    while(sz > 1) {
      if(intersect(st[sz-2], a) < intersect(st[sz-2], st[sz \leftarrow
            -1])) sz--;
       else break;
    st[sz++] = a;
  lli qry(lli x) {
    lli L = 0, R = sz-1;
```

while(L < R) {

lli mid = (L+R)/2;

if(x < intersect(st[mid], st[mid+1])) R = mid;</pre>

```
else L = mid+1;
    return st[L].m*x+st[L].c;
};
LiChaoTree.h
Description: Add line y = mx + c using add line(\{m, c\}). Query for point
x using query(x). For max instead of min, change lines marked with ###.
P > largest point at which query occurs Coordinate compress if too large
point values.
Time: \mathcal{O}(\log N)
                                                       93d5aa, 57 lines
typedef long long ll;
const ll maxn = ll(1e5)+5, inf = ll(1e18)+5;
struct point {
  11 m, c; // y = mx + c
  point(): m(0), c(inf) {} // ### (inf \rightarrow -inf)
  point(ll m, ll c): m(m), c(c) {}
point line[8 * maxn];
struct li chao tree {
    inline int left(int node) { return (node<<1); }</pre>
    inline int right(int node) { return (node<<1)+1; }</pre>
  void add line(point add, int node = 1, ll l = -maxn, ll r ←
       = \max n  {
    11 m = (1 + r) / 2;
    bool lef = (eval(add, 1) < eval(line[node], 1)); // ###
    bool mid = (eval(add, m) < eval(line[node], m)); // ###
    if (mid) {
      swap(add, line[node]);
    if (1 == r - 1) { // Leaf Node
      return;
    } else if (lef != mid) { // Intersection point in [1, m)
      add line(add, left (node), l, m);
    } else { // Intersection point in [m, r)
      add line(add, right(node), m, r);
  ll query(ll x, int node = 1, ll l = -\max, ll r = \maxn) {
    ll ans = eval(line[node], x), m = (l + r) / 2;
    if (l == r - 1) {
      return ans;
    } else if (x < m) {</pre>
      return min(ans, query(x, left(node), l, m)); // ###
      return min(ans, query(x, right(node), m, r)); // ###
  ll eval(point p, ll x) {
    return p.m * x + p.c;
  void clear(int node = 1, ll l = -maxn, ll r = maxn) {
    11 \text{ m} = (1 + r) / 2;
    if (line[node].c == inf) {
      return;
    line[node] = point();
    if (l == r - 1) {
      return;
```

```
clear (left (node), l, m);
    clear(right(node), m, r);
};
PersistentSegtree.h
Description: Persistent segment tree, implemented using pointers.
Time: Query and Update are \mathcal{O}(\log N)
                                                        a2870b, 48 lines
struct pnode {
  int res = 0:
  pnode *L, *R;
  pnode(int v = 0, pnode* l = NULL, pnode* r = NULL) {
    L = 1;
    R = r;
};
typedef pnode* node;
node dummy;
int A[int(1e5)+5], pm[int(1e5)+5];
node root[int(1e5)+5];
node upd(node root, int L, int R, int idx) {
  if(L == R \&\& L == idx)  {
    return new prode(root->res+1, dummy, dummy);
  }
  else {
    if(idx >= L && idx <= (L+R)/2) return new pnode(root->←
         res+1, upd(root->L, L, (L+R)/2, idx), root->R);
    else return new pnode(root->res+1, root->L, upd(root->R, ←
          (L+R)/2+1, R, idx));
 }
int qry(node rootL, node rootR, int L, int R, int k) {
 if(L == R) return L;
  else {
    int left = rootR->L->res-rootL->L->res;
    //cout << L << " " << R << " " << left << " " << k << " \leftarrow
         n";
    if(left >= k) {
      return gry(rootL->L, rootR->L, L, (L+R)/2, k);
    else return gry(rootL->R, rootR->R, (L+R)/2+1, R, k-left \( - \)
// Initialise
dummy = new pnode();
dummy \rightarrow L = dummy \rightarrow R = dummy;
for(int i = 0; i < n; i++) {
 if(i) root[i] = upd(root[i-1], 0, v, A[i]);
  else root[i] = upd(dummy, 0, v, A[i]);
LCA.h
Description: LCA table with offline update support.
Time: Updates are \mathcal{O}(N \log N) total.
const int max = int(2e5)+5, maxlog = 20, inf = int(1e9)+5;
```

```
int n, m, T[maxn][maxlog+1], TT[maxn][maxlog+1], F[maxn][←
     \max(g+1), H[\max n], upar[\max n], P[\max n], taken[\max n], \leftarrow
     res[maxn];
pair <pair <int, int>, pair <int, int>> E[maxn];
vector <pair <int, int>> graph [maxn];
void dfs0(int node, int par, int income, int dep) {
  upar[node] = income;
  T[node][0] = par;
  H[node] = dep;
  if(income != -1) TT[node][0] = E[income].first.first;
  for(auto it: graph[node]) {
    if(it.first != par) dfs0(it.first, node, it.second, dep←
}
void init()
  for(int j = 1; j <= maxlog; j++) {</pre>
    for(int i = 0; i < n; i++) {
      if(T[i][j-1] != -1) {
        T[i][j] = T[T[i][j-1]][j-1];
        TT[i][j] = max(TT[i][j-1], TT[T[i][j-1]][j-1]);
void onit()
  for(int j = maxlog; j > 0; j--) {
    for(int i = 0; i < n; i++) {
      F[i][j-1] = min(F[i][j-1], F[i][j]);
      if(T[i][j-1] != -1) {
        int node = T[i][j-1];
        F[node][j-1] = min(F[node][j-1], F[i][j]);
int qry(int x, int y)
  if(H[x] > H[y]) swap(x, y);
  int res = -inf;
  for(int i = maxlog; i >= 0; i--) {
    if(H[y]-(1<<i)) >= H[x]) {
      res = max(res, TT[y][i]);
      y = T[y][i];
  if(x == y) return res;
  for(int i = maxlog; i \ge 0; i--) {
    if(T[x][i] != T[y][i]) {
      res = max(res, TT[x][i]);
      res = max(res, TT[y][i]);
      x = T[x][i], y = T[y][i];
  res = max(res, TT[x][0]);
  res = max(res, TT[y][0]);
  return res;
void upd(int x, int y, int c)
```

```
if(H[x] > H[y]) swap(x, y);
  for(int i = maxlog; i >= 0; i--) {
    if(H[y]-(1<<i)) >= H[x]) {
     F[y][i] = min(F[y][i], c);
      y = T[y][i];
  if(x == y) return;
  for(int i = maxlog; i >= 0; i--) {
    if(T[x][i] != T[y][i]) {
     F[x][i] = min(F[x][i], c), F[y][i] = min(F[y][i], c);
      x = T[x][i], y = T[y][i];
  F[x][0] = min(F[x][0], c), F[y][0] = min(F[y][0], c);
// Initialise before input
for(int i = 0; i < n; i++) P[i] = i;
for(int i = 0; i < n; i++) {</pre>
  for(int j = 0; j \leftarrow \max(j + 1) T[i][j] = -1, TT[i][j] = F[\leftarrow
       i | [i] = \inf;
// Initialise after input
dfs0(0, -1, -1, 0);
init();
// Update structure
for(all u, v, c) upd(u, v, c);
onit(); // After all upd()s.
HLDSubtree.h
Description: HLD implementation that also supports subtree updates/-
Time: Path query is \mathcal{O}(\log^2 n)
                                                      890b14, 137 lines
typedef long long int lli;
const lli maxn = lli(1e5)+5, maxlog = 17, inf = lli(1e17)+5;
lli n, totchain = 0, curst = 0, inchain[maxn], head[maxn],
     inst [maxn], H[maxn], T[maxn] [maxlog+1], lz [4*maxn], ←
     start[maxn], en[maxn], sz[maxn], lookup[maxn], ptr[maxn
    ];
vector < lli > graph [maxn];
vector < lli > girls [maxn];
pair < lli, lli > st [4*maxn];
inline lli left(lli node) { return (node<<1); }</pre>
inline lli right(lli node) { return (node<<1)+1; }</pre>
void build (lli node, lli L, lli R) {
  if(L == R) {
    lli realnode = lookup[L];
    if(girls[realnode].empty()) st[node] = {inf, realnode};
    else st[node] = {*girls[realnode].begin(), realnode};
  else {
    build (left (node), L, (L+R)/2);
    build (right (node), (L+R)/2+1, R);
    st[node] = min(st[left(node)], st[right(node)]);
void shift (lli node, lli L, lli R) {
```

```
if(lz[node] && L != R) {
    lz[left(node)] += lz[node];
    lz[right(node)] += lz[node];
    st[left(node)].first += lz[node];
    st[right(node)].first += lz[node];
  lz[node] = 0;
void upd(lli node, lli L, lli R, lli a, lli b, lli v) {
 if(a > R \mid | b < L) return;
  else if(a <= L && R <= b) {
    st[node].first += v;
    lz[node] += v;
  else {
    shift (node, L, R);
    upd(left (node), L, (L+R)/2, a, b, v);
    upd(right(node), (L+R)/2+1, R, a, b, v);
    st[node] = min(st[left(node)], st[right(node)]);
pair < lli, lli > qry(lli node, lli L, lli R, lli a, lli b) {
  if(a > R | | b < L) return {inf, -1};
  else if(a <= L && R <= b) return st[node];
    shift (node, L, R);
    return min(qry(left(node), L, (L+R)/2, a, b), qry(right(-
         node), (L+R)/2+1, R, a, b));
}
void init() {
  for(lli j = 1; j <= maxlog; j++) {
    for(lli i = 0; i < n; i++) {
      if(T[i][j-1] != -1) T[i][j] = T[T[i][j-1]][j-1];
 }
lli LCA(lli x, lli y) {
  if(H[x] > H[y]) swap(x, y);
  for(lli i = maxlog; i >= 0; i--) {
    if(H[y]-(1<<i)) >= H[x]) y = T[y][i];
  if(x == y) return x;
  for(lli i = maxlog; i >= 0; i--) {
    if(T[x][i] != T[y][i]) x = T[x][i], y = T[y][i];
  return T[x][0];
// v MUST be an ancestor of u
pair < lli, lli > pathqry(lli node, lli anc)
  lli cur = node;
  pair < lli, lli > res = \{\inf, -1\};
  while(inchain[cur] != inchain[anc])
    res = min(res, qry(1, 0, curst-1, inst[head[inchain[cur←
        ]]], inst[cur]));
    cur = T[head[inchain[cur]]][0];
  res = min(res, qry(1, 0, curst-1, inst[anc], inst[cur]));
  return res;
void dfs0(lli node, lli par, lli ht) {
```

```
sz[node] = 1; H[node] = ht; T[node][0] = par;
  for(auto it: graph[node]) {
    if(it != par) {
      dfs0(it, node, ht+1);
      sz[node] += sz[it];
 }
void dfs1(lli node, lli par, lli chain) {
 inchain[node] = chain;
  if(head[chain] == -1) head[chain] = node;
  inst[node] = curst++;
  lookup[curst-1] = node;
  start[node] = curst-1;
  pair<lli, lli > largest = \{-1, -1\};
  for(auto it: graph[node]) {
    if(it != par) largest = max(largest, {sz[it], it});
  if(largest.second != -1) dfs1(largest.second, node, chain)←
  for(auto it: graph[node]) {
    if(it != par && it != largest.second) {
      dfs1(it, node, totchain++);
 en[node] = curst-1;
// Add k to subtree(node)
upd(1, 0, n-1, start[node], en[node], k);
// initialisation before input
for(lli i = 0; i < maxn; i++) head[i] = -1;
for(lli i = 0; i < maxn; i++) {</pre>
 for(lli j = 0; j <= \max(j; j++) T[i][j] = -1;
// initialisation after input
dfs0(0, -1, 0);
init();
totchain = 1;
dfs1(0, -1, 0);
build(1, 0, curst-1);
```

Numerical (4)

4.1 Polynomials and recurrences

```
};
PolvRoots.h
Description: Finds the real roots to a polynomial.
Usage: poly_roots(\{\{2,-3,1\}\},-1e9,1e9) // solve x^2-3x+2=0
Time: \mathcal{O}\left(n^2\log(1/\epsilon)\right)
"Polynomial.h"
                                                          2cf190, 23 lines
vector <double > poly roots (Poly p, double xmin, double xmax) \leftarrow
  if (sz(p.a) == 2) \{ return \{-p.a[0]/p.a[1]\}; \}
  vector <double> ret;
  Poly der = p;
  der.diff();
  auto dr = poly roots(der, xmin, xmax);
  dr.push back(xmin-1);
  dr.push back(xmax+1);
  sort (all (dr));
  rep(i,0,sz(dr)-1) {
    double l = dr[i], h = dr[i+1];
    bool sign = p(1) > 0;
    if (sign ^ (p(h) > 0)) {
      rep(it,0,60) { // while (h - 1 > 1e-8)
        double m = (l + h) / 2, f = p(m);
        if ((f <= 0) ^ sign) l = m;
        else h = m;
      ret.push back((l + h) / 2);
  return ret;
PolyInterpolate.h
Description: Given n points (x[i], y[i]), computes an n-1-degree polynomial
```

Description: Given n points (x[i], y[i]), computes an n-1-degree polynomial p that passes through them: $p(x) = a[0] * x^0 + ... + a[n-1] * x^{n-1}$. For numerical precision, pick $x[k] = c * \cos(k/(n-1) * \pi), k = 0 \dots n-1$. **Time:** $\mathcal{O}(n^2)$

```
typedef vector<double> vd;
vd interpolate(vd x, vd y, int n) {
  vd res(n), temp(n);
```

```
vd interpolate(vd x, vd y, int n) {
  vd res(n), temp(n);
  rep(k,0,n-1) rep(i,k+1,n)
    y[i] = (y[i] - y[k]) / (x[i] - x[k]);
  double last = 0; temp[0] = 1;
  rep(k,0,n) rep(i,0,n) {
    res[i] += y[k] * temp[i];
    swap(last, temp[i]);
    temp[i] -= last * x[k];
  }
  return res;
}
```

BerlekampMassey.h

ll $d = s[i] \% \mod;$

Description: Recovers any n-order linear recurrence relation from the first 2n terms of the recurrence. Useful for guessing linear recurrences after brute-forcing the first terms. Should work on any field, but numerical stability for floats is not guaranteed. Output will have size $\leq n$.

```
Usage: BerlekampMassey(\{0, 1, 1, 3, 5, 11\}) // \{1, 2\}

Time: \mathcal{O}(N^2)

".../number-theory/ModPow.h"

vector<\{1\} BerlekampMassey(vector<\{1\} s) \{

int n = sz(s), L = 0, m = 0;

vector<\{1\} \mathcal{O}(n), \mathcal{O}(
```

```
rep(j,1,L+1) d = (d + C[j] * s[i - j]) % mod;
    if (!d) continue;
   T = C; 11 coef = d * modpow(b, mod-2) % mod;
    rep(j,m,n) C[j] = (C[j] - coef * B[j - m]) % mod;
    if (2 * L > i) continue;
    L = i + 1 - L; B = T; b = d; m = 0;
 C. resize (L + 1); C. erase (C. begin ());
 trav(x, C) x = (mod - x) \% mod;
 return C:
LinearRecurrence.h
Description: Generates the k'th term of an n-order linear recurrence
S[i] = \sum_{j} S[i-j-1]tr[j], given S[0... \ge n-1] and tr[0...n-1]. Faster
than matrix multiplication. Useful together with Berlekamp-Massey.
Usage: linearRec(\{0, 1\}, \{1, 1\}, k) // k'th Fibonacci number
Time: \mathcal{O}\left(n^2 \log k\right)
                                                       f4e444, 26 lines
typedef vector <11 > Poly;
ll linearRec(Poly S, Poly tr, ll k) {
 int n = sz(tr);
 auto combine = [&](Poly a, Poly b) {
    Poly res(n * 2 + 1);
    rep(i,0,n+1) rep(j,0,n+1)
     res[i + j] = (res[i + j] + a[i] * b[j]) % mod;
    for (int i = 2 * n; i > n; --i) rep(j,0,n)
     res[i - 1 - j] = (res[i - 1 - j] + res[i] * tr[j]) % \leftarrow
          mod:
    res.resize(n + 1):
    return res;
 };
 Poly pol(n + 1), e(pol);
 pol[0] = e[1] = 1;
  for (++k; k; k /= 2) {
    if (k % 2) pol = combine(pol, e);
    e = combine(e, e);
 ll res = 0;
 rep(i,0,n) res = (res + pol[i + 1] * S[i]) % mod;
 return res:
```

4.2 Optimization

GoldenSectionSearch.h

} else {

Description: Finds the argument minimizing the function f in the interval [a,b] assuming f is unimodal on the interval, i.e. has only one local minimum. The maximum error in the result is eps. Works equally well for maximization with a small change in the code. See TernarySearch.h in the Various chapter for a discrete version.

```
Usage: double func(double x) { return 4+x+.3*x*x; } double xmin = gss(-1000,1000,func);

Time: \mathcal{O}(\log((b-a)/\epsilon)) 31d45b, 14 lines

double gss(double a, double b, double (*f)(double)) {
    double r = (sqrt(5)-1)/2, eps = 1e-7;
    double x1 = b - r*(b-a), x2 = a + r*(b-a);
    double f1 = f(x1), f2 = f(x2);
    while (b-a > eps)
    if (f1 < f2) { //change to > to find maximum b = x2; x2 = x1; f2 = f1;
```

x1 = b - r*(b-a); f1 = f(x1);

```
a = x1; x1 = x2; f1 = f2;
      x2 = a + r*(b-a); f2 = f(x2);
  return a;
HillClimbing.h
Description: Poor man's optimization for unimodal functions f40e55, 16 lines
typedef array <double, 2> P;
double func (P p);
pair <double, P> hillClimb (P start) {
  pair <double, P> cur(func(start), start);
  for (double jmp = 1e9; jmp > 1e-20; jmp /= 2) {
    rep(j,0,100) rep(dx,-1,2) rep(dy,-1,2) {
      P p = cur.second;
      p[0] += dx*imp;
      p[1] += dy*jmp;
      cur = min(cur, make pair(func(p), p));
  return cur:
```

Integrate.h

Description: Simple integration of a function over an interval using Simpson's rule. The error should be proportional to h^4 , although in practice you will want to verify that the result is stable to desired precision when epsilon changes.

4756fc, 7 lines

```
template < class F>
double quad (double a, double b, F f, const int n = 1000) {
    double h = (b - a) / 2 / n, v = f(a) + f(b);
    rep(i,1,n*2)
    v += f(a + i*h) * (i&1 ? 4 : 2);
    return v * h / 3;
}
Integrate Adaptive.h
```

```
typedef double d;
#define S(a,b) (f(a) + 4*f((a+b) / 2) + f(b)) * (b-a) / 6

template <class F>
d rec(F& f, d a, d b, d eps, d S) {
    d c = (a + b) / 2;
    d S1 = S(a, c), S2 = S(c, b), T = S1 + S2;
    if (abs(T - S) <= 15 * eps || b - a < 1e-10)
        return T + (T - S) / 15;
    return rec(f, a, c, eps / 2, S1) + rec(f, c, b, eps / 2, \( \to \)
        S2);
}
template < class F>
d quad(d a, d b, F f, d eps = 1e-8) {
    return rec(f, a, b, eps, S(a, b));
}
```

Simplex.h

Description: Solves a general linear maximization problem: maximize c^Tx subject to $Ax \leq b$, $x \geq 0$. Returns -inf if there is no solution, inf if there are arbitrarily good solutions, or the maximum value of c^Tx otherwise. The input vector is set to an optimal x (or in the unbounded case, an arbitrary solution fulfilling the constraints). Numerical stability is not guaranteed. For better performance, define variables such that x = 0 is viable.

```
Usage: vvd A = \{\{1,-1\}, \{-1,1\}, \{-1,-2\}\};
vd b = \{1,1,-4\}, c = \{-1,-1\}, x;
T val = LPSolver(A, b, c).solve(x);
Time: \mathcal{O}(NM * \#pivots), where a pivot may be e.g. an edge relaxation.
\mathcal{O}(2^n) in the general case.
                                                     aa8530, 68 lines
typedef double T; // long double, Rational, double + mod√P←
typedef vector <T> vd;
typedef vector <vd> vvd;
const T eps = 1e-8, inf = 1/.0;
#define MP make pair
#define ltj(X) if(s == -1 || MP(X[j],N[j]) < MP(X[s],N[s])) ←
    s=i
struct LPSolver {
  int m, n;
  vi N, B;
  vvd D;
  LPSolver(const vvd& A, const vd& b, const vd& c):
   m(sz(b)), n(sz(c)), N(n+1), B(m), D(m+2, vd(n+2)) {
      rep(i,0,m) rep(j,0,n) D[i][j] = A[i][j];
      rep(i,0,m) { B[i] = n+i; D[i][n] = -1; D[i][n+1] = b[i \leftarrow
      rep(j,0,n) \{ N[j] = j; D[m][j] = -c[j]; \}
     N[n] = -1; D[m+1][n] = 1;
  void pivot(int r, int s) {
   T *a = D[r].data(), inv = 1 / a[s];
    rep(i,0,m+2) if (i != r && abs(D[i][s]) > eps) {
     T *b = D[i].data(), inv2 = b[s] * inv;
      rep(j,0,n+2) b[j] -= a[j] * inv2;
     b[s] = a[s] * inv2;
    rep(j,0,n+2) if (j != s) D[r][j] *= inv;
    rep(i,0,m+2) if (i != r) D[i][s] *= -inv;
   D[r][s] = inv;
    swap(B[r], N[s]);
  bool simplex(int phase) {
    int x = m + phase - 1;
    for (;;) {
      int s = -1;
      rep(j,0,n+1) if (N[j] != -phase) ltj(D[x]);
      if (D[x][s] >= -eps) return true;
      int r = -1;
      rep(i,0,m) {
        if (D[i][s] <= eps) continue;</pre>
       if (r == -1 || MP(D[i][n+1] / D[i][s], B[i])
                     if (r == -1) return false;
      pivot(r, s);
  T solve(vd &x) {
    int r = 0;
    rep(i,1,m) if (D[i][n+1] < D[r][n+1]) r = i;
    if (D[r][n+1] < -eps) {
      pivot(r, n);
      if (!simplex(2) || D[m+1][n+1] < -eps) return -inf;</pre>
      rep(i,0,m) if (B[i] == -1) {
        int s = 0:
        rep(j,1,n+1) ltj(D[i]);
```

```
pivot(i, s);
      }
    bool ok = simplex(1); x = vd(n);
    rep(i,0,m) if (B[i] < n) x[B[i]] = D[i][n+1];
    return ok ? D[m] [n+1] : inf;
};
       Matrices
Determinant.h
Description: Calculates determinant of a matrix. Destroys the matrix.
Time: \mathcal{O}(N^3)
double det(vector < vector < double >>& a) {
 int n = sz(a); double res = 1;
 rep(i,0,n) {
    int b = i;
    rep(j,i+1,n) if (fabs(a[j][i]) > fabs(a[b][i])) b = j;
    if (i != b) swap(a[i], a[b]), res *= -1;
    res *= a[i][i];
    if (res == 0) return 0;
    rep(j,i+1,n) {
      double v = a[j][i] / a[i][i];
      if (v != 0) rep(k, i+1, n) a[j][k] -= v * a[i][k];
 }
 return res;
IntDeterminant.h
Description: Calculates determinant using modular arithmetics. Modulos
can also be removed to get a pure-integer version.
Time: \mathcal{O}(N^3)
                                                        3313dc, 18 lines
const ll mod = 12345;
ll det(vector < vector < ll >> & a) {
 int n = sz(a); ll ans = 1;
 rep(i,0,n) {
    rep(j,i+1,n) {
      while (a[j][i] != 0) { // gcd step
        11 t = a[i][i] / a[j][i];
        if (t) rep(k,i,n)
          a[i][k] = (a[i][k] - a[j][k] * t) % mod;
        swap(a[i], a[j]);
        ans *= -1;
    ans = ans \star a[i][i] % mod;
    if (!ans) return 0;
 return (ans + mod) % mod;
Description: Solves A * x = b. If there are multiple solutions, an arbitrary
one is returned. Returns rank, or -1 if no solutions. Data in A and b is lost.
Time: \mathcal{O}\left(n^2m\right)
                                                        44c9ab, 38 lines
typedef vector <double> vd;
const double eps = 1e-12;
int solveLinear(vector <vd>& A, vd& b, vd& x) {
 int n = sz(A), m = sz(x), rank = 0, br, bc;
 if (n) assert(sz(A[0]) == m);
 vi col(m); iota(all(col), 0);
```

rep(i,0,n) {

```
if ((v = fabs(A[r][c])) > bv)
        br = r, bc = c, bv = v;
    if (bv <= eps) {
      rep(j,i,n) if (fabs(b[j]) > eps) return -1;
      break;
    swap(A[i], A[br]);
    swap(b[i], b[br]);
    swap(col[i], col[bc]);
    rep(j,0,n) swap(A[j][i], A[j][bc]);
    bv = 1/A[i][i];
    rep(j,i+1,n) {
      double fac = A[j][i] * bv;
      b[j] = fac * b[i];
      rep(k, i+1,m) A[j][k] -= fac*A[i][k];
    rank++;
  }
  x.assign (m, 0);
  for (int i = rank; i--;) {
    b[i] /= A[i][i];
    x[col[i]] = b[i];
    rep(j,0,i) b[j] -= A[j][i] * b[i];
  return rank; // (multiple solutions if rank < m)
SolveLinear2.h
Description: To get all uniquely determined values of x back from Solve-
Linear, make the following changes:
"SolveLinear.h"
                                                        08e495, 7 lines
rep(j,0,n) if (j != i) // instead of rep(j,i+1,n)
// ... then at the end:
x.assign (m, undefined);
rep(i,0,rank) {
  rep(j,rank,m) if (fabs(A[i][j]) > eps) goto fail;
  x[col[i]] = b[i] / A[i][i];
fail:; }
SolveLinearBinary.h
Description: Solves Ax = b over \mathbb{F}_2. If there are multiple solutions, one is
returned arbitrarily. Returns rank, or -1 if no solutions. Destroys A and b.
Time: \mathcal{O}\left(n^2m\right)
                                                       fa2d7a, 34 lines
typedef bitset <1000> bs;
int solveLinear(vector <bs>& A, vi& b, bs& x, int m) {
  int n = sz(A), rank = 0, br;
  assert (m \le sz(x));
  vi col(m); iota(all(col), 0);
  rep(i,0,n) {
    for (br=i; br<n; ++br) if (A[br].any()) break;
    if (br == n) {
      rep(j,i,n) if(b[j]) return -1;
      break;
    int bc = (int)A[br]. Find next(i-1);
    swap(A[i], A[br]);
    swap(b[i], b[br]);
    swap(col[i], col[bc]);
    rep(j,0,n) if (A[j][i] != A[j][bc]) {
      A[j].flip(i); A[j].flip(bc);
    rep(j,i+1,n) if (A[j][i]) {
      b[j] ^= b[i];
      A[j] \sim A[i];
```

double v, bv = 0;

rep(r,i,n) rep(c,i,m)

```
rank++;
x = bs();
for (int i = rank; i--;) {
 if (!b[i]) continue;
 x[col[i]] = 1;
 rep(j,0,i) b[j] ^= A[j][i];
return rank; // (multiple solutions if rank < m)
```

MatrixInverse.h

Description: Invert matrix A. Returns rank; result is stored in A unless singular (rank < n). Can easily be extended to prime moduli; for prime powers, repeatedly set $A^{-1} = A^{-1}(2I - AA^{-1}) \pmod{p^k}$ where A^{-1} starts as the inverse of A mod p, and k is doubled in each step.

```
Time: \mathcal{O}\left(n^3\right)
                                                       ebfff6, 35 lines
int matInv(vector < vector < double >> & A) {
  int n = sz(A); vi col(n);
  vector < vector < double >> tmp(n, vector < double > (n));
  rep(i,0,n) tmp[i][i] = 1, col[i] = i;
  rep(i,0,n) {
    int r = i, c = i;
    rep(j,i,n) rep(k,i,n)
      if (fabs(A[j][k]) > fabs(A[r][c]))
        r = j, c = k;
    if (fabs(A[r][c]) < 1e-12) return i;
   A[i].swap(A[r]); tmp[i].swap(tmp[r]);
    rep(j,0,n)
      swap(A[j][i], A[j][c]), swap(tmp[j][i], tmp[j][c]);
    swap(col[i], col[c]);
    double v = A[i][i];
    rep(j,i+1,n) {
     double f = A[j][i] / v;
     A[j][i] = 0;
     rep(k, i+1,n) A[j][k] = f*A[i][k];
     rep(k,0,n) tmp[j][k] = f*tmp[i][k];
   rep(j, i+1,n) A[i][j] /= v;
   rep(j,0,n) tmp[i][j] /= v;
   A[i][i] = 1;
  for (int i = n-1; i > 0; --i) rep(j,0,i) {
   double v = A[j][i];
   rep(k,0,n) tmp[j][k] -= v*tmp[i][k];
  rep(i,0,n) rep(j,0,n) A[col[i]][col[j]] = tmp[i][j];
  return n;
```

Tridiagonal.h

Description: x = tridiagonal(d, p, q, b) solves the equation system

$$\begin{pmatrix} & b_0 \\ & b_1 \\ & b_2 \\ & b_3 \\ & \vdots \\ & b_{n-1} \end{pmatrix} = \begin{pmatrix} d_0 & p_0 & 0 & 0 & \cdots & 0 \\ q_0 & d_1 & p_1 & 0 & \cdots & 0 \\ 0 & q_1 & d_2 & p_2 & \cdots & 0 \\ \vdots & \vdots & \ddots & \ddots & \ddots & \vdots \\ 0 & 0 & \cdots & q_{n-3} & d_{n-2} & p_{n-2} \\ 0 & 0 & \cdots & 0 & q_{n-2} & d_{n-1} \end{pmatrix} \begin{pmatrix} x_0 \\ x_1 \\ x_2 \\ x_3 \\ \vdots \\ x_{n-1} \end{pmatrix}$$

This is useful for solving problems on the type

$$a_i = b_i a_{i-1} + c_i a_{i+1} + d_i, \ 1 \le i \le n,$$

```
where a_0, a_{n+1}, b_i, c_i and d_i are known. a can then be obtained from
     \{a_i\} = \text{tridiagonal}(\{1, -1, -1, \dots, -1, 1\}, \{0, c_1, c_2, \dots, c_n\},
                        {b_1, b_2, \ldots, b_n, 0}, {a_0, d_1, d_2, \ldots, d_n, a_{n+1}}.
Fails if the solution is not unique.
If |d_i| > |p_i| + |q_{i-1}| for all i, or |d_i| > |p_{i-1}| + |q_i|, or the matrix is positive
definite, the algorithm is numerically stable and neither tr nor the check for
diag[i] == 0 is needed.
Time: \mathcal{O}(N)
                                                              8f9fa8, 26 lines
typedef double T;
vector <T> tridiagonal (vector <T> diag, const vector <T>& super ←
     const vector <T>& sub, vector <T> b) {
  int n = sz(b); vi tr(n);
  rep(i,0,n-1) {
     if (abs(diag[i]) < 1e-9 * abs(super[i])) { // diag[i]} \longrightarrow
       b[i+1] = b[i] * diag[i+1] / super[i];
       if (i+2 < n) b[i+2] = b[i] * sub[i+1] / super[i];
       diag[i+1] = sub[i]; tr[++i] = 1;
       diag[i+1] -= super[i]*sub[i]/diag[i];
       b[i+1] = b[i] * sub[i] / diag[i];
  for (int i = n; i--;) {
    if (tr[i]) {
      swap(b[i], b[i-1]);
       diag[i-1] = diag[i];
      b[i] /= super[i-1];
    } else {
      b[i] /= diag[i];
       if (i) b[i-1] -= b[i]*super[i-1];
  return b;
```

Fourier transforms

FastFourierTransform.h

rolled)

Description: fft(a) computes $\hat{f}(k) = \sum_{x} a[x] \exp(2\pi i \underline{kx/N})$ for all k. Useful for convolution: conv(a, b) = c, where $c[x] = \sum a[i]b[x-i]$. For convolution of complex numbers or more than two vectors: FFT, multiply pointwise, divide by n, reverse(start+1, end), FFT back. Rounding is safe if $(\sum a_i^2 + \sum b_i^2) \log_2 N < 9 \cdot 10^{14}$ (in practice 10^{16} ; higher for random inputs). Otherwise, use long doubles/NTT/FFTMod.

```
Time: \mathcal{O}(N \log N) with N = |A| + |B| (\sim 1s \text{ for } N = 2^{22})
                                                        aca9fa, 35 lines
typedef complex <double> C;
typedef vector <double> vd;
void fft(vector < C>& a) {
 int n = sz(a), L = 31 - builtin clz(n);
 static vector <complex <long double>> R(2, 1);
 static vector <>> rt(2, 1); // (^ 10% faster if double)
 for (static int k = 2; k < n; k *= 2) {
   R. resize(n); rt. resize(n);
    auto x = polar (1.0L, M PI / k); // M PI, lower-case L
    rep(i,k,2*k) rt[i] = R[i] = i&1 ? R[i/2] * x : R[i/2];
 vi rev(n);
 rep(i,0,n) rev[i] = (rev[i / 2] | (i & 1) << L) / 2;
 rep(i,0,n) if (i < rev[i]) swap(a[i], a[rev[i]]);
 for (int k = 1; k < n; k *= 2)
    for (int i = 0; i < n; i += 2 * k) rep(j,0,k) {
      Cz = rt[j+k] * a[i+j+k]; // (25\% faster if hand-<math>\leftarrow
```

```
a[i + j + k] = a[i + j] - z;
      a[i + j] += z;
vd conv(const vd& a, const vd& b) {
 if (a.empty() || b.empty()) return {};
  vd res(sz(a) + sz(b) - 1);
  int L = 32 - builtin clz(sz(res)), n = 1 \ll L;
  vector \langle C \rangle in (n), out (n);
  copy(all(a), begin(in));
  rep(i,0,sz(b)) in[i].imag(b[i]);
  fft (in);
  trav(x, in) x *= x;
  rep(i,0,n) out[i] = in[-i & (n-1)] - conj(in[i]);
  rep(i,0,sz(res)) res[i] = imag(out[i]) / (4 * n);
  return res;
```

FastFourierTransformMod.h

Description: Higher precision FFT, can be used for convolutions modulo arbitrary integers as long as $N \log_2 N \cdot \text{mod} < 8.6 \cdot 10^{14}$ (in practice 10^{16} or higher). Inputs must be in [0, mod].

Time: $\mathcal{O}(N \log N)$, where N = |A| + |B| (twice as slow as NTT or FFT) "FastFourierTransform.h"

```
typedef vector < ll > vl:
template<int M vl convMod(const vl &a, const vl &b) {
 if (a.empty() || b.empty()) return {};
  vl res(sz(a) + sz(b) - 1);
 int B=32- builtin clz(sz(res)), n=1<<B, cut=int(sqrt(M));</pre>
  vector \langle C \rangle L(n), R(n), outs(n), outl(n);
  rep(i,0,sz(a)) L[i] = C((int)a[i] / cut, (int)a[i] % cut);
  rep(i,0,sz(b)) R[i] = C((int)b[i] / cut, (int)b[i] % cut);
  fft(L), fft(R);
  rep(i,0,n) {
    int j = -i & (n - 1);
    outl[j] = (L[i] + conj(L[j])) * R[i] / (2.0 * n);
    outs[j] = (L[i] - conj(L[j])) * R[i] / (2.0 * n) / 1i;
  fft (outl), fft (outs);
  rep(i,0,sz(res)) {
    ll av = ll(real(outl[i])+.5), cv = ll(imag(outs[i])+.5);
    ll bv = ll(imag(outl[i])+.5) + ll(real(outs[i])+.5);
    res[i] = ((av % M * cut + bv) % M * cut + cv) % M;
  return res;
```

NumberTheoreticTransform.h

Description: Can be used for convolutions modulo specific nice primes of the form $2^a b + 1$, where the convolution result has size at most 2^a . Inputs must be in [0, mod).

Time: $\mathcal{O}(N \log N)$

```
"../number-theory/ModPow.h"
const ll mod = (119 << 23) + 1, root = 62; // = 998244353
// For p < 2^30 there is also e.g. 5 \ll 25, 7 \ll 26, 479 \ll \leftarrow
// and 483 \ll 21 (same root). The last two are > 10^9.
typedef vector < ll > vl;
void ntt(vl& a, vl& rt, vl& rev, int n) {
 rep(i,0,n) if (i < rev[i]) swap(a[i], a[rev[i]]);
  for (int k = 1; k < n; k *= 2)
    for (int i = 0; i < n; i += 2 * k) rep(j,0,k) {
        ll z = rt[j + k] * a[i + j + k] % mod, & ai = a[i + j \leftarrow
        a[i + j + k] = (z > ai ? ai - z + mod : ai - z);
        ai += (ai + z >= mod ? z - mod : z);
```

```
vl conv(const vl& a, const vl& b) {
  if (a.empty() || b.empty())
    return {};
  int s = sz(a)+sz(b)-1, B = 32 - builtin clz(s), n = 1 << \leftarrow
  vl L(a), R(b), out(n), rt(n, 1), rev(n);
  L. resize(n), R. resize(n);
  rep(i,0,n) rev[i] = (rev[i / 2] | (i & 1) << B) / 2;
  11 \text{ curL} = \text{mod} / 2, \text{inv} = \text{modpow(n, mod - 2)};
  for (int k = 2; k < n; k *= 2) {
    ll z[] = {1, modpow(root, curL /= 2)};
    rep(i,k,2*k) rt[i] = rt[i / 2] * z[i & 1] % mod;
  ntt(L, rt, rev, n); ntt(R, rt, rev, n);
  rep(i,0,n) out[-i & (n-1)] = L[i] * R[i] % mod * inv % mod←
  ntt(out, rt, rev, n);
  return {out.begin(), out.begin() + s};
FastSubsetTransform.h
Description: Transform to a basis with fast convolutions of the form
```

 $c[z] = \sum_{z = x \oplus y} a[x] \cdot b[y], \text{ where } \oplus \text{ is one of AND, OR, XOR.}$ The size of a must be a power of two.

Time: $\mathcal{O}(N \log N)$

3de473, 16 lines

```
void FST(vi& a, bool inv) {
 for (int n = sz(a), step = 1; step < n; step *= 2) {
   for (int i = 0; i < n; i += 2 * step) rep(j,i,i+step) {
     int &u = a[j], &v = a[j + step]; tie(u, v) =
       inv ? pii(v - u, u) : pii(v, u + v); // AND
       inv ? pii(v, u - v) : pii(u + v, u); // OR
       pii(u + v, u - v);
 if (inv) trav(x, a) x /= sz(a); // XOR only
vi conv(vi a, vi b) {
 FST(a, 0); FST(b, 0);
 rep(i,0,sz(a)) a[i] *= b[i];
 FST(a, 1); return a;
```

Number theory (5)

5.1 Modular arithmetic

Modular Arithmetic.h

Description: Operators for modular arithmetic. You need to set mod to some number first and then you can use the structure. 35bfea, 18 lines

```
const ll mod = 17; // change to something else
struct Mod {
  ll x;
 Mod(ll xx) : x(xx) \{ \}
 Mod operator+(Mod b) { return Mod((x + b.x) % mod); }
  Mod operator-(Mod b) { return Mod((x - b.x + mod) % mod); \leftarrow
  Mod operator*(Mod b) { return Mod((x * b.x) % mod); }
  Mod operator/(Mod b) { return *this * invert(b); }
  Mod invert (Mod a) {
   ll x, y, g = euclid(a.x, mod, x, y);
    assert (g == 1); return Mod((x + mod) \% mod);
```

```
Mod operator^(ll e) {
    if (!e) return Mod(1);
    Mod r = *this ^ (e / 2); r = r * r;
    return e&1 ? *this * r : r;
ModInverse.h
Description: Pre-computation of modular inverses. Assumes LIM ≤ mod
and that mod is a prime.
const ll mod = 1000000007, LIM = 200000;
ll * inv = new ll[LIM] - 1; inv[1] = 1;
rep(i,2,LIM) inv[i] = mod - (mod / i) * inv[mod % i] % mod;
ModPow.h
                                                           b83e45, 8 lines
const ll mod = 1000000007: // faster if const
ll modpow(ll b, ll e) {
  ll ans = 1:
  for (; e; b = b * b % mod, e /= 2)
    if (e & 1) ans = ans * b % mod;
  return ans:
ModLog.h
Description: Returns the smallest x \geq 0 s.t. a^x = b \pmod{m}. a and m
must be coprime.
Time: \mathcal{O}\left(\sqrt{m}\right)
ll modLog(ll a, ll b, ll m) {
  assert(_{gcd}(a, m) == 1);
ll n = (\overline{ll}) sqrt(m) + 1, e = 1, x = 1, res = \underline{HONG\_MAX};
  unordered map<ll, ll>f;
  rep(i,0,n) e = e * a % m;
  rep(i,0,n) x = x * e % m, f.emplace(x, i + 1);
  rep(i,0,n) if (f.count(b = b * a % m))
    res = min(res, f[b] * n - i - 1);
  return res:
ModSum.h
Description: Sums of mod'ed arithmetic progressions.
modsum(to, c, k, m) = \sum_{i=0}^{\text{to}-1} (ki+c)\%m. divsum is similar but for
floored division.
Time: \log(m), with a large constant.
                                                          5c5bc5, 16 lines
```

```
typedef unsigned long long ull;
ull sumsq(ull to) { return to / 2 * ((to-1) | 1); }
ull divsum(ull to, ull c, ull k, ull m) {
 ull res = k / m * sumsq(to) + c / m * to;
 k %= m; c %= m;
 if (!k) return res;
 ull to2 = (to \star k + c) / m;
 return res + (to - 1) * to2 - divsum(to2, m-1 - c, m, k);
ll modsum(ull to, ll c, ll k, ll m) {
 c = ((c \% m) + m) \% m;
 k = ((k \% m) + m) \% m;
 return to * c + k * sumsq(to) - m * divsum(to, c, k, m);
```

ModMulLL.h

Description: Calculate $a \cdot b \mod c$ (or $a^b \mod c$) for $0 \le a, b < c < 2^{63}$. **Time:** $\mathcal{O}(1)$ for mod_mul, $\mathcal{O}(\log b)$ for mod_pow 88c37a, 12 lines

```
typedef unsigned long long ull;
```

```
typedef long double ld;
ull mod mul(ull a, ull b, ull M) {
 ll ret = a * b - M * ull(ld(a) * ld(b) / ld(M));
 return ret + M * (ret < 0) - M * (ret >= (11)M);
ull mod pow(ull b, ull e, ull mod) {
 ull ans = 1;
 for (; e; b = mod mul(b, b, mod), e /= 2)
   if (e & 1) ans = mod mul(ans, b, mod);
ModSgrt.h
```

Description: Tonelli-Shanks algorithm for modular square roots. Finds x s.t. $x^2 = a \pmod{p}$ (-x gives the other solution).

Time: $\mathcal{O}(\log^2 p)$ worst case, $\mathcal{O}(\log p)$ for most p"ModPow.h"

19a793, 24 lines

```
ll sgrt(ll a, ll p) {
 a \% = p; if (a < 0) a += p;
 if (a == 0) return 0;
 assert (modpow(a, (p-1)/2, p) == 1); // else no solution
 if (p % 4 == 3) return modpow(a, (p+1)/4, p);
 // a^{(n+3)/8} or 2^{(n+3)/8} * 2^{(n-1)/4} works if p % 8 == 5
 ll \ s = p - 1, n = 2;
 int r = 0, m;
 while (s \% 2 == 0)
   ++r s /= 2:
 while (modpow(n, (p - 1) / 2, p) != p - 1) ++n;
 11 x = modpow(a, (s + 1) / 2, p);
 11 b = modpow(a, s, p), g = modpow(n, s, p);
 for (;; r = m) {
   11 t = b;
   for (m = 0; m < r && t != 1; ++m)
    t = t * t % p;
   if (m == 0) return x;
   ll gs = modpow(g, 1LL \ll (r - m - 1), p);
   g = gs * gs % p;
   x = x * gs % p;
   b = b * g % p;
```

Primality

MillerRabin.h

Description: Deterministic Miller-Rabin primality test. Guaranteed to work for numbers up to 2^{64} ; for larger numbers, extend A randomly.

Time: 7 times the complexity of $a^{\overline{b}} \mod c$.

```
"ModMulLL.h"
                                                        6ab8e1, 12 lines
bool isPrime(ull n) {
 if (n < 2 | | n % 6 % 4 != 1) return n - 2 < 2;
  ull A[] = \{2, 325, 9375, 28178, 450775, 9780504, \leftarrow
       1795265022},
      s = builtin ctzll(n-1), d = n >> s;
  trav(a, \bar{A})^{-}\{ // count trailing zeroes
    ull p = mod pow(a, d, n), i = s;
    while (p != 1 && p != n - 1 && a % n && i--)
     p = mod mul(p, p, n);
    if (p != \overline{n}-1 && i != s) return 0;
 return 1;
```

Description: Pollard-rho randomized factorization algorithm. Returns prime factors of a number, in arbitrary order (e.g. 2299 -> {11, 19, 11}).

Time: $\mathcal{O}\left(n^{1/4}\right)$ gcd calls, less for numbers with small factors.

"ModMulLL.h", "MillerRabin.h" f5adaa, 18 lines

```
ull pollard(ull n) {
    auto f = [n](ull x) { return (mod_mul(x, x, n) + 1) % n; ←
    };
    if (!(n & 1)) return 2;
    for (ull i = 2;; i++) {
        ull x = i, y = f(x), p;
        while ((p = __gad(n + y - x, n)) == 1)
            x = f(x), y = f(f(y));
        if (p != n) return p;
    }
}
vector<ull>    factor(ull n) {
    if (n == 1) return {};
    if (isPrime(n)) return {n};
    ull x = pollard(n);
    auto l = factor(x), r = factor(n / x);
    l.insert(l.end(), all(r));
    return l;
}
```

5.3 Divisibility

euclid.h

Description: Finds two integers x and y, such that $ax + by = \gcd(a, b)$. If you just need gcd, use the built in __gcd instead. If a and b are coprime, then x is the inverse of $a \pmod{b}$.

```
ll euclid(ll a, ll b, ll &x, ll &y) {
  if (b) { ll d = euclid(b, a % b, y, x);
    return y -= a/b * x, d; }
  return x = 1, y = 0, a;
}
```

EuclidBigInt.java

Description: Finds $\{x, y, d\}$ s.t. ax + by = d = gcd(a, b). _{6aba01, 11 lines}

```
static BigInteger[] euclid(BigInteger a, BigInteger b) {
   BigInteger x = BigInteger.ONE, yy = x;
   BigInteger y = BigInteger.ZERO, xx = y;
   while (b.signum() != 0) {
      BigInteger q = a.divide(b), t = b;
      b = a.mod(b); a = t;
      t = xx; xx = x.subtract(q.multiply(xx)); x = t;
      t = yy; yy = y.subtract(q.multiply(yy)); y = t;
   }
   return new BigInteger[]{x, y, a};
}
```

CRT.h

 ${\bf Description:} \ {\bf Chinese} \ {\bf Remainder} \ {\bf Theorem}.$

 $crt(a,\ m,\ b,\ n)$ computes x such that $x\equiv a\pmod m,\ x\equiv b\pmod n.$ If |a|< m and $|b|< n,\ x$ will obey $0\le x< \mathrm{lcm}(m,n).$ Assumes $mn<2^{62}.$ $crt(x,\ a)$ computes z such that $z\pmod x|_i=a_i\forall i.$ Note that the solution is unique modulo $M=\mathrm{lcm}(x_i)$. Return $(z,\ M).$ Note that we do not require the a_i to be relatively prime.

```
Time: N \log(N)
```

```
for(int i = 1; i < int(x.size()); i++) {
    ret = crt(ret.second, ret.first, a[i], x[i]);
}
return ret;</pre>
```

5.3.1 Bézout's identity

For $a \neq b \neq 0$, then d = gcd(a, b) is the smallest positive integer for which there are integer solutions to

$$ax + by = d$$

If (x, y) is one solution, then all solutions are given by

$$\left(x + \frac{kb}{\gcd(a,b)}, y - \frac{ka}{\gcd(a,b)}\right), \quad k \in \mathbb{Z}$$

phiFunction.h

Description: Euler's ϕ function is defined as $\phi(n) := \#$ of positive integers $\leq n$ that are coprime with n. $\phi(1) = 1$, p prime $\Rightarrow \phi(p^k) = (p-1)p^{k-1}$, m, n coprime $\Rightarrow \phi(mn) = \phi(m)\phi(n)$. If $n = p_1^{k_1}p_2^{k_2}...p_r^{k_r}$ then $\phi(n) = (p_1-1)p_1^{k_1-1}...(p_r-1)p_r^{k_r-1}$. $\phi(n) = n \cdot \prod_{p|n} (1-1/p)$. $\sum_{d|n} \phi(d) = n$, $\sum_{1 \leq k \leq n, \gcd(k,n)=1} k = n\phi(n)/2, n > 1$

Euler's thm: a, n coprime $\Rightarrow a^{\phi(n)} \equiv 1 \pmod{n}$. Fermat's little thm: p prime $\Rightarrow a^{p-1} \equiv 1 \pmod{p} \ \forall a$.

```
const int LIM = 5000000;
int phi[LIM];

void calculatePhi() {
  rep(i,0,LIM) phi[i] = i&1 ? i : i/2;
  for(int i = 3; i < LIM; i += 2) if(phi[i] == i)</pre>
```

for(int j = i; j < LIM; j += i) phi[j] -= phi[j] / i;</pre>

5.4 Fractions

ContinuedFractions.h

Description: Given N and a real number $x \ge 0$, finds the closest rational approximation p/q with $p,q \le N$. It will obey $|p/q - x| \le 1/qN$.

For consecutive convergents, $p_{k+1}q_k - q_{k+1}p_k = (-1)^k$. $(p_k/q_k$ alternates between > x and < x.) If x is rational, y eventually becomes ∞ ; if x is the root of a degree 2 polynomial the a's eventually become cyclic.

Time: $\mathcal{O}(\log N)$ dd6c5e, 21 line

```
make_pair(NP, NQ) : make_pair(P, Q);
}
if (abs(y = 1/(y - (d)a)) > 3*N) {
    return {NP, NQ};
}
LP = P; P = NP;
LQ = Q; Q = NQ;
}
```

FracBinarySearch.h

Description: Given f and N, finds the smallest fraction $p/q \in [0,1]$ such that f(p/q) is true, and $p, q \leq N$. You may want to throw an exception from f if it finds an exact solution, in which case N can be removed. Usage: fracBS($\lceil \rceil$ (Frac f) { return f.p>=3*f.q; }, 10); // {1,3}

Usage: fracBS([](Frac f) { return f.p>=3*f.q; }, 10); // {1,3} $\underline{\text{Time: } \mathcal{O}(\log(N))}$ $\underline{\phantom{\text{Time: } \mathcal{O}(\log(N))}}$ $\underline{\phantom{\text{Time: } \mathcal{O}(\log(N))}}$

```
struct Frac { ll p, q; };
template<class F>
Frac fracBS(F f, 11 N) {
  bool dir = 1, A = 1, B = 1;
  Frac lo (0, 1), hi (1, 1); // Set hi to 1/0 to search (0, N)
  if (f(lo)) return lo;
  assert(f(hi));
  while (A | | B) {
    ll adv = 0, step = 1; // move hi if dir, else lo
    for (int si = 0; step; (step *= 2) >>= si) {
      Frac mid{lo.p * adv + hi.p, lo.q * adv + hi.q};
      if (abs(mid.p) > N | | mid.q > N | | dir == !f(mid)) {
        adv -= step; si = 2;
    hi \cdot p += lo \cdot p * adv:
    hi.q += lo.q * adv;
    dir = !dir;
    swap(lo, hi);
   A = B; B = !!adv;
  return dir ? hi : lo;
```

5.5 Pythagorean Triples

The Pythagorean triples are uniquely generated by

```
a = k \cdot (m^2 - n^2), b = k \cdot (2mn), c = k \cdot (m^2 + n^2),
```

with m > n > 0, k > 0, $m \perp n$, and either m or n even.

5.6 Primes

p=962592769 is such that $2^{21}\mid p-1$, which may be useful. For hashing use 970592641 (31-bit number), 31443539979727 (45-bit), 3006703054056749 (52-bit). There are 78498 primes less than $1\,000\,000$.

Primitive roots exist modulo any prime power p^a , except for p=2, a>2, and there are $\phi(\phi(p^a))$ many. For p=2, a>2, the group $\mathbb{Z}_{2^a}^{\times}$ is instead isomorphic to $\mathbb{Z}_2 \times \mathbb{Z}_{2^{a-2}}$.

Some other primes are:

```
 \begin{vmatrix} 10^9 + 7, 10^9 + 9, 10^9 + 21, 10^9 + 33, 10^9 + 87, 10^9 + 93, 10^9 + 97, 10^9 + 103, 10^9 + 123, 10^9 + 181, 10^9 + 207, 10^9 + 223 \end{vmatrix}
```

Estimates

 $\sum_{d|n} d = O(n \log \log n).$

The number of divisors of n is at most around 100 for n < 5e4, 500 for n < 1e7, 2000 for n < 1e10, 200 000 for n < 1e19.

Mobius function

$$\mu(n) = \begin{cases} 0 & n \text{ is not square free} \\ 1 & n \text{has even number of prime factors} \\ -1 & n \text{has odd number of prime factors} \end{cases}$$

We note that $\mu(ab) = \mu(a)\mu(b)$ if a and b are relatively prime.

Inversion formula: If $g(n) = \sum_{d|n} f(d)$, then $f(n) = \sum_{d|n} \mu(d)g(n/d).$

Combinatorial (6)

6.1 Permutations

6.1.1 Factorial

	n	1 2 3	4	5 6	7	8	9	10	
_	n!	1 2 6	24 1	20 72	0 5040	0 40320	362880	3628800	_
	n	11	12	13	14	1	5 16	17	
_	n!	4.0e7	7 4.8e	8 6.2e	9 8.7e	e10 1.3€	e12 2.1e	13 3.6e14	-
	n	20	25	30	40	50 1	00 15	50 17	1
_	n!	2e18	2e25	3e32	8e47	3e64 9e	$157 \ 6e2$	$262 > DBL_$	MAX

6.1.2 Cycles

Let $q_S(n)$ be the number of n-permutations whose cycle lengths all belong to the set S. Then

$$\sum_{n=0}^{\infty} g_S(n) \frac{x^n}{n!} = \exp\left(\sum_{n \in S} \frac{x^n}{n}\right)$$

6.1.3 Burnside's lemma

Given a group G of symmetries and a set X, the number of elements of X up to symmetry equals

$$\frac{1}{|G|} \sum_{g \in G} |X^g|,$$

where X^g are the elements fixed by g (g.x = x).

If f(n) counts "configurations" (of some sort) of length n, we can ignore rotational symmetry using $G = \mathbb{Z}_n$ to get

$$g(n) = \frac{1}{n} \sum_{k=0}^{n-1} f(\gcd(n,k)) = \frac{1}{n} \sum_{k|n} f(k)\phi(n/k).$$

The number of orbits of a set X under the group action Gequals the average number of elements of X fixed by the elements of G.

Here's an example. Consider a square of $2n \times 2n$ cells. How many ways are there to color it into X colors, up to rotations and/or reflections? Here, the group has only 8 elements (rotations by 0, 90, 180, 270 degrees, reflections over two diagonals, over a vertical line and over a horizontal line). Every coloring stays itself after rotating by 0 degrees, so that rotation has X^{4n^2} fixed points. Rotation by 180 degrees and reflections over a horizonal/vertical line split all cells in pairs that must be of the same color for a coloring to be unaffected by such rotation/reflection, thus there exist X^{2n^2} such colorings for each of them. Rotations by 90 and 270 degrees split cells in groups of four, thus yielding X^n fixed colorings. Reflections over diagonals split cells into 2ngroups of 1 (the diagonal itself) and $2n^2 - n$ groups of 2 (all remaining cells), thus yielding $X^{2n^2-n+2n} = X^{2n^2+n}$ affected colorings. So, the answer is: $\underline{X^{4n^2} \! + \! 3X^{2n^2} \! + \! 2X^{n^2} \! + \! 2X^{2n^+n}}$

Partitions and subsets

6.2.1 Partition function

Number of ways of writing n as a sum of positive integers, disregarding the order of the summands.

$$p(0) = 1, \ p(n) = \sum_{k \in \mathbb{Z} \setminus \{0\}} (-1)^{k+1} p(n - k(3k - 1)/2)$$

6.2.2 Binomials

binomialModPrime.h

Description: Lucas' thm: Let n, m be non-negative integers and p a prime. Write $n = n_k p^k + ... + n_1 p + n_0$ and $m = m_k p^k + ... + m_1 p + m_0$. Then $\binom{n}{m} \equiv \prod_{i=0}^k \binom{n_i}{m_i} \pmod{p}$. fact and invfact must hold pre-computed factorials / inverse factorials, e.g. from ModInverse.h. Time: $\mathcal{O}\left(\log_n n\right)$

```
ll chooseModP(ll n, ll m, int p, vi& fact, vi& invfact) {
 ll c = 1;
 while (n | | m) {
    ll a = n \% p, b = m \% p;
    if (a < b) return 0;
    c = c * fact[a] % p * invfact[b] % p * invfact[a - b] % \leftarrow
   n /= p; m /= p;
 return c;
```

```
multinomial.h
Description: Computes \binom{k_1 + \dots + k_n}{k_1, k_2, \dots, k_n}
ll multinomial(vi& v) {
  ll c = 1, m = v.empty() ? 1 : v[0];
  rep(i,1,sz(v)) rep(j,0,v[i])
     c = c * ++m / (j+1);
```

General purpose numbers

6.3.1 Bernoulli numbers

EGF of Bernoulli numbers is $B(t) = \frac{t}{e^t - 1}$ (FFT-able). $B[0,\ldots] = [1,-\frac{1}{2},\frac{1}{6},0,-\frac{1}{30},0,\frac{1}{42},\ldots]$

Sums of powers:

return c;

$$\sum_{i=1}^{n} n^{m} = \frac{1}{m+1} \sum_{k=0}^{m} {m+1 \choose k} B_{k} \cdot (n+1)^{m+1-k}$$

Euler-Maclaurin formula for infinite sums:

$$\sum_{i=m}^{\infty} f(i) = \int_{m}^{\infty} f(x)dx - \sum_{k=1}^{\infty} \frac{B_k}{k!} f^{(k-1)}(m)$$

$$\approx \int_{0}^{\infty} f(x)dx + \frac{f(m)}{2} - \frac{f'(m)}{12} + \frac{f'''(m)}{720} + O(f^{(5)}(m))$$

6.3.2 Stirling numbers of the first kind

Number of permutations on n items with k cycles.

$$c(n,k) = c(n-1,k-1) + (n-1)c(n-1,k), \ c(0,0) = 1$$

$$\sum_{k=0}^{n} c(n,k)x^{k} = x(x+1)\dots(x+n-1)$$

c(8, k) = 8, 0, 5040, 13068, 13132, 6769, 1960, 322, 28, 1 $c(n,2) = 0, 0, 1, 3, 11, 50, 274, 1764, 13068, 109584, \dots$

6.3.3 Eulerian numbers

Number of permutations $\pi \in S_n$ in which exactly k elements are greater than the previous element. k j:s s.t. $\pi(j) > \pi(j+1), k+1 \text{ } j:s \text{ s.t. } \pi(j) \geq j, k \text{ } j:s \text{ s.t.}$

$$\pi(j) > \pi(j+1), \ k+1 \ j$$
:s s.t. $\pi(j) \ge j, \ k \ j$:s $\pi(j) > j$.

$$E(n,k) = (n-k)E(n-1,k-1) + (k+1)E(n-1,k)$$

$$E(n,0) = E(n,n-1) = 1$$

$$E(n,k) = \sum_{j=0}^{k} (-1)^{j} \binom{n+1}{j} (k+1-j)^{n}$$

UCSD

6.3.4 Stirling numbers of the second kind

Partitions of n distinct elements into exactly k groups.

$$S(n,k) = S(n-1,k-1) + kS(n-1,k)$$

$$S(n,1) = S(n,n) = 1$$

$$S(n,k) = \frac{1}{k!} \sum_{j=0}^{k} (-1)^{k-j} \binom{k}{j} j^{n}$$

6.3.5 Bell numbers

Total number of partitions of n distinct elements. B(n) = 1, 1, 2, 5, 15, 52, 203, 877, 4140, 21147, For p prime,

$$B(p^{m} + n) \equiv mB(n) + B(n + 1) \pmod{p}$$
$$B(n) = \sum_{k=0}^{n} \binom{n}{k} B_{k}$$

6.3.6 Labeled unrooted trees(Cayley)

on n vertices: n^{n-2} # on k existing trees of size n_i : $n_1 n_2 \cdots n_k n^{k-2}$ # with degrees d_i : $(n-2)!/((d_1-1)!\cdots(d_n-1)!)$

6.3.7 Catalan numbers

$$C_n = \frac{1}{n+1} {2n \choose n} = {2n \choose n} - {2n \choose n+1} = \frac{(2n)!}{(n+1)!n!}$$

$$C_0 = 1, \ C_{n+1} = \frac{2(2n+1)}{n+2} C_n, \ C_{n+1} = \sum_{n=1}^{\infty} C_n C_{n-n}$$

 $C_n = 1, 1, 2, 5, 14, 42, 132, 429, 1430, 4862, 16796, 58786, \dots$

- $\bullet\,$ sub-diagonal monotone paths in an $n\times n$ grid.
- \bullet strings with n pairs of parenthesis, correctly nested.
- binary trees with with n+1 leaves (0 or 2 children).
- ordered trees with n+1 vertices.
- ways a convex polygon with n+2 sides can be cut into triangles by connecting vertices with straight lines.
- \bullet permutations of [n] with no 3-term increasing subseq.

6.4 General purpose theorems - 1

6.4.1 Identities

Vandermonde Convolution: $\binom{m+n}{r} = \sum_{k=0}^{r} \binom{m}{k} \cdot \binom{r}{n-k}$. Hockey Stick: $\binom{n+1}{r+1} = \sum_{i=r}^{n} \binom{i}{r}$.

6.4.2 Cycle Lemma

Any sequence of mX's and nY's, where m > n has exactly m-n cyclic permutations which are dominating, and m-kn which are k-dominating. To find them, arrange sequence in a circle and repeatedly remove adjacent pairs XY. The remaining X's were each the start of a dominating permutation.

6.4.3 Sprague Grundy theorem

Every impartial game is equivalent to a nimber. Nimbers are de-fined inductively as $*0 = \{\}, *1 = *0, *2 = *0, *1, *(n+1) = *n \cup n$, and

*0 = {},*1 = *0,*2 = *0,*1,*(n+1) = *n \cup n, and corresponds to a heap of size n. The formula for adding positions is $S + S' = S + s'|s' \in S' \cup s + S'|s \in S$. $a + b = a \oplus b + 2(a\&b)$.

Define minimum exclusion $M: \phi(N) \to N$ by M(S) = the least non-negative integer not in S. Let $C = (M(A) \oplus B) \cup (M(B) \oplus A)$. Then $M(C) = M(A) \oplus M(B)$. Define $SG(S) = M(\{SG(s)|s \in S\})$. $SG(Nim_k) = k$ by strong induction. Game is losing iff SG(S) = 0. Theorem:

6.4.4 Partisan Game

 $SG(A+B) = SG(A) \oplus SG(B).$

Can define the negative of a game by interchanging L and R's possible moves. Define G=0 if first player loses. G=H if G+(-H)=0. A cold game is one which moving only hurts players. In this case we never have G fuzzy 0, so G is representable as an integer, thus calculable by DP.

6.4.5 Matrices for operators

Matrices for xor, and, and or are: $\frac{1}{\sqrt{2}}\begin{bmatrix} 1 & 1\\ 1 & -1 \end{bmatrix}, \begin{bmatrix} 0 & 1\\ 1 & 1 \end{bmatrix}$ with inverses: $\begin{bmatrix} -1 & 1\\ 1 & 0 \end{bmatrix}, \begin{bmatrix} 1 & 1\\ 1 & 0 \end{bmatrix}$.

6.5 General purpose theorems - 2

6.5.1 Prufer sequences

The set of labeled trees on n vertices corresponds bijectively to the set of Prufer sequences of length n-2. To convert a tree into a Prufer sequence, repeatedly remove the leaf with the smallest label, and write down its neighbor. To convert sequence to tree, first set the degree of each vertex to $n_v + 1$, where n_v is the number of times the vertex appears in the sequence. Then for each i, find lowest j with degree 1, add edge a_i, j , and decrease the degrees of a_i and j by 1. After this, two nodes of degree 1 remain - connect them.

This can be used to calculate number of labeled trees in a complete bipartite graph - $l^{r-1} \cdot r^{l-1}$.

12

6.5.2 Tournament Graphs

There exists a Hamiltonian path on any tournament graphs - use induction to find. Cycle if strongly connected. TFAE:

- 1. T is transitive.
- 2. T is strict total ordering.
- 3. T is acyclic.
- 4. T has no cycle of length 3.
- 5. The outdegrees are $\{0, 1, \dots, n-1\}$.
- 6. T has exactly one Hamiltonian path.

Landau's theorem: A sequence of numbers is called a score sequence if for each subset S, sum of numbers in S is at least $\binom{|S|}{2}$ and sum of all numbers is $\binom{n}{2}$.

This score sequence represents the outdegrees of a vertex in

6.6 terrogeneral purpose theorems - 3

6.6.1 Dilworth's / Hall's / Mirsky's theorem

Maximum antichain has same size as minimum chain decomposition.

Maximum chain size has same size as minimum antichain decomposition.

To compute size, model as bipartite graph with two copies of vertices - $v_i n$ and $v_o ut$. Distinct representatives can be chosen for a family of sets S iff every subfamily W of S has at least |W| elements in their union. E.g. Left side of bipartite graph can be fully matched iff each subset has sufficient "degree".

6.6.2 Laplacian Matrix and Kirchoff's Theorem

Laplacian matrix is defined as L = D - A, where D is the degree matrix (diagonal), and A the adjacency matrix.

Kirchoff's Theorem states that the number of spanning trees in a graph is any cofactor of the Laplacian.

To calculate that, remove the first row and column and calculate the determinant of the remaining matrix.

6.6.3 Derangements

Permutations of a set such that none of the elements appear in their original position.

$$D_n = n! \sum_{k=0}^{n} \frac{(-1)^k}{k!} = (n-1)(D_{n-2} - D_{n-1}) = \left\lfloor \frac{n!}{e} \right\rfloor$$

Graph (7)

7.1 Euler walk

EulerWalk.h

Description: Eulerian undirected/directed path/cycle algorithm. Input should be a vector of (dest, global edge index), where for undirected graphs, forward/backward edges have the same index. Returns a list of nodes in the Eulerian path/cycle with src at both start and end, or empty list if no cycle/path exists. To get edge indices back, add .second to s and ret.

```
Time: \mathcal{O}(V+E)
vi eulerWalk(vector < vector < pii >>& gr, int nedges, int src=0) ←
  int n = sz(gr);
  vi D(n), its(n), eu(nedges), ret, s = {src};
  D[src]++; // to allow Euler paths, not just cycles
  while (!s.emptv()) {
    int x = s.back(), y, e, &it = its[x], end = sz(gr[x]);
    if (it == end){ ret.push back(x); s.pop back(); continue←
    tie(y, e) = gr[x][it++];
    if (!eu[e]) {
     D[x] --, D[y] ++;
      eu[e] = 1; s.push back(y);
  trav(x, D) if (x < 0 \mid \mid sz(ret) != nedges+1) return {};
  return {ret.rbegin(), ret.rend()};
```

Network flow

PushRelabel.h

Description: Push-relabel using the highest label selection rule and the gap heuristic. Quite fast in practice. To obtain the actual flow, look at positive values only.

```
Time: \mathcal{O}\left(V^2\sqrt{E}\right)
                                                          3df61b, 50 lines
typedef ll Flow;
struct Edge {
  int dest, back;
  Flow f, c;
};
struct PushRelabel {
  vector < vector < Edge >> g;
  vector <Flow> ec;
  vector < Edge*> cur;
  vector <vi> hs; vi H;
  PushRelabel(int n): g(n), ec(n), cur(n), hs(2*n), H(n) {}
  void add edge(int s, int t, Flow cap, Flow rcap=0) {
    if (s == t) return;
    g[s].push back({t, sz(g[t]), 0, cap});
    g[t].push back({s, sz(g[s])-1, 0, rcap});
```

```
void add flow (Edge& e, Flow f) {
    Edge &back = g[e.dest][e.back];
    if (!ec[e.dest] && f) hs[H[e.dest]].push back(e.dest);
    e.f += f; e.c -= f; ec[e.dest] += f;
    back.f -= f; back.c += f; ec[back.dest] -= f;
  Flow maxflow(int s, int t) {
    int v = sz(g); H[s] = v; ec[t] = 1;
    vi co(2*v); co[0] = v-1;
    rep(i,0,v) cur[i] = g[i].data();
    trav(e, g[s]) add flow(e, e.c);
    for (int hi = 0;;) {
      while (hs[hi].empty()) if (!hi--) return -ec[s];
      int u = hs[hi].back(); hs[hi].pop back();
      while (ec[u] > 0) // discharge u
        if (\operatorname{cur}[u] == g[u] \cdot \operatorname{data}() + \operatorname{sz}(g[u])) {
          H[u] = 1e9;
           trav(e, g[u]) if (e.c && H[u] > H[e.dest]+1)
             H[u] = H[e.dest] + 1, cur[u] = &e;
           if (++co[H[u]], !--co[hi] && hi < v)
             rep(i,0,v) if (hi < H[i] && H[i] < v)
                --co[H[i]], H[i] = v + 1;
           hi = H[u]:
        } else if (\operatorname{cur}[u] \rightarrow \operatorname{c \&\& H[u]} == \operatorname{H[cur}[u] \rightarrow \operatorname{dest}] + 1)
          add flow(*cur[u], min(ec[u], cur[u]->c));
         else ++cur[u];
  bool leftOfMinCut(int a) { return H[a] >= sz(g); }
MinCostMaxFlow.h
Description: Min-cost max-flow. cap[i][j] != cap[j][i] is allowed; double
edges are not. If costs can be negative, call setpi before maxflow, but note
that negative cost cycles are not supported. To obtain the actual flow, look
at positive values only.
Time: Approximately \mathcal{O}(E^2)
                                                          6915ce, 81 lines
#include <bits/extc++.h>
const ll INF = numeric limits<ll>::max() / 4;
typedef vector < ll > VL;
struct MOMF {
  int N:
  vector <vi> ed, red;
  vector <VL> cap, flow, cost;
  vi seen:
  VL dist, pi;
  vector < pii > par;
 MCMF(int N):
    N(N), ed(N), red(N), cap(N, VL(N)), flow(cap), cost(cap)\leftarrow
    seen(N), dist(N), pi(N), par(N) {}
  void addEdge(int from, int to, ll cap, ll cost) {
    this->cap[from][to] = cap;
    this->cost[from][to] = cost;
    ed[from].push back(to);
    red[to].push back(from);
  void path(int s) {
    fill (all (seen), 0);
    fill (all (dist), INF);
    dist[s] = 0; ll di;
```

```
gnu pbds::priority queue <pair <11, int>> q;
    vector < decltype (q)::point iterator > its (N);
    q.push({0, s});
    auto relax = [&](int i, ll cap, ll cost, int dir) {
      ll val = di - pi[i] + cost;
      if (cap && val < dist[i]) {
        dist[i] = val;
        par[i] = \{s, dir\};
        if (its[i] == q.end()) its[i] = q.push(\{-dist[i], i \leftarrow
        else q.modify(its[i], {-dist[i], i});
    };
    while (!q.emptv()) {
      s = q.top().second; q.pop();
      seen[s] = 1; di = dist[s] + pi[s];
      trav(i, ed[s]) if (!seen[i])
        relax(i, cap[s][i] - flow[s][i], cost[s][i], 1);
      trav(i, red[s]) if (!seen[i])
        relax(i, flow[i][s], -cost[i][s], 0);
    rep(i,0,N) pi[i] = min(pi[i] + dist[i], INF);
  pair < ll, ll > maxflow(int s, int t) {
    ll totflow = 0, totcost = 0:
    while (path(s), seen[t]) {
      11 fl = INF:
      for (int p,r,x = t; tie(p,r) = par[x], x != s; x = p)
        fl = min(fl, r ? cap[p][x] - flow[p][x] : flow[x][p\leftarrow
      totflow += fl;
      for (int p,r,x = t; tie(p,r) = par[x], x != s; x = p)
        if (r) flow[p][x] += fl;
        else flow[x][p] -= fl;
    rep(i,0,N) rep(j,0,N) totcost += cost[i][j] * flow[i][j\leftarrow
    return {totflow, totcost};
  // If some costs can be negative, call this before maxflow-
  void setpi(int s) { // (otherwise, leave this out)
    fill(all(pi), INF); pi[s] = 0;
    int it = \hat{N}, ch = 1; \hat{l}l v;
    while (ch-- && it--)
      rep(i,0,N) if (pi[i] != INF)
        trav(to, ed[i]) if (cap[i][to])
          if ((v = pi[i] + cost[i][to]) < pi[to])
            pi[to] = v, ch = 1;
    assert(it >= 0); // negative cost cycle
};
```

EdmondsKarp.h

Description: Flow algorithm with guaranteed complexity $O(VE^2)$. To get edge flow values, compare capacities before and after, and take the positive values only.

```
template < class T > T edmonds Karp (vector < unordered map < int, T -
    >>& graph, int source, int sink) {
  assert (source != sink);
 T flow = 0;
 vi par(sz(graph)), q = par;
  for (;;) {
```

```
fill (all (par), -1);
    par[source] = 0;
    int ptr = 1;
    q[0] = source;
    rep(i,0,ptr) {
      int x = q[i];
      trav(e, graph[x]) {
        if (par[e.first] == -1 && e.second > 0) {
          par[e.first] = x;
          q[ptr++] = e.first;
          if (e.first == sink) goto out;
   }
    return flow;
    T inc = numeric limits <T>::max();
    for (int y = sink; y != source; y = par[y])
     inc = min(inc, graph[par[y]][y]);
    flow += inc:
    for (int y = sink; y != source; y = par[y]) {
      int p = par[v];
     if ((graph[p][y] -= inc) \le 0) graph[p].erase(y);
      graph[y][p] += inc;
Dinic.h
Description: Flow algorithm with complexity O(VE \log U) where U =
max |cap|. O(\min(E^{1/2}, V^{2/3})E) if U = 1; O(\sqrt{V}E) for bipartite match-
ing.
                                                      f688cf, 41 lines
struct Dinic {
  struct Edge {
    int to, rev;
    ll c, oc;
    ll flow() { return max(oc - c, 0LL); } // if you need ←
         flows
  };
  vi lvl, ptr, q;
  vector < vector < Edge >> adj;
  Dinic(int n): lvl(n), ptr(n), q(n), adj(n) {}
  void addEdge(int a, int b, ll c, int rcap = 0) {
    adj[a].push back({b, sz(adj[b]), c, c});
    adj[b].push back({a, sz(adj[a]) - 1, rcap, rcap});
  ll dfs(int v, int t, ll f) {
    if (v == t \mid | !f) return f;
    for (int& i = ptr[v]; i < sz(adj[v]); i++) {
      Edge& e = adj[v][i];
     if (lvl[e.to] == lvl[v] + 1)
        if (ll p = dfs(e.to, t, min(f, e.c))) {
          e.c -= p, adj[e.to][e.rev].c += p;
          return p;
    return 0;
  ll calc(int s, int t) {
    ll flow = 0; q[0] = s;
    rep(L,0,31) do { // 'int L=30' maybe faster for random ←
         data
      lvl = ptr = vi(sz(q));
```

int qi = 0, qe = lvl[s] = 1;

while (qi < qe && !lvl[t]) {

int v = q[qi++];

```
trav(e, adj[v])
         if (!lvl[e.to] && e.c >> (30 - L))
           q[qe++] = e.to, lvl[e.to] = lvl[v] + 1;
     while (ll p = dfs(s, t, LLONG MAX)) flow += p;
   } while (lvl[t]);
    return flow;
};
GlobalMinCut.h
```

Description: Find a global minimum cut in an undirected graph, as represented by an adjacency matrix.

```
Time: \mathcal{O}\left(V^3\right)
                                                       03261f, 31 lines
pair <int, vi > GetMinCut(vector < vi > & weights) {
  int N = sz(weights);
  vi used(N), cut, best cut;
  int best weight = -1;
  for (int phase = N-1; phase >= 0; phase--) {
    vi w = weights[0], added = used;
    int prev, k = 0;
    rep(i,0,phase){
      prev = k;
      k = -1:
      rep(j,1,N)
        if (!added[j] && (k == -1 || w[j] > w[k])) k = j;
      if (i == phase-1) {
        rep(j,0,N) weights[prev][j] += weights[k][j];
        rep(j,0,N) weights[j][prev] = weights[prev][j];
        used[k] = true:
        cut.push back(k);
        if (best_weight == -1 || w[k] < best_weight) {</pre>
          best cut = cut;
          best weight = w[k];
      } else {
        rep(j,0,N)
          w[j] += weights[k][j];
        added[k] = true;
 return {best weight, best cut};
```

7.3 Matching

hopcroftKarp.h

Description: Fast bipartite matching algorithm. Graph g should be a list of neighbors of the left partition, and btoa should be a vector full of -1's of the same size as the right partition. Returns the size of the matching. btoa[i]will be the match for vertex i on the right side, or -1 if it's not matched. Usage: vi btoa(m, -1); hopcroftKarp(g, btoa);

```
Time: \mathcal{O}\left(\sqrt{V}E\right)
bool dfs(int a, int L, vector < vi>& g, vi& btoa, vi& A, vi& B←
 if (A[a] != L) return 0;
 A[a] = -1;
 trav(b, g[a]) if (B[b] == L + 1) {
   B[b] = 0;
    if (btoa[b] == -1 || dfs(btoa[b], L + 1, g, btoa, A, B))
      return btoa[b] = a, 1;
 }
 return 0;
```

```
int hopcroftKarp(vector < vi > & g, vi & btoa) {
  int res = 0;
  vi A(g.size()), B(btoa.size()), cur, next;
  for (;;) {
    fill (all (A), 0);
    fill (all (B), 0);
    cur.clear();
    trav(a, btoa) if(a != -1) A[a] = -1;
    rep(a,0,sz(g)) if(A[a] == 0) cur.push back(a);
    for (int lay = 1;; lay++) {
      bool islast = 0;
      next.clear();
      trav(a, cur) trav(b, g[a]) {
        if (btoa[b] == -1) {
          B[b] = lav;
          islast = 1;
        else if (btoa[b] != a && !B[b]) {
          B[b] = lay;
          next.push back(btoa[b]);
      if (islast) break;
      if (next.empty()) return res;
      trav(a, next) A[a] = lay;
      cur.swap(next);
    rep(a,0,sz(g))
      res += dfs(a, 0, g, btoa, A, B);
}
```

DFSMatching.h

Description: Simple bipartite matching algorithm. Graph g should be a list of neighbors of the left partition, and btoa should be a vector full of -1's of the same size as the right partition. Returns the size of the matching. btoa[i]will be the match for vertex i on the right side, or -1 if it's not matched.

Usage: vi btoa(m, -1); dfsMatching(g, btoa);

```
Time: \mathcal{O}(VE)
                                                      6a3472, 22 lines
bool find(int j, vector < vi>& g, vi& btoa, vi& vis) {
 if (btoa[j] == -1) return 1;
  vis[i] = 1; int di = btoa[i];
  trav(e, g[di])
    if (!vis[e] && find(e, g, btoa, vis)) {
      btoa[e] = di;
      return 1;
  return 0;
int dfsMatching(vector < vi>& g, vi& btoa) {
  vi vis:
  rep(i,0,sz(g)) {
    vis.assign(sz(btoa), 0);
    trav(j,g[i])
      if (find(j, g, btoa, vis)) {
        btoa[j] = i;
        break:
 return sz(btoa) - (int)count(all(btoa), -1);
```

MinimumVertexCover.h

Description: Finds a minimum vertex cover in a bipartite graph. The size is the same as the size of a maximum matching, and the complement is a maximum independent set.

```
"DFSMatching.h"
                                                             d0b3f2, 20 lines
vi cover(vector < vi>& g, int n, int m) {
```

vi match (m, -1);

while (!q.empty()) {

vi q, cover;

int res = dfsMatching(g, match);

vector <bool> lfound(n, true), seen(m);

rep(i,0,n) if (lfound[i]) q.push back(i);

trav(it, match) if (it != -1) lfound[it] = false;

```
int i = q.back(); q.pop back();
    lfound[i] = 1;
    trav(e, g[i]) if (!seen[e] && match[e] != -1) {
      seen[e] = true;
      q.push back(match[e]);
  rep(i,0,n) if (!lfound[i]) cover.push back(i);
  rep(i,0,m) if (seen[i]) cover.push back(n+i);
  assert(sz(cover) == res);
  return cover;
WeightedMatching.h
Description: Min cost bipartite matching. Negate costs for max cost.
Time: \mathcal{O}(N^3)
                                                       055ca9, 75 lines
typedef vector <double> vd;
bool zero(double x) { return fabs(x) < 1e-10; }</pre>
double MinCostMatching(const vector < vd>& cost, vi& L, vi& R) ←
  int n = sz(cost), mated = 0;
  vd dist(n), u(n), v(n);
  vi dad(n), seen(n);
  rep(i,0,n) {
    u[i] = cost[i][0];
    rep(j,1,n) u[i] = min(u[i], cost[i][j]);
  rep(j,0,n) {
   v[j] = cost[0][j] - u[0];
   rep(i,1,n) v[j] = min(v[j], cost[i][j] - u[i]);
 L = R = vi(n, -1);
  rep(i,0,n) rep(j,0,n) {
    if (R[j] != -1) continue;
    if (zero(cost[i][j] - u[i] - v[j])) {
     L[i] = j;
     R[j] = i;
     mated++;
     break;
  for (; mated < n; mated++) { // until solution is feasible
    int s = 0;
    while (L[s] != -1) s++;
    fill (all (dad), -1);
    fill (all (seen), 0);
    rep(k,0,n)
     dist[k] = cost[s][k] - u[s] - v[k];
    int j = 0;
    for (;;) {
     j = -1;
      rep(k,0,n){
        if (seen[k]) continue;
        if (j == -1 \mid | \operatorname{dist}[k] < \operatorname{dist}[j]) j = k;
      seen[j] = 1;
      int i = R[j];
      if (i == -1) break;
```

```
rep(k,0,n) {
        if (seen[k]) continue;
        auto new dist = dist[j] + cost[i][k] - u[i] - v[k];
        if (dist[k] > new dist) {
           dist[k] = new dist;
           dad[k] = j;
    rep(k,0,n) {
      if (k == j || !seen[k]) continue;
      auto w = dist[k] - dist[i];
      v[k] += w, u[R[k]] -= w;
    u[s] += dist[j];
    while (dad[j] >= 0) {
      int d = dad[j];
      R[i] = R[d]:
      L[R[j]] = j;
      j = d;
    R[i] = s;
    L[s] = j;
  auto value = vd(1)[0];
  rep(i,0,n) value += cost[i][L[i]];
  return value;
GaleShaplev.h
Description: Gale-Shapley algorithm for the stable marriage problem.
madj[i][j] is the jth highest ranked woman for man i. fpref[i][j] is the rank
woman i assigns to man j. Returns a pair of vectors(mpart, fpart), where
mpart[i] gives the partner of man i, and fpart is analogous.
Time: \mathcal{O}(n^2)
                                                        ac5ecc, 26 lines
pair < vector < int >, vector < int >> stable marriage (vector < vector ←
     <int>> & madj, vector <vector <int>>& fpref) {
  int n = madj.size();
  vector <int> mpart(n, -1), fpart(n, -1);
  vector <int> midx(n);
  queue<int> mfree:
  for (int i = 0; i < n; i++) {
    mfree.push(i);
  while (!mfree.emptv()) {
    int m = mfree.front();
    mfree.pop();
    int f = madj[m][midx[m]++];
    if (fpart[f] == -1) {
      mpart[m] = f;
       fpart[f] = m;
    } else if (fpref[f][m] < fpref[f][fpart[f]]) {</pre>
      mpart[fpart[f]] = -1;
      mfree.push(fpart[f]);
      mpart[m] = f;
       fpart[f] = m;
    } else {
      mfree.push (m);
  return {mpart, fpart};
General Matching.h
Description: Matching for general graphs. Fails with probability N/mod.
Time: \mathcal{O}(N^3)
"../numerical/MatrixInverse-mod.h"
                                                        bb8be4, 40 lines
```

```
vector <pii > generalMatching(int N, vector <pii > & ed) {
 vector < vector < ll >> mat(N, vector < ll > (N)), A;
 trav(pa, ed) {
   int a = pa.first, b = pa.second, r = rand() % mod;
   mat[a][b] = r, mat[b][a] = (mod - r) \% mod;
 int r = matInv(A = mat), M = 2*N - r, fi, fj;
  assert (r % 2 == 0);
 if (M != N) do {
   mat.resize (M, vector < ll > (M));
   rep(i,0,N) {
     mat[i].resize(M);
      rep(j,N,M) {
        int r = rand() % mod;
        mat[i][i] = r, mat[i][i] = (mod - r) \% mod;
 } while (matInv(A = mat) != M);
  vi has (M, 1); vector <pii > ret;
 rep(it,0,M/2) {
   rep(i,0,M) if (has[i])
      rep(j, i+1,M) if (A[i][j] && mat[i][j]) {
        fi = i; fj = j; goto done;
    } assert(0); done:
    if (fj < N) ret.emplace back(fi, fj);</pre>
    has[fi] = has[fj] = 0;
    rep(sw,0,2) {
      ll a = modpow(A[fi][fj], mod-2);
      rep(i,0,M) if (has[i] && A[i][fj]) {
        ll b = A[i][fj] * a % mod;
        rep(j,0,M) A[i][j] = (A[i][j] - A[fi][j] * b) % mod;
      swap(fi,fj);
 return ret;
```

7.4 DFS algorithms

ArticulationPointAndBridges.h

Description: Computes dfs_low = minimum dfs number in subtree and dfs num. Note AP special case for dfs root.

```
Time: \mathcal{O}(N+E)
                                                        952683, 37 lines
void dfs(int u) {
  dfs low[u] = dfs num[u] = dfsNumberCounter++; // dfs low[u\leftarrow
       | \leq dfs \quad num[u]
  for(int j = 0; j < (int) AdjList[u].size(); j++) {
    ii v = AdjList[u][j];
    if(dfs num[v.first] == UNVISITED) { // a tree edge
      dfs parent[v.first] = u;
      if (u == dfsRoot) rootChildren++; // special case if u←
             is a root
      dfs(v.first);
      if (dfs low[v.first] >= dfs num[u]) {
        articulation vertex[u] = true;
      if (dfs low[v.first] > dfs num[u]) {
         printf(" Edge (%d, %d) is a bridge\n", u, v.first);
      dfs low[u] = min(dfs low[u], dfs low[v.first]); // \leftarrow
           update dfs low[\overline{u}]
```

```
else if(v.first != dfs parent[u]) // a back edge and not←
          direct cycle
      dfs low[u] = min(dfs low[u], dfs num[v.first]); // ←
           update dfs low[u]
}
// inside int main()
dfsNumberCounter = 0;
dfs num.assign(V, UNVISITED);
dfs low.assign(V, 0);
dfs parent.assign(V, 0);
articulation vertex.assign(V, 0);
for(int i = 0; i < V; i++) {
  if (dfs num[i] == UNVISITED) {
    dfsRoot = i;
    rootChildren = 0;
    dfs(i);
    articulation vertex[dfsRoot] = (rootChildren > 1); // ←
         special case
}
```

2sat.h

Description: Calculates a valid assignment to boolean variables a, b, c,... to a 2-SAT problem, so that an expression of the type (a|||b)&&(!a|||c)&&(d|||!b)&&... becomes true, or reports that it is unsatisfiable. Negated variables are represented by bit-inversions $(\sim x)$.

Usage: TwoSat ts(number of boolean variables); ts.either(0, ~3); // Var 0 is true or var 3 is false ts.set_value(2); // Var 2 is true ts.at_most_one($\{0, \sim 1, 2\}$); // <= 1 of vars 0, ~ 1 and 2 are true ts.solve(); // Returns true iff it is solvable ts.values[0..N-1] holds the assigned values to the vars

Time: $\mathcal{O}\left(N+E\right)$, where N is the number of boolean variables, and E is the number of clauses.

```
struct TwoSat {
  int N;
  vector <vi> gr;
  vi values; // 0 = false, 1 = true
 TwoSat(int n = 0) : N(n), gr(2*n) {}
  int add var() { // (optional)
    gr.emplace back();
    gr.emplace back();
   return N++;
  void either(int f, int j) {
    f = \max(2*f, -1-2*f);
   j = max(2*j, -1-2*j);
   gr[f].push back(j^1);
   gr[j].push back(f^1);
  void set value(int x) { either(x, x); }
  void at most one(const vi& li) { // (optional)
    if (sz(li) <= 1) return;
    int cur = \simli[0];
    rep(i,2,sz(li)) {
     int next = add var();
      either (cur, ~li[i]);
      either (cur, next);
      either (~li[i], next);
      cur = \simnext;
    either (cur, ~li[1]);
```

```
vi val, comp, z; int time = 0;
 int dfs(int i) {
    int low = val[i] = ++time, x; z.push back(i);
    trav(e, gr[i]) if (!comp[e])
     low = min(low, val[e] ?: dfs(e));
    if (low == val[i]) do {
     x = z.back(); z.pop back();
     comp[x] = low;
      if (values[x>>1] == -1)
        values[x>>1] = x&1;
   } while (x != i);
    return val[i] = low;
 bool solve() {
    values.assign(N, -1);
    val.assign(2*N, 0); comp = val;
    rep(i,0,2*N) if (!comp[i]) dfs(i);
   rep(i,0,N) if (comp[2*i] == comp[2*i+1]) return 0;
    return 1:
};
```

7.5 Heuristics

MaximalCliques.h

Description: Runs a callback for all maximal cliques in a graph (given as a symmetric bitset matrix; self-edges not allowed). Callback is given a bitset representing the maximal clique.

```
Time: \mathcal{O}\left(3^{n/3}\right), much faster for sparse graphs
```

b0d5b1, 12 lines

MaximumClique.h

Description: Quickly finds a maximum clique of a graph (given as symmetric bitset matrix; self-edges not allowed). Can be used to find a maximum independent set by finding a clique of the complement graph.

Time: Runs in about 1s for n=155 and worst case random graphs (p=.90). Runs faster for sparse graphs.

```
fbbef1, 49 lines
typedef vector <br/>
vector <br/>
vitset <200>> vb;
struct Maxclique {
 double limit=0.025, pk=0;
 struct Vertex { int i, d=0; };
 typedef vector (Vertex) vv;
 vb e;
 vv V;
 vector <vi>C;
 vi qmax, q, S, old;
 void init(vv& r) {
   trav(v,r) v.d = 0;
    trav(v, r) trav(j, r) v.d += e[v.i][j.i];
    sort(all(r), [](auto a, auto b) { return a.d > b.d; });
    int mxD = r[0].d;
    rep(i,0,sz(r)) r[i].d = min(i, mxD) + 1;
```

```
void expand(vv& R, int lev = 1) {
    S[lev] += S[lev - 1] - old[lev];
    old[lev] = S[lev - 1];
    while (sz(R)) {
      if (sz(q) + R.back().d <= sz(qmax)) return;
      q.push back(R.back().i);
      vv T;
      trav(v,R) if (e[R.back().i][v.i]) T.push back({v.i});
      if (sz(T)) {
        if (S[lev]++ / ++pk < limit) init(T);
        int j = 0, mxk = 1, mnk = max(sz(qmax) - sz(q) + 1, \leftarrow
        C[1].clear(), C[2].clear();
        trav(v, T) {
          int k = 1;
          auto f = [&](int i) { return e[v.i][i]; };
          while (any of(all(C[k]), f)) k++;
          if (k > mx\bar{k}) mxk = k, C[mxk + 1].clear();
          if (k < mnk) T[j++] \cdot i = v \cdot i;
          C[k].push back(v.i);
        if (j > 0) T[j - 1] \cdot d = 0;
        rep(k,mnk,mxk + 1) trav(i, C[k])
          T[j] \cdot i = i, T[j++] \cdot d = k;
        expand (T, lev + 1);
      } else if (sz(q) > sz(qmax)) qmax = q;
      q.pop back(), R.pop back();
 vi maxClique() { init(V), expand(V); return qmax; }
 Maxclique(vb conn): e(conn), C(sz(e)+1), S(sz(C)), old(S) \leftarrow
    rep(i,0,sz(e)) V.push_back({i});
};
```

7.6 Trees

CompressTree.h

Description: Given a rooted tree and a subset S of nodes, compute the minimal subtree that contains all the nodes by adding all (at most |S|-1) pairwise LCA's and compressing edges. Returns a list of (par, orig_index) representing a tree rooted at 0. The root points to itself.

```
Time: \mathcal{O}\left(|S|\log|S|\right)
```

```
dabd75, 20 lines
vpi compressTree(LCA& lca, const vi& subset) {
 static vi rev: rev.resize(sz(lca.dist));
 vi li = subset, &T = lca.time;
 auto cmp = [&](int a, int b) { return T[a] < T[b]; };</pre>
  sort(all(li), cmp);
 int m = sz(li)-1;
 rep(i,0,m) {
   int a = li[i], b = li[i+1];
   li.push back(lca.query(a, b));
 sort(all(li), cmp);
 li.erase(unique(all(li)), li.end());
 rep(i,0,sz(li)) rev[li[i]] = i;
 vpi ret = {pii(0, li[0])};
 rep(i,0,sz(li)-1) {
   int a = li[i], b = li[i+1];
   ret.emplace back(rev[lca.query(a, b)], b);
 return ret;
```

LinkCutTree.h

Description: Represents a forest of unrooted trees. You can add and remove edges (as long as the result is still a forest), and check whether two nodes are in the same tree.

Time: All operations take amortized $\mathcal{O}(\log N)$.

693483, 90 lines

```
struct Node { // Splay tree. Root's pp contains tree's ←
  Node *p = 0, *pp = 0, *c[2];
  bool flip = 0;
  Node() { c[0] = c[1] = 0; fix(); }
  void fix() {
    if (c[0]) c[0] - p = this;
    if (c[1]) c[1] \rightarrow p = this;
    // (+ update sum of subtree elements etc. if wanted)
  void push flip() {
    if (!flip) return;
    flip = 0; swap(c[0], c[1]);
    if (c[0]) c[0]->flip ^= 1;
    if (c[1]) c[1]->flip ^= 1;
  int up() { return p ? p->c[1] == this : -1; }
  void rot(int i, int b) {
    int h = i ^ b:
    Node \star x = c[i], \star y = b == 2 ? x : x \rightarrow c[h], \star z = b ? y : \leftarrow
    if ((y-p = p)) p-c[up()] = y;
    c[i] = z -> c[i ^ 1];
    if (b < 2) {
      x\rightarrow c[h] = y\rightarrow c[h ^ 1];
      z \rightarrow c[h ^1] = b ? x : this;
    y - c[i ^1] = b ? this : x;
    fix(); x\rightarrow fix(); y\rightarrow fix();
    if (p) p->fix();
    swap(pp, y \rightarrow pp);
  void splay() {
    for (push flip(); p; ) {
      if (p\rightarrow p) p\rightarrow p\rightarrow push flip();
      p->push flip(); push_flip();
      int c1 = up(), c2 = p \rightarrow up();
      if (c2 == -1) p \rightarrow rot(c1, 2);
      else p->p->rot(c2, c1 != c2);
  Node* first() {
    return c[0] ? c[0]->first() : (splay(), this);
};
struct LinkCut {
  vector < Node > node:
  LinkCut(int N) : node(N) {}
  void link(int u, int v) \{ // add an edge (u, v) \}
    assert (!connected(u, v));
    make root(&node[u]);
    node[u].pp = &node[v];
  void cut(int u, int v) { // remove an edge (u, v)
    Node *x = &node[u], *top = &node[v];
    make root(top); x->splay();
    assert (top == (x-pp ?: x-c[0]));
    if (x-pp) x-pp = 0;
    else {
      x - c[0] = top - p = 0;
```

```
x\rightarrow fix();
  bool connected(int u, int v) { // are u, v in the same ←
    Node* nu = access(&node[u])->first();
     return nu == access(&node[v])->first();
  void make root(Node* u) {
     access(u);
    u->splay();
    if(u->c[0]) {
       u - c[0] - p = 0;
       u->c[0]->flip ^= 1;
       u - c[0] - pp = u;
       u \rightarrow c[0] = 0;
       u->fix();
  Node* access(Node* u) {
    u->splay();
    while (Node* pp = u-pp) {
       pp \rightarrow splay(); u \rightarrow pp = 0;
       if (pp->c[1]) {
         pp \rightarrow c[1] \rightarrow p = 0; pp \rightarrow c[1] \rightarrow pp = pp; 
       pp \rightarrow c[1] = u; pp \rightarrow fix(); u = pp;
    return u;
};
DirectedMST.h.
Description: Edmonds' algorithm for finding the weight of the minimum
spanning tree/arborescence of a directed graph, given a root node. If no
MST exists, returns -1.
Time: \mathcal{O}\left(E\log V\right)
"../data-structures/UnionFind.h"
struct Edge { int a, b; ll w; };
struct Node {
  Edge kev:
  Node *1, *r;
  ll delta;
  void prop() {
    kev.w += delta:
    if (1) 1->delta += delta;
    if (r) r->delta += delta;
     delta = 0;
  Edge top() { prop(); return key; }
Node *merge(Node *a, Node *b) {
  if (!a || !b) return a ?: b;
  a->prop(), b->prop();
  if (a\rightarrow key.w \rightarrow b\rightarrow key.w) swap(a, b);
  swap(a\rightarrow l, (a\rightarrow r = merge(b, a\rightarrow r)));
  return a;
void pop(Node*& a) { a \rightarrow prop(); a = merge(a \rightarrow l, a \rightarrow r); }
ll dmst(int n, int r, vector < Edge > & g) {
 UF uf(n);
  vector < Node*> heap(n);
  trav(e, g) heap[e.b] = merge(heap[e.b], new Node{e});
  ll res = 0;
  vi seen(n, -1), path(n);
  seen[r] = r;
  rep(s,0,n) {
    int u = s, qi = 0, w;
```

while (seen[u] < 0) {

```
path[qi++] = u, seen[u] = s;
    if (!heap[u]) return -1;
    Edge e = heap[u] \rightarrow top();
    heap[u]->delta -= e.w, pop(heap[u]);
    res += e.w, u = uf.find(e.a);
    if (seen[u] == s) {
      Node* cvc = 0;
      do cyc = merge(cyc, heap[w = path[--qi]]);
      while (uf.join(u, w));
      u = uf.find(u);
      heap[u] = cyc, seen[u] = -1;
 }
return res;
```

Math-0

7.7.1 Number of Spanning Trees

Create an $N \times N$ matrix mat, and for each edge $a \to b \in G$. do mat[a][b]-, mat[b][b]++ (and mat[b][a]-, $\mathsf{mat}[\mathsf{a}][\mathsf{a}] + \mathsf{if} G$ is undirected). Remove the *i*th row and column and take the determinant; this yields the number of directed spanning trees rooted at i (if G is undirected, remove any row/column).

7.7.2 Erdős–Gallai theorem

A simple graph with node degrees $d_1 \geq \cdots \geq d_n$ exists iff $d_1 + \cdots + d_n$ is even and for every $k = 1 \dots n$,

$$\sum_{i=1}^{k} d_i \le k(k-1) + \sum_{i=k+1}^{n} \min(d_i, k).$$

7.7.3 Konig's theorem

In any bipartite graph, the number of edges in a maximum mathing equals the number of vertices in a minimum vertex cover. To exhibit the vertex cover:

- 1. Find a maximum matching.
- 2. Change each edge **used** in the matching into a directed edge from right to left.
- 3. Change each edge **not used** in the matching into a directed edge from left to right.
- 4. Compute the set T of all vertices reachable from unmatched vertices on the left (including themselves).
- 5. The vertex cover consists of all vertices on the right that are in T, and all vertices on the left that are not in T.

Math-2

7.8.1 Minumum Edge cover

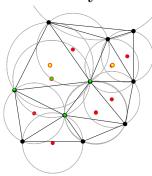
If a minimum edge cover contains C edges, and a maximum matching contains M edges, then C + M = |V|. To obtain the edge cover, start with a maximum matching, and then, for every vertex not matched, just select some edge incident upon it and add it to the edge set.

7.8.2 Maximum Independent set

To obtain a maximum independent set of a graph, find a max clique of the complement. If the graph is bipartite, see MinimumVertexCover.

Geometry (8)

8.1 Voronoi - Delanauy dual



The Delaunay triangulation with all the circumcircles and their centers (in red).



Connecting the centers of the circumcircles produces the Voronoi diagram (in red).

Green highlighted points show that circumcenter can be outside triangle. Yellow highlighted points show that Voronoi cell edge is present only triangles have shared edge, not shared vertex.

8.2 Geometric primitives

if (s==e) return (p-s).dist();

return ((p-s)*d-(e-s)*t).dist()/d;

auto $d = (e-s) \cdot dist2()$, $t = min(d, max(.0, (p-s) \cdot dot(e-s)))$;

```
Point.h
Description: Class to handle points in the plane. T can be e.g. double or
long long. (Avoid int.)
template \langle class T \rangle int sgn(T x) \{ return (x > 0) - (x < 0); \leftarrow
template<class T>
struct Point {
 typedef Point P;
 Тх, у;
  explicit Point (T x=0, T y=0): x(x), y(y) {}
  bool operator<(P p) const { return tie(x,y) < tie(p.x,p.y) \leftarrow
  bool operator==(P p) const { return tie(x,y)==tie(p.x,p.y)} \leftarrow
  P operator+(P p) const { return P(x+p.x, y+p.y); }
 P operator-(P p) const { return P(x-p.x, y-p.y); }
 P operator*(T d) const { return P(x*d, y*d); }
 P operator/(T d) const { return P(x/d, y/d); }
 T dot(P p) const { return x*p.x + y*p.y; }
 T cross(P p) const { return x*p.v - y*p.x; }
 T cross(P a, P b) const { return (a-*this).cross(b-*this);←
 T dist2() const { return x*x + y*y; }
  double dist() const { return sart((double)dist2()); }
  // angle to x-axis in interval [-pi, pi]
  double angle() const { return atan2(y, x); }
 P unit() const { return *this/dist(); } // makes dist()=1
 P perp() const { return P(-y, x); } // rotates +90 degrees
 P normal() const { return perp().unit(); }
  // returns point rotated 'a' radians ccw around the origin
 P rotate (double a) const {
    return P(x*\cos(a)-y*\sin(a),x*\sin(a)+y*\cos(a)); }
  friend ostream& operator<<(ostream& os, P p) {
    return os << "(" << p.x << "," << p.y << ")"; }
lineDistance.h
Description:
Returns the signed distance between point p and the line con-
taining points a and b. Positive value on left side and negative
on right as seen from a towards b. a==b gives nan. P is sup-
posed to be Point<T> or Point3D<T> where T is e.g. double
or long long. It uses products in intermediate steps so watch
out for overflow if using int or long long. Using Point3D will
always give a non-negative distance. For Point3D, call .dist /S
on the result of the cross product.
                                                         f6bf6b, 4 lines
template<class P>
double lineDist(const P& a, const P& b, const P& p) {
 return (double)(b-a).cross(p-a)/(b-a).dist();
SegmentDistance.h
Description:
Returns the shortest distance between point p and the line
segment from point s to e.
Usage: Point<double> a, b(2,2), p(1,1);
bool onSegment = segDist(a,b,p) < 1e-10;</pre>
                                                         5c88f4, 6 lines
typedef Point <double> P;
double segDist(P& s, P& e, P& p) {
```

```
SegmentIntersection.h
```

Description:

If a unique intersection point between the line segments going from s1 to e1 and from s2 to e2 exists then it is returned. If no intersection point exists an empty vector is returned. If infinitely many exist a vector with 2 elements is returned, containing the endpoints of the common line segment. The wrong position will be returned if P is Point<|l> and the intersection point does not have integer coordinates. Products of three coordinates are used in intermediate steps so watch out for overflow if using int or long long. Usage: vector<P> inter = segInter(s1,e1,s2,e2);



```
if (sz(inter)==1)
cout << "segments intersect at " << inter[0] << endl;</pre>
"Point.h", "OnSegment.h"
                                                      9d57f2, 13 lines
template < class P> vector < P> segInter (P a, P b, P c, P d) {
  auto oa = c.cross(d, a), ob = c.cross(d, b),
       oc = a.cross(b, c), od = a.cross(b, d);
  // Checks if intersection is single non-endpoint point.
 if (sgn(oa) * sgn(ob) < 0 && sgn(oc) * sgn(od) < 0)
   return \{(a * ob - b * oa) / (ob - oa)\};
  set <P> s;
 if (onSegment(c, d, a)) s.insert(a);
 if (onSegment(c, d, b)) s.insert(b);
 if (onSegment(a, b, c)) s.insert(c);
 if (onSegment(a, b, d)) s.insert(d);
 return {all(s)};
```

lineIntersection.h

Description:

If a unique intersection point of the lines going through s1,e1 and s2,e2 exists {1, point} is returned. If no intersection point exists $\{0, (0,0)\}\$ is returned and if infinitely many exists $\{-1,$ (0,0)} is returned. The wrong position will be returned if P is Point<|l> and the intersection point does not have integer coordinates. Products of three coordinates are used in intermediate steps so watch out for overflow if using int or ll. Does not always return true if the intersection is at the boundary point(s). Usage: auto res = lineInter(s1,e1,s2,e2);



3af81c, 9 lines

```
if (res.first == 1)
cout << "intersection point at " << res.second << endl;</pre>
template<class P>
pair <int, P> lineInter (P s1, P e1, P s2, P e2) {
  auto d = (e1 - s1).cross(e2 - s2);
  if (d == 0) // if parallel
    return \{-(s1.cross(e1, s2) == 0), P(0, 0)\};
  auto p = s2.cross(e1, e2), q = s2.cross(e2, s1);
  return \{1, (s1 * p + e1 * q) / d\};
```

sideOf.h

Description: Returns where p is as seen from s towards e. $1/0/-1 \Leftrightarrow \text{left/on}$ line/right. If the optional argument eps is given 0 is returned if p is within distance eps from the line. P is supposed to be Point<T> where T is e.g. double or long long. It uses products in intermediate steps so watch out for overflow if using int or long long.

```
Usage: bool left = sideOf(p1,p2,q)==1;
"Point.h"
```

```
template<class P>
int sideOf(P s, P e, P p) { return sgn(s.cross(e, p)); }
template<class P>
int sideOf(const P& s, const P& e, const P& p, double eps) {
  auto a = (e-s) \cdot cross(p-s);
 double l = (e-s) \cdot dist() *eps;
 return (a > l) - (a < -l);
```

OnSegment.h

Description: Returns true iff p lies on the line segment from s to e. Use (segDist(s,e,p)<=epsilon) instead when using Point<double>.

c597e8, 3 lines template < class P> bool on Segment (P s, P e, P p) { return p.cross(s, e) == 0 && (s - p). $dot(e - p) \le 0$;

linearTransformation.h Description:

scaling) which takes line p0-p1 to line q0-q1 to point r.

Apply the linear transformation (translation, rotation and 03a306, 6 lines

```
typedef Point <double> P;
P linearTransformation(const P& p0, const P& p1,
    const P& q0, const P& q1, const P& r) {
  P dp = p1-p0, dq = q1-q0, num(dp.cross(dq), dp.dot(dq));
  return q0 + P((r-p0).cross(num), (r-p0).dot(num))/dp.dist2
```

LineProjectionReflection.h

Description: Projects point p onto line ab. Set refl=true to get reflection of point p across line ab insted. The wrong point will be returned if P is an integer point and the desired point doesn't have integer coordinates. Products of three coordinates are used in intermediate steps so watch out for overflow. "Point.h" b5562d, 5 lines

```
template<class P>
P lineProj (P a, P b, P p, bool refl=false) {
 P v = b - a;
  return p - v.perp()*(1+refl)*v.cross(p-a)/v.dist2();
```

Angle.h

between

Description: A class for ordering angles (as represented by int points and a number of rotations around the origin). Useful for rotational sweeping. Sometimes also represents points or vectors.

Usage: vector<Angle $> v = \{w[0], w[0].t360() ...\}; // sorted$ int j = 0; rep(i,0,n) { while (v[j] < v[i].t180()) ++j; } // sweeps j such that (j-i) represents the number of positively oriented triangles with vertices at 0 and i 0f0602, 35 lines

```
struct Angle {
  int x, y;
  int t;
  Angle(int x, int y, int t=0): x(x), y(y), t(t) {}
  Angle operator-(Angle b) const { return \{x-b.x, y-b.y, t\}; \leftarrow
  int half() const {
    assert(x | | y);
    return y < 0 \mid | (y == 0 && x < 0);
  Angle t90() const { return \{-y, x, t + (half() \&\& x \ge 0) \leftarrow \}
      }; }
  Angle t180() const { return {-x, -y, t + half()}; }
 Angle t360() const { return {x, y, t + 1}; }
bool operator<(Angle a, Angle b) {</pre>
  // add a.dist2() and b.dist2() to also compare distances
  return make tuple(a.t, a.half(), a.y * (ll)b.x) <
         make tuple(b.t, b.half(), a.x * (ll)b.y);
// Given two points, this calculates the smallest angle ←
```

```
// them, i.e., the angle that covers the defined line \
pair < Angle, Angle > segment Angles (Angle a, Angle b) {
 if (b < a) swap(a, b);
 return (b < a.t180() ?
         make pair(a, b): make pair(b, a.t360()));
Angle operator+(Angle a, Angle b) { // point a + vector b
 Angle r(a.x + b.x, a.y + b.y, a.t);
 if (a.t180() < r) r.t--;
 return r.t180() < a ? r.t360() : r;
Angle angleDiff(Angle a, Angle b) { // angle b - angle a
 int tu = b.t - a.t; a.t = b.t;
 return \{a.x*b.x + a.y*b.y, a.x*b.y - a.y*b.x, tu - (b < a) \leftarrow
```

Circles

CircleIntersection.h

Description: Computes the pair of points at which two circles intersect. Returns false in case of no intersection.

```
84d6d3, 11 lines
typedef Point <double> P;
bool circleInter (P a, P b, double r1, double r2, pair <P, P>* out←
  if (a == b) { assert(r1 != r2); return false; }
 P \text{ vec} = b - a;
  double d2 = \text{vec.dist2}(), sum = r1+r2, dif = r1-r2,
          p = (d2 + r1*r1 - r2*r2)/(d2*2), h2 = r1*r1 - p*p*\leftarrow
  if (sum*sum < d2 || dif*dif > d2) return false;
  P mid = a + vec*p, per = vec.perp() * sqrt(fmax(0, h2) / \leftarrow
  \star \text{out} = \{ \text{mid} + \text{per}, \text{mid} - \text{per} \};
  return true;
```

CircleTangents.h

Description: Finds the external tangents of two circles, or internal if r2 is negated. Can return 0, 1, or 2 tangents – 0 if one circle contains the other (or overlaps it, in the internal case, or if the circles are the same); 1 if the circles are tangent to each other (in which case .first = .second and the tangent line is perpendicular to the line between the centers). .first and .second give the tangency points at circle 1 and 2 respectively. To find the tangents of a circle with a point set r2 to 0.

```
b0153<u>d</u>, 13 lines
"Point.h"
template<class P>
vector <pair <P, P>> tangents (P c1, double r1, P c2, double r2 ←
 P d = c2 - c1;
 double dr = r1 - r2, d2 = d \cdot dist2(), h2 = d2 - dr * dr;
 if (d2 == 0 | | h2 < 0) return {};
 vector <pair <P, P>> out;
 for (double sign : {-1, 1}) {
   P v = (d * dr + d \cdot perp() * sqrt(h2) * sign) / d2;
    out.push back(\{c1 + v * r1, c2 + v * r2\});
 if (h2 == 0) out.pop back();
 return out:
```

CircleLine.h

Description: Finds the intersection between a circle and a line. Returns a vector of either 0, 1, or 2 intersection points. P is intended to be Point<double>

```
"Point.h", "lineDistance.h", "LineProjectionReflection.h"
```

debf86, 8 lines

```
template<class P>
vector <P> circleLine (P c, double r, P a, P b) {
 double h2 = r*r - a.cross(b,c)*a.cross(b,c)/(b-a).dist2();
 if (h2 < 0) return {};
 P p = lineProj(a, b, c), h = (b-a).unit() * sqrt(h2);
 if (h2 == 0) return {p};
 return {p - h, p + h};
```

CirclePolygonIntersection.h

Description: Returns the area of the intersection of a circle with a ccw polygon.

```
Time: \mathcal{O}(n)
```

"../../content/geometry/Point.h"

```
typedef Point <double> P;
#define arg(p, q) atan2(p.cross(q), p.dot(q))
double circlePoly(P c, double r, vector <P> ps) {
  auto tri = [&](P p, P q) {
    auto r2 = r * r / 2;
    Pd = q - p;
    auto a = d \cdot dot(p)/d \cdot dist2(), b = (p \cdot dist2()-r*r)/d \cdot dist2 \leftarrow
         ();
    auto det = a * a - b;
    if (det \le 0) return arg(p, q) * r2;
    auto s = \max(0., -a-\operatorname{sqrt}(\det)), t = \min(1., -a+\operatorname{sqrt}(\det))
    if (t < 0 | | 1 \le s) return arg(p, q) * r2;
    Pu = p + d * s, v = p + d * t;
    return arg(p,u) * r2 + u.cross(v)/2 + arg(v,q) * r2;
  auto sum = 0.0;
  rep(i,0,sz(ps))
    sum += tri(ps[i] - c, ps[(i + 1) % sz(ps)] - c);
  return sum;
```

circumcircle.h Description:

"Point.h"

The circumcirle of a triangle is the circle intersecting all three vertices. ccRadius returns the radius of the circle going through points A. B and C and ccCenter returns the center of the same circle.



1caa3a, 9 lines

```
typedef Point <double> P;
double ccRadius(const P& A, const P& B, const P& C) {
 return (B-A).dist()*(C-B).dist()*(A-C).dist()/
      abs((B-A).cross(C-A))/2;
P ccCenter (const P& A, const P& B, const P& C) {
 P b = C-A, c = B-A;
 return A + (b*c.dist2()-c*b.dist2()).perp()/b.cross(c)/2;
```

MinimumEnclosingCircle.h

Description: Computes the minimum circle that encloses a set of points. **Time:** expected $\mathcal{O}(n)$

```
"circumcircle.h"
                                                      09dd0a, 17 lines
pair <P, double> mec(vector <P> ps) {
 shuffle(all(ps), mt19937(time(0)));
 P \circ = ps[0];
 double r = 0, EPS = 1 + 1e-8;
 rep(i,0,sz(ps)) if ((o - ps[i]).dist() > r * EPS) {
    o = ps[i], r = 0;
    rep(j,0,i) if ((o - ps[j]).dist() > r * EPS) {
      o = (ps[i] + ps[j]) / 2;
      r = (o - ps[i]) \cdot dist();
      rep(k,0,j) if ((o - ps[k]).dist() > r * EPS) {
```

```
return {o, r};
kIntersection.h
Description: Given n circles, for all k <= n, computes the area of regions
part of atleast k circles.
Time: \mathcal{O}\left(n^2\right)
const int N = 22222;
const double EPS = 1e-8:
const double PI = acos(-1.0);
typedef complex <double> Point;
double r[N], result[N];
Point c[N];
pair <double, int> event[N];
int sgn (double x) {return x < -EPS? -1: x < EPS? 0: 1;}
double det (const Point &a, const Point &b) { return a.real ←
     () * b.imag() - a.imag()
  * b.real();}
  void addEvent (double a, int v) {
    event[m ++] = make pair(a, v);
void addPair (double a, double b) {
  if (sgn(a - b) <= 0) {
    addEvent(a, +1);
    addEvent(b, -1);
  } else {
    addPair(a, +PI);
    addPair(-PI, b);
Point polar (double t) { return Point(cos(t), sin(t)); }
Point radius (int i, double t) {
  return c[i] + polar(t) * r[i];
void solve () {
  // result[k]: the total area covered no less than k times
  memset(result, 0, sizeof(result));
  for (int i = 0; i < n; ++ i) {
   m = 0:
    addEvent(-PI, 0);
    addEvent(+PI, 0);
    for (int j = 0; j < n; ++ j) {
      if (i != j) {
        if (\operatorname{sgn}(\operatorname{abs}(c[i] - c[j]) - \operatorname{abs}(r[i] - r[j])) \le 0)
          if (sgn(r[i] - r[j]) <= 0) {
             addPair(-PI, +PI);
        } else {
          if (sgn(abs(c[i] - c[j]) - (r[i] + r[j])) >= 0) {
            continue;
          double d = abs(c[j] - c[i]);
          Point b = (c[j] - c[i]) / d * r[i];
          double t = acos((r[i] * r[i] + d * d - r[j] * r[j \leftarrow
               ]) / (2 *
                 r[i] * d));
           Point a = b * polar(-t);
          Point c = b * polar(+t);
          addPair(arg(a), arg(c));
    sort (event, event + m);
```

o = ccCenter(ps[i], ps[j], ps[k]);

r = (o - ps[i]).dist();

8.4 Polygons

InsidePolygon.h

Description: Returns true if p lies within the polygon. If strict is true, it returns false for points on the boundary. The algorithm uses products in intermediate steps so watch out for overflow.

```
Usage: vector P> v = P\{4,4\}, P\{1,2\}, P\{2,1\}\}; bool in = inPolygon(v, P{3, 3}, false);
Time: O(n)
```

"Point.h", "OnSegment.h", "SegmentDistance.h" 2bf504, 11 lines

```
template < class P>
bool inPolygon(vector < P> & p, P a, bool strict = true) {
  int cnt = 0, n = sz(p);
  rep(i,0,n) {
    P q = p[(i + 1) % n];
    if (onSegment(p[i], q, a)) return ! strict;
    //or: if (segDist(p[i], q, a) <= eps) return ! strict;
    cnt ^= ((a.y < p[i].y) - (a.y < q.y)) * a.cross(p[i], q) > \leftarrow
    0;
  }
  return cnt;
```

PolygonArea.h

Description: Returns twice the signed area of a polygon. Clockwise enumeration gives negative area. Watch out for overflow if using int as T!

"Point.h"

f12300, 6 lines

```
template<class T>
T polygonArea2(vector<Point<T>& v) {
  T a = v.back().cross(v[0]);
  rep(i,0,sz(v)-1) a += v[i].cross(v[i+1]);
  return a;
}
```

PolygonCenter.h

Description: Returns the center of mass for a polygon.

Time: $\mathcal{O}\left(n\right)$

```
typedef Point<double> P;
P polygonCenter(const vector<P>& v) {
   P res(0, 0); double A = 0;
   for (int i = 0, j = sz(v) - 1; i < sz(v); j = i++) {
      res = res + (v[i] + v[j]) * v[j].cross(v[i]);
      A += v[j].cross(v[i]);
}
return res / A / 3;</pre>
```

PolygonCut.h Description:

Returns a vector with the vertices of a polygon with everything to the left of the line going from s to e cut away.

```
Usage: vector<P> p = ...;
p = polygonCut(p, P(0,0), P(1,0));
"Point.h", "lineIntersection.h"
```



9706dc, 9 lines

```
typedef Point<double> P;
vector <P> polygonCut(const vector <P>& poly, P s, P e) {
  vector <P> res;
  rep(i,0,sz(poly)) {
    P cur = poly[i], prev = i ? poly[i-1] : poly.back();
    bool side = s.cross(e, cur) < 0;
    if (side != (s.cross(e, prev) < 0))
      res.push_back(lineInter(s, e, cur, prev).second);
    if (side)
      res.push_back(cur);
  }
  return res;
}</pre>
```

ConvexHull.h

Description:

Returns a vector of the points of the convex hull in counterclockwise order. Points on the edge of the hull between two other points are not considered part of the hull. **Time:** $\mathcal{O}(n \log n)$



```
"Point.h" 26a0a9, 13 lines
```

```
typedef Point<ll> P;
vector <P> convexHull(vector <P> pts) {
   if (sz(pts) <= 1) return pts;
   sort(all(pts));
   vector <P> h(sz(pts)+1);
   int s = 0, t = 0;
   for (int it = 2; it--; s = --t, reverse(all(pts)))
      trav(p, pts) {
      while (t >= s + 2 && h[t-2].cross(h[t-1], p) <= 0) t ←
      --;
      h[t++] = p;
   }
   return {h.begin(), h.begin() + t - (t == 2 && h[0] == h ←
      [1])};
}</pre>
```

HullDiameter.h

Description: Returns the two points with max distance on a convex hull (ccw, no duplicate/colinear points).

```
typedef Point<ll> P;
array<P, 2> hullDiameter(vector<P> S) {
  int n = sz(S), j = n < 2 ? 0 : 1;
  pair<ll, array<P, 2>> res({0, {S[0], S[0]}});
  rep(i,0,j)
  for (;; j = (j + 1) % n) {
    res = max(res, {(S[i] - S[j]).dist2(), {S[i], S[j]}});
    if ((S[(j + 1) % n] - S[j]).cross(S[i + 1] - S[i]) >= 
      0)
      break;
  }
  return res.second;
}
```

PointInsideHull.h

Description: Determine whether a point t lies inside a convex hull (CCW order, with no colinear points). Returns true if point lies within the hull. If strict is true, points on the boundary aren't included.

Time: $\mathcal{O}(\log N)$

```
"Point.h", "sideOf.h", "OnSegment.h" 71446b, 14 lines

typedef Point<ll> P;
```

```
return false;
while (abs(a - b) > 1) {
  int c = (a + b) / 2;
  (sideOf(l[0], l[c], p) > 0 ? b : a) = c;
}
return sgn(l[a].cross(l[b], p)) < r;</pre>
```

LineHullIntersection.h

Description: Line-convex polygon intersection. The polygon must be ccw and have no colinear points. lineHull(line, poly) returns a pair describing the intersection of a line with the polygon: \bullet (-1,-1) if no collision, \bullet (i,-1) if touching the corner i, \bullet (i,i) if along side $(i,i+1), \bullet$ (i,j) if crossing sides (i,i+1) and (j,j+1). In the last case, if a corner i is crossed, this is treated as happening on side (i,i+1). The points are returned in the same order as the line hits the polygon. extrVertex returns the point of a hull with the max projection onto a line.

```
Time: \mathcal{O}\left(N + Q \log n\right)
```

```
"Point.h"
                                                     758f22, 39 lines
typedef array <P, 2> Line;
#define cmp(i,j) sgn(dir.perp().cross(poly[(i)%n]-poly[(j)%n←
#define extr(i) cmp(i + 1, i) >= 0 && cmp(i, i - 1 + n) < 0
int extrVertex(vector < P>& poly, P dir) {
  int n = sz(poly), lo = 0, hi = n;
  if (extr(0)) return 0:
  while (lo + 1 < hi) {</pre>
    int m = (lo + hi) / 2;
    if (extr(m)) return m;
    int ls = cmp(lo + 1, lo), ms = cmp(m + 1, m);
    (ls < ms | | (ls == ms && ls == cmp(lo, m)) ? hi : lo) = ←
  return lo:
#define cmpL(i) sgn(line[0].cross(poly[i], line[1]))
array <int, 2> line Hull (Line line, vector <P> poly) {
  int endA = extrVertex(poly, (line[0] - line[1]).perp());
  int endB = extrVertex(poly, (line[1] - line[0]).perp());
  if (cmpL(endA) < 0 || cmpL(endB) > 0)
    return {-1, -1};
  array<int, 2> res;
  rep(i,0,2) {
    int lo = endB, hi = endA, n = sz(poly);
    while ((lo + 1) % n != hi) {
     int m = ((lo + hi + (lo < hi ? 0 : n)) / 2) % n;
      (cmpL(m) == cmpL(endB) ? lo : hi) = m;
    res[i] = (lo + !cmpL(hi)) % n;
   swap (endA, endB);
  if (res[0] == res[1]) return \{res[0], -1\};
  if (!cmpL(res[0]) && !cmpL(res[1]))
    switch ((res[0] - res[1] + sz(poly) + 1) % sz(poly)) {
      case 0: return {res[0], res[0]};
      case 2: return {res[1], res[1]};
  return res;
```

8.5 Misc. Point Set Problems

ManhattanMST.h

Description: Given N points, returns up to 4*N edges, which are guaranteed to contain a minimum spanning tree for the graph with edge weights w(p, q) = |p.x - q.x| + |p.y - q.y|. Edges are in the form (distance, src, dst). Use a standard MST algorithm on the result to find the final MST.

```
Time: \mathcal{O}(NlogN)
                                                     995288, 24 lines
typedef Point < int > P:
vector <array <int, 3>> manhattan MST (vector <P> ps) {
  vi id(sz(ps));
  iota(all(id), 0);
  vector <array <int, 3>> edges;
  rep(k,0,4) {
    sort(all(id), [&](int i, int j) {
         return (ps[i]-ps[j]).x < (ps[j]-ps[i]).y;);
    map<int, int> sweep:
    trav(i,id) {
      for (auto it = sweep.lower bound(-ps[i].y);
                it != sweep.end(); sweep.erase(it++)) {
        int j = it ->second;
        P d = ps[i] - ps[j];
        if (d.y > d.x) break;
        edges.push back({d.y + d.x, i, j});
      sweep[-ps[i].y] = i;
    if (k & 1) trav(p,ps) p.x = -p.x;
    else trav(p,ps) swap(p.x, p.y);
  return edges;
kdTree.h
Description: KD-tree (2d, can be extended to 3d)
                                                     bac5b0, 63 lines
typedef long long T;
typedef Point <T> P;
const T INF = numeric limits <T>::max();
bool on x(const P& a, const P& b) { return a.x < b.x; }
bool on y(const P& a, const P& b) { return a.y < b.y; }
struct Node {
 P pt; // if this is a leaf, the single point in it
  T x0 = INF, x1 = -INF, y0 = INF, y1 = -INF; // bounds
  Node *first = 0, *second = 0;
  T distance (const P& p) { // min squared distance to a ←
    T x = (p.x < x0 ? x0 : p.x > x1 ? x1 : p.x);
    T y = (p.y < y0 ? y0 : p.y > y1 ? y1 : p.y);
    return (P(x,y) - p) \cdot dist2();
  Node(vectorP&& vp) : pt(vp[0]) {
    for (P p : vp) {
      x0 = min(x0, p.x); x1 = max(x1, p.x);
      y0 = min(y0, p.y); y1 = max(y1, p.y);
    if (vp.size() > 1) {
      // split on x if width >= height (not ideal...)
      sort(all(vp), x1 - x0 >= y1 - y0 ? on x : on y);
      // divide by taking half the array for each child (not
      // best performance with many duplicates in the middle←
      int half = sz(vp)/2;
      first = new Node({vp.begin(), vp.begin() + half});
      second = new Node({vp.begin() + half, vp.end()});
};
struct KDTree {
```

Node* root;

```
KDTree(const vector <P>& vp): root(new Node({all(vp)})) {}
  pair <T, P> search (Node *node, const P& p) {
    if (!node->first) {
       // uncomment if we should not find the point itself:
       // if (p = node > pt) return \{INF, P()\};
       return make pair((p - node->pt).dist2(), node->pt);
    Node \star f = \text{node} - \text{first}, \star s = \text{node} - \text{second};
    T bfirst = f->distance(p), bsec = s->distance(p);
    if (bfirst > bsec) swap(bsec, bfirst), swap(f, s);
    // search closest side first, other side if needed
     auto best = search(f, p);
    if (bsec < best.first)</pre>
       best = min(best, search(s, p));
     return best;
  // find nearest point to a point, and its squared distance
  // (requires an arbitrary operator< for Point)
  pair <T, P> nearest (const P& p) {
    return search (root, p);
};
FastDelaunav.h
Description: Fast Delaunay triangulation. Each circumcircle contains none
of the input points. There must be no duplicate points. If all points are on a
line, no triangles will be returned. Should work for doubles as well, though
there may be precision issues in 'circ'. Returns triangles in order {t[0][0],
t[0][1], t[0][2], t[1][0], \dots\}, all counter-clockwise.
Time: \mathcal{O}(n \log n)
"Point.h"
                                                            bf87ec, 88 lines
typedef Point<ll> P:
typedef struct Quad* Q;
typedef int128 t lll; // (can be ll if coords are < 2e4)
P arb (LLONG MAXILONG MAX); // not equal to any other point
struct Quad {
  bool mark; Qo, rot; Pp;
  P F() { return r()->p; }
  Q r() { return rot->rot; }
  Q prev() { return rot->o->rot; }
  Q next() { return r()->prev(); }
bool circ(P p, P a, P b, P c) { // is p in the circumcircle?
  111 p2 = p.dist2(), A = a.dist2()-p2,
       B = b \cdot dist2()-p2, C = c \cdot dist2()-p2;
  return p.cross(a,b)*C + p.cross(b,c)*A + p.cross(c,a)*B > ←
Q makeEdge(P orig, P dest) {
  Q q[] = \{\text{new Quad}\{0,0,0,\text{orig}\}, \text{new Quad}\{0,0,0,\text{arb}\},
             new Quad{0,0,0,dest}, new Quad{0,0,0,arb}};
    q[i] \rightarrow o = q[-i \& 3], q[i] \rightarrow rot = q[(i+1) \& 3];
  return *q;
void splice (Q a, Q b) {
  swap(a\rightarrow o\rightarrow rot\rightarrow o, b\rightarrow o\rightarrow rot\rightarrow o); swap(a\rightarrow o, b\rightarrow o);
Q connect (Q a, Q b) {
  Q q = makeEdge(a->F(), b->p);
  splice(q, a->next());
  splice(q->r(), b);
  return q;
```

```
pair <Q,Q> rec(const vector <P>& s) {
 if (sz(s) <= 3) {
   Q a = makeEdge(s[0], s[1]), b = makeEdge(s[1], s.back()) \leftarrow
    if (sz(s) == 2) return \{a, a \rightarrow r()\};
    splice(a->r(), b);
    auto side = s[0].cross(s[1], s[2]);
   Q c = side ? connect(b, a) : 0;
    return {side < 0 ? c \rightarrow r() : a, side < 0 ? c : b \rightarrow r() };
#define H(e) e->F(), e->p
#define valid(e) (e->F().cross(H(base)) > 0)
 QA, B, ra, rb;
  int half = sz(s) / 2;
  tie(ra, A) = rec({all(s) - half});
  tie(B, rb) = rec(\{sz(s) - half + all(s)\});
  while ((B->p.cross(H(A)) < 0 && (A = A->next())) ||
         (A->p.cross(H(B)) > 0 && (B = B->r()->o)));
 Q base = connect(B->r(), A);
  if (A\rightarrow p == ra\rightarrow p) ra = base\rightarrow r();
  if (B\rightarrow p == rb\rightarrow p) rb = base;
#define DEL(e, init, dir) 0 e = init->dir; if (valid(e)) \
    while (circ(e->dir->F(), H(base), e->F())) { }
     Q t = e \rightarrow dir; \
      splice(e, e->prev()); \
      splice(e->r(), e->r()->prev()); \
      e = t; \
  for (;;) {
   DEL(LC, base->r(), o); DEL(RC, base, prev());
    if (!valid(LC) && !valid(RC)) break;
    if (!valid(LC) | (valid(RC) && circ(H(RC), H(LC))))
      base = connect(RC, base->r());
    else
      base = connect(base->r(), LC->r());
  return { ra, rb };
vector <P> triangulate(vector <P> pts) {
  sort(all(pts)); assert(unique(all(pts)) == pts.end());
  if (sz(pts) < 2) return {};
 Q e = rec(pts).first;
  vector < Q > q = {e};
  int qi = 0;
  while (e->o->F().cross(e->F(), e->p) < 0) e = e->o;
#define ADD { 0 c = e; do { c->mark = 1; pts.push back(c->p) \leftarrow
  q.push back(c\rightarrow r()); c = c\rightarrow next(); } while (c != e); }
 ADD; pts.clear();
  while (qi < sz(q)) if (!(e = q[qi++]) \rightarrow mark) ADD;
  return pts;
```

8.6 3D

PolyhedronVolume.h

Description: Magic formula for the volume of a polyhedron. Faces should point outwards.

```
template<class V, class L>
double signed_poly_volume(const V& p, const L& trilist) {
  double v = 0;
  trav(i, trilist) v += p[i.a].cross(p[i.b]).dot(p[i.c]);
  return v / 6;
```

```
Point3D.h
Description: Class to handle points in 3D space. T can be e.g. double or
template<class T> struct Point3D {
  typedef Point3D P;
  typedef const P& R;
  T x, y, z;
  explicit Point3D(T x=0, T y=0, T z=0): x(x), y(y), z(z) \leftarrow
  bool operator<(R p) const {</pre>
    return tie(x, y, z) < tie(p.x, p.y, p.z); }
  bool operator==(R p) const {
    return tie(x, y, z) == tie(p.x, p.y, p.z); }
  P operator+(R p) const { return P(x+p.x, y+p.y, z+p.z); }
  P operator-(R p) const { return P(x-p.x, y-p.y, z-p.z); }
  P operator*(T d) const { return P(x*d, y*d, z*d); }
  P operator/(T d) const { return P(x/d, y/d, z/d); }
  T dot(R p) const { return x*p.x + y*p.y + z*p.z; }
  P cross(R p) const {
    return P(y*p.z - z*p.y, z*p.x - x*p.z, x*p.y - y*p.x);
  T dist2() const { return x*x + y*y + z*z; }
  double dist() const { return sqrt((double)dist2()); }
  //Azimuthal angle (longitude) to x-axis in interval [-pi, \leftarrow
  double phi() const { return atan2(y, x); }
  //Zenith angle (latitude) to the z-axis in interval [0, pi
  double theta() const { return atan2(sqrt(x*x+v*v),z); }
  P unit() const { return *this/(T) dist(); } //makes dist()←
  //returns unit vector normal to *this and p
  P normal(P p) const { return cross(p).unit(); }
  //returns point rotated 'angle' radians ccw around axis
  P rotate (double angle, P axis) const {
    double s = \sin(\text{angle}), c = \cos(\text{angle}); P u = axis.unit() \leftarrow
    return u*dot(u)*(1-c) + (*this)*c - cross(u)*s;
};
Description: Computes all faces of the 3-dimension hull of a point set. *No
four points must be coplanar*, or else random results will be returned. All
faces will point outwards.
Time: \mathcal{O}\left(n^2\right)
"Point3D.h"
                                                       c172e9, 49 lines
typedef Point3D<double> P3:
struct PR {
  void ins(int x) \{ (a == -1 ? a : b) = x; \}
  void rem(int x) { (a == x ? a : b) = -1; }
  int cnt() { return (a != -1) + (b != -1); }
  int a, b;
struct F { P3 q; int a, b, c; };
vector <F> hull3d(const vector <P3>& A) {
  assert(sz(A) >= 4);
  vector vector PR>> E(sz(A), vector PR>(sz(A), {-1, -1}));
#define E(x,y) E[f.x][f.y]
  vector <F> FS;
  auto mf = [&](int i, int j, int k, int l) {
    P3 q = (A[j] - A[i]) \cdot cross((A[k] - A[i]));
    if (q.dot(A[1]) > q.dot(A[i]))
```

```
q = q * -1;
   F f{q, i, j, k};
   E(a,b).ins(k); E(a,c).ins(j); E(b,c).ins(i);
   FS.push back(f);
 rep(i,0,4) rep(j,i+1,4) rep(k,j+1,4)
   mf(i, j, k, 6 - i - j - k);
 rep(i,4,sz(A)) {
   rep(j,0,sz(FS)) {
     F f = FS[j];
     if(f.q.dot(A[i]) > f.q.dot(A[f.a])) {
       E(a,b).rem(f.c);
       E(a,c).rem(f.b);
       E(b,c).rem(f.a);
       swap(FS[j--], FS.back());
       FS.pop back();
   int nw = sz(FS);
   rep(j,0,nw) {
     F f = FS[j];
#define C(a, b, c) if (E(a,b).cnt() != 2) mf(f.a, f.b, i, f.←
     C(a, b, c); C(a, c, b); C(b, c, a);
 trav(it, FS) if ((A[it.b] - A[it.a]).cross(
   A[it.c] - A[it.a]).dot(it.q) <= 0) swap(it.c, it.b);
 return FS;
```

sphericalDistance.h

Description: Returns the shortest distance on the sphere with radius radius between the points with azimuthal angles (longitude) f1 (ϕ_1) and f2 (ϕ_2) from x axis and zenith angles (latitude) t1 (θ_1) and t2 (θ_2) from z axis. All angles measured in radians. The algorithm starts by converting the spherical coordinates to cartesian coordinates so if that is what you have you can use only the two last rows. dx*radius is then the difference between the two points in the x direction and d*radius is the total distance between the points θ_1 lines

```
double sphericalDistance(double f1, double t1,
    double f2, double t2, double radius) {
    double dx = sin(t2)*cos(f2) - sin(t1)*cos(f1);
    double dy = sin(t2)*sin(f2) - sin(t1)*sin(f1);
    double dz = cos(t2) - cos(t1);
    double d = sqrt(dx*dx + dy*dy + dz*dz);
    return radius*2*asin(d/2);
}
```

Strings (9)

KMP.h

Description: pi[x] computes the length of the longest prefix of s that ends at x, other than s[0...x] itself (abacaba -> 0010123). Can be used to find all occurrences of a string.

Time: $\mathcal{O}(n)$

```
vi pi(const string& s) {
  vi p(sz(s));
  rep(i,1,sz(s)) {
    int g = p[i-1];
    while (g && s[i] != s[g]) g = p[g-1];
    p[i] = g + (s[i] == s[g]);
}
  return p;
}
vi match(const string& s, const string& pat) {
```

```
UCSD
  vi p = pi(pat + '\0' + s), res;
  rep(i,sz(p)-sz(s),sz(p))
    if (p[i] == sz(pat)) res.push back(i - 2 * sz(pat));
  return res;
Zfunc.h
Description: z[x] computes the length of the longest common prefix of s[i:]
and s, except z[0] = 0. (abacaba -> 0010301)
Time: \mathcal{O}(n)
                                                         3ae526, 12 lines
vi Z(string S) {
  vi z(sz(S));
  int l = -1, r = -1;
  rep(i,1,sz(S)) {
    z[i] = i >= r ? 0 : min(r - i, z[i - l]);
    while (i + z[i] < sz(S) && S[i + z[i]] == S[z[i]])
      z[i]++;
    if (i + z[i] > r)
     l = i, r = i + z[i];
  return z;
Manacher.h
Description: For each position in a string, computes p[0][i] = half length
of longest even palindrome around pos i, p[1][i] = longest odd (half rounded
down).
Time: \mathcal{O}(N)
                                                        e7ad79, 13 lines
array < vi, 2> manacher (const string& s) {
  int n = sz(s):
  array \langle vi, 2 \rangle p = \{vi(n+1), vi(n)\};
  rep(z,0,2) for (int i=0,l=0,r=0; i < n; i++) {
    int t = r-i+!z;
    if (i<r) p[z][i] = min(t, p[z][l+t]);
    int L = i-p[z][i], R = i+p[z][i]-!z;
    while (L>=1 && R+1<n && s[L-1] == s[R+1])
      p[z][i]++, L--, R++;
    if (R>r) l=L, r=R;
  return p;
MinRotation.h
Description: Finds the lexicographically smallest rotation of a string.
Usage: rotate(v.begin(), v.begin()+min_rotation(v), v.end());
Time: \mathcal{O}(N)
int min rotation(string s) {
  int a=0, N=sz(s); s += s;
  rep(b,0,N) rep(k,0,N) {
    if (a+k == b | | s[a+k] < s[b+k]) {b += max(0, k-1); \leftarrow
         break;}
    if (s[a+k] > s[b+k]) { a = b; break; }
  return a;
SuffixArray.h
Description: Builds suffix array for a string. sa[i] is the starting index
```

Description: Builds suffix array for a string. sa[i] is the starting index of the suffix which is i'th in the sorted suffix array. The returned vector is of size n+1, and sa[0] = n. The lcp array contains longest common prefixes for neighbouring strings in the suffix array: lcp[i] = lcp(sa[i], sa[i-1]), lcp[0] = 0. The input string must not contain any zero bytes. Time: $\mathcal{O}(n \log n)$

```
struct SuffixArray {
  vi sa, lcp;
  SuffixArray(string& s, int lim=256) { // or basic_string<
  int>
```

```
int n = sz(s) + 1, k = 0, a, b;
    vi x(all(s)+1), y(n), ws(max(n, lim)), rank(n);
    sa = lcp = y, iota(all(sa), 0);
    for (int j = 0, p = 0; p < n; j = \max(1, j * 2), \lim = p \leftarrow
      p = j, iota(all(y), n - j);
      rep(i,0,n) if (sa[i] >= j) y[p++] = sa[i] - j;
      fill (all (ws), 0);
      rep(i,0,n) ws[x[i]]++;
      rep(i,1,\lim) ws[i] += ws[i-1];
      for (int i = n; i--;) sa[-ws[x[y[i]]]] = y[i];
      swap(x, y), p = 1, x[sa[0]] = 0;
      rep(i,1,n) a = sa[i - 1], b = sa[i], x[b] =
        (y[a] == y[b] \&\& y[a + j] == y[b + j]) ? p - 1 : p \leftarrow
   }
   rep(i,1,n) rank[sa[i]] = i;
    for (int i = 0, j; i < n - 1; lcp[rank[i++]] = k)</pre>
      for (k && k--, j = sa[rank[i] - 1];
          s[i + k] == s[j + k]; k++);
 }
};
```

SuffixTree.h

Description: Ukkonen's algorithm for online suffix tree construction. Each node contains indices $[l,\,r)$ into the string, and a list of child nodes. Suffixes are given by traversals of this tree, joining $[l,\,r)$ substrings. The root is 0 (has $l=-1,\,r=0$), non-existent children are -1. To get a complete tree, append a dummy symbol – otherwise it may contain an incomplete path (still useful for substring matching, though).

Time: $\mathcal{O}\left(26N\right)$ aae0b8, 50 lines struct SuffixTree {

```
enum { N = 200010, ALPHA = 26 }; //N \sim 2*maxlen+10
int toi(char c) { return c - 'a'; }
string a; // v = cur node, q = cur position
int t[N][ALPHA], l[N], r[N], p[N], s[N], v=0, q=0, m=2;
void ukkadd(int i, int c) { suff:
  if (r[v]<=q) {
    if (t[v][c]==-1) { t[v][c]=m; l[m]=i;
      p[m++]=v; v=s[v]; q=r[v]; goto suff; }
    v=t[v][c]; q=l[v];
  if (q==-1 || c==toi(a[q])) q++; else {
    l[m+1]=i; p[m+1]=m; l[m]=l[v]; r[m]=q;
    p[m]=p[v]; t[m][c]=m+1; t[m][toi(a[q])]=v;
    l[v]=q; p[v]=m; t[p[m]][toi(a[l[m]])]=m;
    v=s[p[m]]; q=l[m];
    while (q < r[m]) { v = t[v][toi(a[q])]; q + = r[v] - l[v]; }
    if (q==r[m]) s[m]=v; else s[m]=m+2;
    q=r[v]-(q-r[m]); m+=2; goto suff;
}
SuffixTree(string a): a(a) {
  fill (r, r+N, sz(a));
  memset(s, 0, sizeof s);
  memset(t, -1, sizeof t);
  fill(t[1],t[1]+ALPHA,0);
  s[0] = 1; l[0] = l[1] = -1; r[0] = r[1] = p[0] = p[1] = \leftarrow
  rep(i,0,sz(a)) ukkadd(i, toi(a[i]));
// example: find longest common substring (uses ALPHA = ←
pii best;
```

int lcs(int node, int i1, int i2, int olen) {

if (l[node] <= i1 && i1 < r[node]) return 1;</pre>

```
if (l[node] <= i2 && i2 < r[node]) return 2;</pre>
    int mask = 0, len = node ? olen + (r[node] - l[node]) : ←
    rep(c,0,ALPHA) if (t[node][c] != -1)
      mask = lcs(t[node][c], i1, i2, len);
    if (mask == 3)
      best = max(best, {len, r[node] - len});
    return mask;
  static pii LCS(string s, string t) {
    SuffixTree st(s + (char)('z' + 1) + t + (char)('z' + 2))\leftarrow
    st.lcs(0, sz(s), sz(s) + 1 + sz(t), 0);
    return st.best;
 }
};
Hashing.h
Description: Self-explanatory methods for string hashing.
// Arithmetic mod 2^64-1. 2x slower than mod 2^64 and more
// code, but works on evil test data (e.g. Thue-Morse, where
// ABBA... and BAAB... of length 2^10 hash the same mod 2^←
// "typedef ull H;" instead if you think test data is random←
// or work mod 10^9+7 if the Birthday paradox is not a ←
    problem.
struct H {
  typedef uint64 t ull;
  ull x: H(ull x=0) : x(x) {}
#define OP(0,A,B) H operator O(H o) { ull r = x; asm \
  (A "addq %%rdx, %0\n adcq $0,%0" : "+a"(r) : B); return r;←
  OP(+,,"d"(o.x)) OP(*,"mul %1\n", "r"(o.x) : "rdx")
  H operator-(H o) { return *this + \sim0.x; }
  ull get() const { return x + !\sim x; }
  bool operator==(H o) const { return get() == o.get(); }
  bool operator<(H o) const { return get() < o.get(); }</pre>
static const H C = (11)1e11+3; // (order ~ 3e9; random also ←
struct HashInterval {
  vector <H> ha, pw;
  HashInterval(string& str): ha(sz(str)+1), pw(ha) {
    pw[0] = 1;
    rep(i,0,sz(str))
      ha[i+1] = ha[i] * C + str[i],
      pw[i+1] = pw[i] * C;
  H hashInterval(int a, int b) { // hash [a, b)
    return ha[b] - ha[a] * pw[b - a];
 }
};
vector <H> getHashes(string& str, int length) {
  if (sz(str) < length) return {};</pre>
  H h = 0, pw = 1;
  rep(i,0,length)
   h = h * C + str[i], pw = pw * C;
  vector <H> ret = {h};
  rep(i,length,sz(str)) {
    ret.push back(h = h * C + str[i] - pw * str[i-length]);
  return ret;
H hashString(string& s) { H h{}; trav(c,s) h=h*C+c; return h←
```

AhoCorasick DivideAndConquerDP MosAlgo

AhoCorasick.h

Description: Aho-Corasick tree is used for multiple pattern matching. Initialize the tree with create(patterns). $\operatorname{find}(\operatorname{word})$ returns for each position the index of the longest word that ends there, or -1 if none. $\operatorname{findAll}(_, \operatorname{word})$ finds all words (up to $N\sqrt{N}$ many if no duplicate patterns) that start at each position (shortest first). Duplicate patterns are allowed; empty patterns are not. To find the longest words that start at each position, reverse all input. **Time:** create is $\mathcal{O}(26N)$ where N is the sum of length of patterns. find is $\mathcal{O}(M)$ where M is the length of the word. findAll is $\mathcal{O}(NM) \cdot_{716\text{ac4}, 67}$ lines

```
struct AhoCorasick {
  enum {alpha = 26, first = 'A'};
  struct Node {
    // (nmatches is optional)
    int back, next[alpha], start = -1, end = -1, nmatches = -
   Node(int v) { memset(next, v, sizeof(next)); }
  };
  vector < Node> N;
  vector <int> backp;
  void insert(string& s, int j) {
    assert(!s.empty());
    int n = 0;
    trav(c, s) {
     int& m = N[n].next[c - first];
     if (m == -1) { n = m = sz(N); N.emplace back(-1); }
    if (N[n].end == -1) N[n].start = j;
    backp.push back(N[n].end);
   N[n] \cdot end = j;
   N[n].nmatches++;
  AhoCorasick(vector<string>& pat) {
   N. emplace back (-1);
    rep(i,0,sz(pat)) insert(pat[i], i);
   N[0].back = sz(N);
   N.emplace back(0);
    queue<int> q;
    for (q.push(0); !q.empty(); q.pop()) {
      int n = q.front(), prev = N[n].back;
      rep(i,0,alpha) {
        int &ed = N[n].next[i], y = N[prev].next[i];
        if (ed == -1) ed = y;
        else {
          N[ed] \cdot back = y;
          (N[ed].end == -1 ? N[ed].end : backp[N[ed].start])
            = N[v] end;
          N[ed].nmatches += N[y].nmatches;
          q.push(ed);
  vi find (string word) {
    vi res; // 11 count = 0;
    trav(c, word) {
     n = N[n] \cdot next[c - first];
     res.push back(N[n].end);
      // count += N[n]. nmatches;
   return res;
  vector <vi> find All (vector < string > & pat, string word) {
    vi r = find(word);
    vector <vi> res(sz(word));
    rep(i,0,sz(word)) {
```

```
int ind = r[i];
  while (ind != -1) {
    res[i - sz(pat[ind]) + 1].push_back(ind);
    ind = backp[ind];
  }
}
return res;
};
```

Various (10)

10.1 Dynamic programming

DivideAndConquerDP.h

Description: Given $a[i] = \min_{lo(i) \le k < hi(i)} (f(i, k))$ where the (minimal) optimal k increases with i, computes a[i] for i = L..R - 1. **Time:** $\mathcal{O}\left((N + (hi - lo))\log N\right)$

```
struct DP { // Modify at will:
    int lo(int ind) { return 0; }
    int hi(int ind) { return dp[ind][k]; }
    void store(int ind, int k, ll v) { res[ind] = pii(k, v); }

void rec(int L, int R, int LO, int HI) {
    if (L >= R) return;
    int mid = (L + R) >> 1;
    pair<ll, int> best(HONG_MAX, LO);
    rep(k, max(LO, lo(mid)), min(HI, hi(mid)))
        best = min(best, make_pair(f(mid, k), k));
    store(mid, best.second, best.first);
    rec(L, mid, LO, best.second+1);
    rec(mid+1, R, best.second, HI);
}

void solve(int L, int R) { rec(L, R, INT_MIN, INT_MAX); }
```

When doing DP on intervals:

 $a[i][j] = \min_{i < k < j} (a[i][k] + a[k][j]) + f(i,j), \text{ where the (minimal) optimal } k \text{ increases with both } i \text{ and } j, \text{ one can solve intervals in increasing order of length, and search } k = p[i][j] \text{ for } a[i][j] \text{ only between } p[i][j-1] \text{ and } p[i+1][j].$ This is known as Knuth DP. Sufficient criteria for this are Monotonicity: $f(b,c) \leq f(a,d)$ and Quadrangle inequality: $f(a,c) + f(b,d) \leq f(a,d) + f(b,c)$ for all $a \leq b \leq c \leq d$. Consider also: LineContainer (ch. Data structures), monotone queues, ternary search.

10.2 Debugging tricks

- signal(SIGSEGV, [](int) { _Exit(0); }); converts segfaults into Wrong Answers. Similarly one can catch SIGABRT (assertion failures) and SIGFPE (zero divisions). _GLIBCXX_DEBUG failures generate SIGABRT (or SIGSEGV on gcc 5.4.0 apparently).
- feenableexcept(29); kills the program on NaNs (1), 0-divs (4), infinities (8) and denormals (16).

10.3 Optimization tricks

10.3.1 Bit hacks

- x & -x is the least bit in x.
- for (int x = m; x;) { --x &= m; ... } loops over all subset masks of m (except m itself).
- c = x&-x, r = x+c; $(((r^x) >> 2)/c) \mid r$ is the next number after x with the same number of bits set.
- rep(b,0,K) rep(i,0,(1 << K))
 if (i & 1 << b) D[i] += D[i^(1 << b)];
 computes all sums of subsets.</pre>

10.3.2 Pragmas

- #pragma GCC optimize ("Ofast") will make GCC auto-vectorize for loops and optimizes floating points better (assumes associativity and turns off denormals).
- #pragma GCC target ("avx,avx2") can double performance of vectorized code, but causes crashes on old machines.
- #pragma GCC optimize ("trapv") kills the program on integer overflows (but is really slow).

MosAlgo.h

return idx;

Description: Mo's on trees with path queries to print the smallest non-negative integer on simple path a_i and b_i .

```
Time: \mathcal{O}\left(N\sqrt{N}\log(N)\right)
                                                          b5ce43, 115 lines
const int maxn = int(2e5)+5, inf = int(1e9)+5, block = 500;
int A[maxn], BIT[maxn+5], ans[maxn], start[maxn], en[maxn], \leftarrow
     occ[maxn], cnt[maxn];
pair < pair < int, int>, int> Q[maxn];
vector <int> dis;
vector <pair <int, int>> graph [maxn];
inline int left(int node) { return (node<<1); }</pre>
inline int right(int node) { return (node<<1)+1; }</pre>
void upd(int idx, int v) {
     while(idx < maxn) {</pre>
         BIT[idx] += v;
         idx += (idx\&-idx);
}
int gry() {
     int idx = 0, b = 16, s = 0;
    while(b >= 0) {
         if(BIT[idx+(1<<b)]+s == idx+(1<<b)) {
             idx += (1<<b);
             s += BIT[idx];
         b--:
```

```
void dfs0(int node, int par, int into) {
    A[node] = into, start[node] = int(dis.size());
    dis.push back(node);
    for(auto it: graph[node]) {
        if(it.first != par) dfs0(it.first, node, it.second);
    en[node] = int(dis.size());
    dis.push back(node);
inline int cmp(pair < pair < int, int>, int>& a, pair < pair < int, ←
     int>, int>& b) {
    if(a.first.first/block != b.first.first/block) return a.←
         first.first < b.first.first;</pre>
    else if((a.first.first/block)%2) return a.first.second >←
          b. first.second;
    else return a.first.second < b.first.second;</pre>
void rem(int v) {
    v++:
    cnt[v]--;
    if(cnt[v] == 0) upd(v, -1);
void add(int v) {
    v++:
    cnt[v]++;
    if(cnt[v] == 1) upd(v, 1);
void act(int node) {
    if(A[node] >= maxn) return;
    occ[node]++;
    if(occ[node] == 2) rem(A[node]);
    else add(A[node]);
void deact(int node) {
    if(A[node] >= maxn) return;
    occ[node]--;
    if(occ[node] == 1) add(A[node]);
    else rem(A[node]);
}
int main(void) {
    int n, q, u, v, x;
    scanf("%d%d", &n, &q);
    for(int i = 1; i < n; i++) {
        scanf("%d%d%d", &u, &v, &x);
        graph[u].push back(\{v, x\}), graph[v].push back(\{u, x \leftarrow
             });
    dfs0(0, -1, inf);
    //for(auto it: dis) cout << it << " ";
     //cout << "\n";
    for(int i = 0; i < q; i++) {
        scanf("%d%d", &u, &v);
        if(start[u] > start[v]) swap(u, v);
         if(start[u] \leftarrow start[v] && start[v] \leftarrow en[u]) Q[i] =\leftarrow
               {{start[u]+1, start[v]}, i};
         else Q[i] = {{en[u], start[v]}, i};
```

```
//cout \ll Q[i]. first. first \ll ":" \ll Q[i]. first. \leftarrow
             second \ll " " \ll Q[i].second \ll " n";
    sort (Q, Q+q, cmp);
    int L = 0, R = 0;
    act (dis [0]);
    for(int i = 0; i < q; i++) {
        int ql = Q[i].first.first, qr = Q[i].first.second;
        if(ql <= qr) {
             while(R < qr) act(dis[++R]);
            while(L < ql) deact(dis[L++]);
            while(L > ql) act(dis[--L]);
            while(R > qr) deact(dis[R--]);
            ans[Q[i].second] = qry();
        else ans[Q[i].second] = 0;
    for(int i = 0; i < q; i++) printf("%d\n", ans[i]);
PairHash.h
Description: Demonstrates hashing for pairs for use in unordered map
                                                        fa756e, 7 lines
template < typename T1 , typename T2 >
struct pair hash {
size t operator () ( const pair < T1 , T2 > & p ) const {
return hash \langle T1 \rangle() ( p . first ) ^ hash \langle T2 \rangle() ( p . \leftarrow
};
unordered map < pair < int , int > , int , pair hash < int , ←
      int > M
FastScanner.java
Description: Fast scanner class
                                                        f05af3, 16 lines
public class MyScanner {
    BufferedReader br = new BufferedReader(new ←
         InputStreamReader(System.in));
    PrintWriter out = new PrintWriter(new 	
         BufferedOutputStream(System.out));
    String next() {
        while (st == null || !st.hasMoreElements()) {
            try { st = new StringTokenizer(br.readLine()); }
            catch (IOException e) { e.printStackTrace(); }
        return st.nextToken();
    int nextInt() { return Integer.parseInt(next()); }
    long nextLong() { return Long.parseLong(next()); }
    double nextDouble() { return Double.parseDouble(next()); ←
```