# UC San Diego

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# MnM

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# Mathematics (2)

# 2.1 Equations

$$ax + by = e$$

$$cx + dy = f$$

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

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

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.

#### 2.2Recurrences

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

# 2.4 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  $\theta$ , 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  $A = \sqrt{(p-a)(p-b)(p-c)(p-d)}$ .

# Geometry - 2

# 2.5.1 Spherical coordinates



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

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

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

# 2.6 Sums

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

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

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

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

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

# 2.6.1 Discrete distributions Binomial distribution

The number of successes in n independent ves/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  $\kappa$  and independently of the time since the last event is  $Po(\lambda), \lambda = t\kappa.$ 

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

# 2.6.2 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.

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

# OrderStatisticTree HashMap Matrix Treap

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

# **Exponential distribution**

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

$$f(x) = \begin{cases} \lambda e^{-\lambda x} & x \ge 0\\ 0 & x < 0 \end{cases}$$

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

# Normal distribution

Most real random values with mean  $\mu$  and variance  $\sigma^2$  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

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

# 2.7 Markov 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.

 $\pi$  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.  $\pi_j/\pi_i$  is the expected number of visits in state j between two visits in state i.

For a connected, undirected and non-bipartite graph, where the transition probability is uniform among all neighbors,  $\pi_i$  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  $\mathbf{A}$  and  $\mathbf{G}$ , such that all states in  $\mathbf{A}$  are absorbing  $(p_{ii}=1)$ , and all states in  $\mathbf{G}$  leads to an absorbing state in  $\mathbf{A}$ . 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 polygon.

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

```
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:**  $O(\log N)$ 

782797, 16 li

#### HashMap.h

**Description:** Hash map with the same API as unordered map, but  $\sim 3x$  faster. Initial capacity must be a power of 2 (if provided).

```
#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 number
   ll operator()(ll x) const { return __builtin_bswap64(x*C); }
};
__gnu_pbds::gp_hash_table<ll,int,chash> h({},{},{},{},{1<<16});</pre>
```

# 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

 $if(t\rightarrow L) t\rightarrow L\rightarrow P = t;$ 

 $if(t\rightarrow R) t\rightarrow R\rightarrow P = t;$ 

```
template<class T, int N> struct Matrix {
  typedef Matrix M;
  array<array<T, N>, N> d{};
```

```
M operator*(const M& m) const {
    Ma;
    rep(i,0,N) rep(j,0,N)
      rep(k,0,N) a.d[i][j] += d[i][k]*m.d[k][j];
  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 >= 0);
    Ma, 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 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 = 1:
    R = r:
    P = NULL;
    v = val;
    sz = 1;
    lz = 0;
    prior = rand();
};
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)
```

```
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\rightarrow L); upd lz(t\rightarrow 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 \overline{P}(t); upd \overline{P}(1); upd \underline{P}(r);
  if(!t)
     l = r = NULL;
     return;
  int cur key = add+get sz(t->L);
  if(cur key < key)
     split(t\rightarrow R, t\rightarrow R, r, key, cur_key+1);
     l = t;
  else
     split(t\rightarrow L, l, t\rightarrow 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(l); upd lz(r); upd lz(t);
  upd P(1); upd_P(r); upd_P(t);
  if(!l || !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 \overline{P}(1); upd \overline{P}(r); upd P(t);
int climber (node t, bool add)
  int res = (add?get sz(t->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 << " " << t
          ->prior << " " << add+get sz(t->L) << "\n";
     printer(t->R, add+get sz(t->L)+1);
void fix (node& t)
  if(t)
    if(t->P) t->P = NULL;
// Initialise
srand (time (NULL));
// Call after each qry/upd, root is overall root of running
     treap.
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>ys; 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()); }

```
ft[x].update(ind(x, y), dif);
  11 query(int x, int y) {
    11 \text{ 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, 43 lines
typedef long long int lli;
const lli maxn = lli(1e5)+5;
struct line {
  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-1])
      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

93d5aa, 57 lines

point values.

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

void update(int x, int y, ll dif) {

for (; x < sz(ys); x = x + 1)

```
typedef long long 11;
const ll \max = ll(1e5)+5, inf = ll(1e18)+5;
struct point {
  11 m, c; // y = mx + c
  point(): m(0), c(inf) {} // ### (inf -> -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 =
      maxn) {
    ll m = (l + 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 (l == 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);
  Il 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)); // ###
    } else {
      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, 56 lines

struct pnode { int res = 0; pnode \*L, \*R; pnode(int v = 0, pnode\* l = NULL, pnode\* r = NULL)

```
res = v;
   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 pnode(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 << " \n";
    if(left >= k)
      return gry(rootL->L, rootR->L, L, (L+R)/2, k);
    else return qry(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++)</pre>
  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.
                                                      8a1f54, 121 lines
const int maxn = int(2e5)+5, maxlog = 20, inf = int(1e9)+5;
int n, m, T[maxn][maxlog+1], TT[maxn][maxlog+1], F[maxn][maxlog
     +1], H[maxn], upar[maxn], P[maxn], taken[maxn], res[maxn];
pair < pair < int, int>, pair < int, int>> E[maxn];
vector <pair <int, int>> graph [maxn];
inline int root(int x) { return ((P[x] == x)?x:(P[x] = root(P[x]
    ]))); }
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+1);
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 \ge 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 >= 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 \le \max\{j; j++\} T[i][j] = -1, TT[i][j] = F[i][
      j] = 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)
                                                      52ddb1, 167 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 | | 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];
  else
    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++)</pre>
    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++)
{
    for(lli j = 0; j <= maxlog; j++) T[i][j] = -1;
}

// initialisation after input
dfs0(0, -1, 0);
init();
totchain = 1;
dfs1(0, -1, 0);
build(1, 0, curst-1);</pre>
```

# Numerical (4)

# 4.1 Polynomials and recurrences

```
Polynomial.h
                                                       c9b7b0, 17 lines
struct Poly {
  vector <double> a;
  double operator()(double x) const {
    double val = 0;
    for(int i = sz(a); i --;) (val *= x) += a[i];
    return val;
  void diff() {
    rep(i,1,sz(a)) a[i-1] = i*a[i];
    a.pop back();
  void divroot(double x0) {
    double b = a.back(), c; a.back() = 0;
    for(int i=sz(a)-1; i--;) c = a[i], a[i] = a[i+1]*x0+b, b=c;
    a.pop back();
};
PolyRoots.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) {
  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;
```

```
PolvInterpolate.h
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);
  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
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}\left(N^2\right)
"../number-theory/ModPow.h"
                                                         40387d, 20 lines
vector <11 > BerlekampMassey(vector <11 > s) {
 int n = sz(s), L = 0, m = 0;
  vector < ll > C(n), B(n), T;
  C[0] = B[0] = 1;
  ll b = 1;
  rep(i,0,n) { ++m;
    ll d = s[i] \% mod;
    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_{i} 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 <ll > Poly;
ll linearRec(Poly S, Poly tr, ll k) {
  int n = sz(tr);
  auto combine = [&] (Poly a, Poly b) {
    Poly res(n \star 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]) % 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

**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; }

#### HillClimbing.h

Description: Poor man's optimization for unimodal functions<sub>f40e55, 16 lines</sub>

```
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*jmp;
        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;
IntegrateAdaptive.h
Description: Fast integration using an adaptive Simpson's rule.
Usage: double sphereVolume = quad(-1, 1, [](double x) {
return quad(-1, 1, [&](double y) {
return quad(-1, 1, [&](double z) {
return x*x + y*y + z*z < 1; {});{});{});
                                                     92dd79, 15 lines
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) \le 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, 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  $c^T x$ 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  $c^T x$  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=j
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];}
      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] \cdot 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])
                    < MP(D[r][n+1] / D[r][s], B[r])) r = 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;
```

#### 4.3Matrices

#### Determinant.h

Description: Calculates determinant of a matrix. Destroys the matrix. Time:  $\mathcal{O}(N^3)$ bd5cec, 15 lines

```
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
        ll 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;
```

#### SolveLinear.h

**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) {
   double v, bv = 0;
   rep(r,i,n) rep(c,i,m)
     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:

```
08e495, 7 lines
```

```
rep(j,0,n) if (j != i) // instead of <math>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:  $\mathcal{O}(n^2m)$ 

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;</pre>
   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] ^= A[i];
   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:  $\mathcal{O}(n^3)$ 

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];
```

# Tridiagonal.h

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

```
x_0
            0
                                                     x_1
b_2
                                                     x_2
b_3
                                                     x_3
                 0 \quad \cdots \quad q_{n-3} \quad d_{n-2} \quad p_{n-2}
            0
```

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, ..., -1, 1\}, \{0, c_1, c_2, ..., 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)
```

```
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] == 0}
     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;
```

# 4.4 Fourier transforms

FastFourierTransform.h

**Description:** fft(a) computes  $\hat{f}(k) = \sum_{x} a[x] \exp(2\pi i \cdot kx/N)$  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: O(N \log N) with N = |A| + |B| (\sim 1s \text{ for } N = 2^{22})
```

```
typedef complex <double> C:
typedef vector <double> vd;
void fft(vector < C>& a) {
                           __builtin clz(n);
 int n = sz(a), L = 31 -
  static vector < complex < long double >> R(2, 1);
  static vector <C> rt(2, 1); // (^ 10% faster if double)
  for (static int k = 2; k < n; k \neq = 2) {
   R. resize (n); rt. resize (n);
    auto x = polar (1.0L, M Pl / k); // M Pl, 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-rolled)
      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]);
  fft (out);
  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" b82773, 22 lines

```
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));
vector<C> 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  $2^ab+1$ , 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"

```
d75aad, 32 lines
const ll mod = (119 << 23) + 1, root = 62; // = 998244353
// For p < 2^30 there is also e.g. 5 << 25, 7 << 26, 479 << 21
// 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) {
        11 z = rt[j + k] * a[i + j + k] % mod, &ai = a[i + j];
       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 << B;
  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 curL = mod / 2, inv = 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  $c[z] = \sum_{z=x \oplus y} a[x] \cdot b[y]$ , where  $\oplus$  is one of AND, OR, XOR. The size of a must be a power of two.

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

```
fvi 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

 $\bf Description:$  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); }
 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  $\leq$  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  $x \ge 0$  s.t.  $a^x = b \pmod{m}$ . a and m must be coprime.

```
Time: \mathcal{O}(\sqrt{m})

11 modLog(11 a, 11 b, 11 m) {
    assert(_gcd(a, m) == 1);
    11 n = (11) sqrt(m) + 1, e = 1, x = 1, res = HONG_MAX;
    unordered map<11, 11> 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;
```

```
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 * 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 \star c + k \star sumsq(to) - m \star divsum(to, c, k, m);
ModMulLL.h
Description: Calculate a \cdot b \mod c (or a^b \mod c) for 0 \le a, b \le c \le 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) {
```

# ModSqrt.h

ull ans = 1;

return ans;

**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

for (; e; b = mod mul(b, b, mod), e /= 2)

if (e & 1) ans = mod mul(ans, b, mod);

```
"ModPow.h"
                                                      19a793, 24 lines
ll sqrt(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
 11 s = p - 1, n = \hat{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);
 ll 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;
}
}
```

# 5.2 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  $a^{\overline{b}} \mod c$ .

#### Factor.h

**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"
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 = __gcd(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 1;
```

# 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). Gaba01, 11 lines

static BigInteger[] euclid(BigInteger a, BigInteger b) {

BigInteger x = BigInteger.ONE, yy = x;

BigInteger y = BigInteger.ZERO, xx = y;

white (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

Description: Chinese Remainder Theorem.

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

#### Time: $N \log(N)$

# 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 \phi 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^{\phi(n)} \equiv 1 \pmod{n}.

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

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

rep(i,0,LIM) phi[i] = i&1 ? i : i/2;

for(int i = 3; i < LIM; i += 2) if(phi[i] == 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  $\infty$ ; if x is the root of a degree 2 polynomial the a's eventually become cyclic.

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

typedef double d; // for  $N \sim 1e7$ ; long double for  $N \sim 1e9$ 

# 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([](Frac f) { return f.p>=3\*f.q; }, 10); // {1,3}

Time:  $\mathcal{O}(\log(N))$  27ab3e, 25 lines

```
struct Frac { ll p, q; };
template<class F>
Frac fracBS(F f, ll 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) {
      adv += step;
      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.p += lo.p * adv;
    hi.q += lo.q * adv;
    dir = !dir;
    swap(lo, hi);
   A = B; B = !!adv;
 return dir ? hi : lo;
```

# 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 1000000.

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:

$$10^9 + 7, 10^9 + 9, 10^9 + 21, 10^9 + 33, 10^9 + 87, 10^9 + 93, 10^9 + 97, 10^9 + 103, 10^9 + 123, 10^9 + 181, 10^9 + 207, 10^9 + 223$$

# 5.7 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.

#### 5.8 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

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

	I					9		
							3628800	
	I					5 16		
n!	4.0e7	4.8e	8 6.26	e9 8.7€	10 1.3e	$12 \ 2.1e1$	3.6e14	
n	20	25	30	40	50 1	00 150	171	
n!	2e18	2e25	3e32	8e47 3	Be64 9e	$157 \ 6e26$	$52 > \text{DBL\_MA}$	X

# **6.1.2** Cycles

Let  $g_S(n)$  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:  $X^{4n^2} + 3X^{2n^2} + 2X^{n^2} + 2X^{2n+n}$ 

# 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.

$$p(0) = 1, \ p(n) = \sum_{k \in \mathbb{Z} \setminus \{0\}} (-1)^{k+1} p(n - k(3k - 1)/2)$$
$$p(n) \approx 0.145/n \cdot \exp(2.56\sqrt{n})$$

# 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) { 11 a = n % p, b = m % p; if (a < b) return 0: c = c \* fact[a] % p \* invfact[b] % p \* invfact[a - b] % p; n /= p; m /= p; return c;

multinomial.h

Description: Computes 
$$\binom{k_1 + \dots + k_n}{k_1, k_2, \dots, k_n} = \frac{(\sum k_i)!}{k_1! k_2! \dots k_n!}$$
.

11 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);
 return c;
}

# 6.3 General purpose numbers

# 6.3.1 Bernoulli numbers

EGF of Bernoulli numbers is  $B(t) = \frac{t}{e^t - 1}$  (FFT-able).  $B[0,\ldots] = [1,-\frac{1}{2},\frac{1}{6},0,-\frac{1}{20},0,\frac{1}{42},\ldots]$ 

Sums of powers:

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

Euler-Maclaurin formula for infinite sums:

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

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

# 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.  $\pi(j) > \pi(j+1), k+1 \text{ } j:\text{s s.t. } \pi(j) \geq j, k \text{ } j:\text{s s.t.}$  $\pi(j) > j$ .

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

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

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

# 6.3.4 Stirling numbers of the second kind

Partitions of n distinct elements into exactly kgroups.

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

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

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

# 6.3.5 Bell numbers

Total number of partitions of n distinct elements. B(n) = $1, 1, 2, 5, 15, 52, 203, 877, 4140, 21147, \dots$  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

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

$$C_0 = 1, \ C_{n+1} = \frac{2(2n+1)}{n+2}C_n, \ C_{n+1} = \sum C_i C_{n-i}$$

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

- sub-diagonal monotone paths in an  $n \times n$  grid.
- 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.

#### General purpose theorems - 1 6.4

# 6.4.1 Identities

Vandermonde Convolution:

$$\binom{m+n}{1} = \sum_{k=0}^{r} \binom{m}{k} \cdot \binom{r}{n-k}.$$
6.4.2. Cycle Lemma  $n$  (in the context of the context)

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 6.4.3 Sprague Grundy theorem

Every impartial game is equivalent to a nimber. Nimbers are de-fined inductively as

 $*0 = \{\}, *1 = *0, *2 = *0, *1, *(n+1) = *n \cup n, \text{ and } \}$ 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:

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

# 6.4.4 Partisan Game

Can define the negative of a game by interchanging L and R's possible moves. Define G=0 if first player loses. G=Hif 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 604r5senMatricesnforteneratorsalculable by DP.

13

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:  $\begin{bmatrix} -1 & 1 \\ 1 & 0 \end{bmatrix}$ ,  $\begin{bmatrix} 1 & 1 \\ 1 & 0 \end{bmatrix}$ .

# 6.5General 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  $n_n$  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 6.5.2 let Toipanament of raphs.  $r^{l-1}$ .

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 a tournament graph.

# General 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.empty()) {
    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 = \bar{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;
        } 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),
    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<ll, 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});
        else q.modify(its[i], {-dist[i], i});
    };
    while (!q.empty()) {
      s = q \cdot top() \cdot second; q \cdot 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]);
      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];
    return {totflow, totcost};
```

void addEdge(int a, int b, ll c, int rcap = 0) {

# 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.

979bb9, 35 lines

```
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) <= 0) graph[p].erase(y);
      graph[y][p] += inc;
```

# Dinic.h

**Description:** Flow algorithm with complexity  $O(VE \log U)$  where  $U = \max |\text{cap}|$ .  $O(\min(E^{1/2}, V^{2/3})E)$  if U = 1;  $O(\sqrt{V}E)$  for bipartite matching.

```
struct Dinic {
    struct Edge {
        int to, rev;
        ll c, oc;
        ll flow() { return max(oc - c, OLL); } // if you need flows
    };
    vi lvl, ptr, q;
    vector<vector<Edge>> adj;
    Dinic(int n): lvl(n), ptr(n), q(n), adj(n) {}
```

```
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 \mid !f) return f;
    for (int& i = ptr[v]; i < sz(adj[v]); i++) {</pre>
      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, adi[v])
          if (!lvl[e.to] && e.c >> (30 - L))
            q[qe++] = e \cdot to, lvl[e \cdot 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 repre-
sented by an adjacency matrix.
Time: \mathcal{O}(V^3)
                                                      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) {
          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: O(VE)

bool find(int j, vector < vi>& g, vi& btoa, vi& vis) {

if (btoa[j] == -1) return 1;

vis[j] = 1; int di = btoa[j];

trav(e, g[di])

if (!vis[e] && find(e, g, btoa, vis)) {

btoa[e] = di;

return 1;

}

6a3472, 22 lines
```

```
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);
 int res = dfsMatching(g, match);
  vector <bool> lfound(n, true), seen(m);
 vi q, cover;
  rep(i,0,n) if (lfound[i]) q.push back(i);
  while (!q.empty()) {
   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:  $\mathcal{O}\left(N^3\right)$ 

```
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 i = 0;
    for (;;) {
     i = -1;
     rep(k,0,n){
       if (seen[k]) continue;
       if (j == -1 || dist[k] < dist[j]) j = k;</pre>
     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[j];
     v[k] += w, u[R[k]] -= w;
   u[s] += dist[j];
    while (dad[j] >= 0) {
     int d = dad[j];
     R[j] = R[d];
     L[R[j]] = j;
     j = d;
   R[j] = 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:  $\mathcal{O}(n^2)$ 

```
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.empty()) {
    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};
GeneralMatching.h
Description: Matching for general graphs. Fails with probability N/mod.
Time: \mathcal{O}(N^3)
"../numerical/MatrixInverse-mod.h"
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, fi;
  assert (r % 2 == 0);
  if (M != N) do {
    mat.resize (M, vector < 11 > (M));
    rep(i,0,N) {
      mat[i].resize(M);
      rep(j,N,M) {
        int r = rand() % mod;
        mat[i][j] = r, mat[j][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);
    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.
```

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]); // update
    dfs_low[u]
}
else if(v.first != dfs_parent[u]) // a back edge and not
```

# 2sat.h

dfs(i);

number of clauses

special case

**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  $(\sim x)$ .

articulation vertex[dfsRoot] = (rootChildren > 1); //

```
Usage: TwoSat ts(number of boolean variables); ts.either(0, \sim3); // 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, \sim1 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}(N+E), where N is the number of boolean variables, and E is the
```

```
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 = \sim next;
    either (cur, \simli[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

0911c1, 56 lines

**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
```

```
typedef bitset<128> B;
template<class F>
void cliques(vector<B>& eds, F f, B P = ~B(), B X={}, B R={}) {
```

```
if (!P.any()) { if (!X.any()) f(R); return; }
auto q = (P | X)._Find_first();
auto cands = P & ~eds[q];
rep(i,0,sz(eds)) if (cands[i]) {
   R[i] = 1;
   cliques(eds, f, P & eds[i], X & eds[i], R);
   R[i] = P[i] = 0; X[i] = 1;
}
```

#### 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 < bitset < 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, 1);
        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 > mxk) 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].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) {
    rep(i,0,sz(e)) V.push_back({i});
```

# CompressTree LinkCutTree DirectedMST

#### Trees 7.6

# 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, origi index) representing a tree rooted at 0. The root points to itself.

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

```
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]; \};
  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.

if  $(p\rightarrow p)$   $p\rightarrow p\rightarrow push flip();$ 

```
Time: All operations take amortized \mathcal{O}(\log N).
                                                          693483, 90 lines
struct Node { // Splay tree. Root's pp contains tree's parent.
  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] -> 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\rightarrow c[1] == this : -1; }
  void rot(int i, int b) {
    int h = i ^ b;
    Node *x = c[i], *y = b == 2 ? x : x -> c[h], *z = b ? y : x;
    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 \rightarrow c[i ^1] = b ? this : x;
    fix(); x\rightarrow fix(); y\rightarrow fix();
    if (p) p->fix();
    swap(pp, y->pp);
  void splay() {
    for (push flip(); p; ) {
```

```
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() {
    push flip();
    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 tree?
    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 \rightarrow c[0] \rightarrow 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;
```

#### Directed MST.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"
                                                                   a69883, 48 lines
struct Edge { int a, b; ll w; };
struct Node {
  Edge key;
  Node *1, *r;
```

```
ll delta;
  void prop() {
    key.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 kev.w \rightarrow b\rightarrow kev.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] - top();
      heap[u]->delta -= e.w, pop(heap[u]);
      res += e.w, u = uf.find(e.a);
      if (seen[u] == s) {
        Node* cyc = 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;
```

#### 7.7Math

# 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 rokinn midskothnaeterminant; this yields the number of directed spanning trees deoretes diff G is undirected iff  $\mathcal{U}_1 + \mathcal{U}_n$  and row even and for every  $k = 1 \dots n$ ,

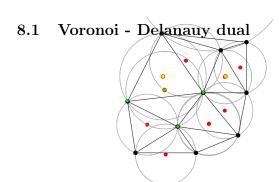
$$\sum_{i=1}^{k} d_i \le k(k-1) + \sum_{i=k+1}^{n} \min(d_i, k).$$

# 7.7.3 Konig's theorem

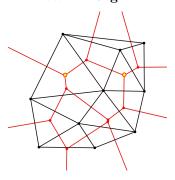
In any bipartite graph, the number of edges in a maximum mathing equals the number of vertices in a minimum vertex cover. To exhibit the vertex cover:

- 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 7.7.4 th Minumum, Edgell coveres on the left that are not If a miniform 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 Maximum Independent sets of a graph, find a max clique of the complement. If the graph is bipartite, see Minimum Vertex Cover.

# Geometry (8)



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

#### Point.h

**Description:** Class to handle points in the plane. T can be e.g. double or long long. (Avoid int.) 47ec0a. 28 lines

```
template \langle class T \rangle int sgn(T x) \{ return (x > 0) - (x < 0); \}
template<class T>
struct Point {
 typedef Point P;
 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); }
  bool operator==(P p) const { return tie(x,y)==tie(p.x,p.y); }
 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.y - 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 sqrt((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 containing 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 supposed 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 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 on Segment = segDist(a,b,p) < 1e-10;

"Point.h" 5c88f4, 6 lines

```
typedef Point<double> P;
double segDist(P& s, P& e, P& p) {
   if (s==e) return (p-s).dist();
   auto d = (e-s).dist2(), t = min(d,max(.0,(p-s).dot(e-s)));
   return ((p-s)*d-(e-s)*t).dist()/d;
}
```

# 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<II> 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"
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

a1ee63, 19 lines

#### 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<ll> 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 \(^{\sigma}\)! not always return true if the intersection is at the boundary point(s). Usage: auto res = lineInter(s1,e1,s2,e2);

```
if (res.first == 1)
cout << "intersection point at " << res.second << endl;</pre>
"Point.h"
                                                         a01f81, 8 lines
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;
```

```
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).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>.

```
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:

Apply the linear transformation (translation, rotation and scaling) which takes line p0-p1 to line q0-q1 to point r. "Point.h"



3af81c, 9 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.

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

#### Angle.h

**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

```
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}; }
 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)\}; \}
 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 between
// them, i.e., the angle that covers the defined line segment.
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)\};
```

#### 8.3Circles

# CircleIntersection.h

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

```
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*d2;
 if (sum*sum < d2 | | dif*dif > d2) return false;
```

```
P mid = a + vec*p, per = vec.perp() * sqrt(fmax(0, h2) / d2);
\starout = {mid + per, mid - 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. "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)
```

return sum;

```
"../../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();
    auto det = a * a - b;
    if (det \le 0) return arg(p, q) * r2;
    auto s = max(0., -a-sqrt(det)), t = min(1., -a+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);
```

#### circumcircle.h Description:

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. "Point.h" 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"

```
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[i]) / 2;
     r = (o - ps[i]).dist();
     rep(k,0,j) if ((o - ps[k]).dist() > r * EPS) {
       o = ccCenter(ps[i], ps[j], ps[k]);
        r = (o - ps[i]).dist();
   }
  return {o, r};
```

#### kIntersection.h

**Description:** Given n circles, for all k <= n, computes the area of regions part of atleast k circles.

Time:  $\mathcal{O}\left(n^2\right)$ 

0e6b2d, 65 lines

```
const int N = 22222;
const double EPS = 1e-8;
const double PI = acos(-1.0);
typedef complex <double > Point;
int n, m;
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) \le 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) {
    addEvent(-PI, 0);
    addEvent(+PI, 0);
    for (int j = 0; j < n; ++ j) {
      if (i != i) {
        if (\operatorname{sgn}(\operatorname{abs}(c[i] - c[j]) - \operatorname{abs}(r[i] - r[j])) \le 0) {
          if (\operatorname{sgn}(r[i] - r[i]) \le 0) {
             addPair(-PI, +PI);
        } else {
          if (sgn(abs(c[i] - c[j]) - (r[i] + r[j])) >= 0) {
          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]) /
                (2 *
                r[i] * d));
          Point a = b * polar(-t);
          Point c = b * polar(+t);
          addPair(arg(a), arg(c));
     }
   }
    sort (event, event + m);
    int count = event[0].second;
    for (int j = 1; j < m; ++ j) {
      double delta = event[j].first - event[j - 1].first;
      result[count] += r[i] * r[i] * (delta - sin(delta));
      result[count] += det(radius(i, event[j - 1].first),
           radius (i,
            event[j].first));
      count += event[j].second;
```

# 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\langle P \rangle v = \{P\{4,4\}, P\{1,2\}, P\{2,1\}\};
bool in = inPolygon(v, P{3, 3}, false);
Time: \mathcal{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) \le eps) return !strict;
    cnt ^= ((a.y<p[i].y) - (a.y<q.y)) * a.cross(p[i], q) > 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"

template<class T> T polygonArea2(vector < Point < T>>& v) { T = 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:  $\mathcal{O}(n)$ 

"Point.h" 9706dc, 9 lines typedef Point <double> P; P polygonCenter(const vector <P>& v) {  $\hat{P}$  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]);

# PolygonCut.h

return res / A / 3;

#### 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" f2b7d4, 13 lines

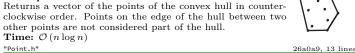
```
typedef Point <double> P;
vector <P> polygonCut(const vector <P>& poly, P s, P e) {
 vector <P> res:
  rep(i,0,sz(poly)) {
    P cur = poly[i], prev = i ? poly[i-1] : poly.back();
    bool side = s.cross(e, cur) < 0;
    if (side != (s.cross(e, prev) < 0))
      res.push back(lineInter(s, e, cur,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

# ConvexHull.h

# Description:

clockwise 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"



```
typedef Point<ll> P;
vector <P> convexHull(vector <P> pts) {
 if (sz(pts) <= 1) return pts;</pre>
 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])\};
```

#### HullDiameter.h

Description: Returns the two points with max distance on a convex hull (ccw, no duplicate/colinear points). c571b8, 12 lines

```
typedef Point<ll> P;
array <P. 2> hullDiameter (vector <P> S) {
  int n = sz(S), j = n < 2 ? 0 : 1;
  pair < 11, 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)
  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;
bool inHull(const vector <P>& l, P p, bool strict = true) {
  int a = 1, b = sz(1) - 1, r = !strict;
  if (sz(l) < 3) return r && onSegment(l[0], l.back(), p);
  if (sideOf(l[0], l[a], l[b]) > 0) swap(a, b);
  if (sideOf(l[0], l[a], p) >= r || sideOf(l[0], l[b], p) <= -r)
   return false;
  while (abs(a - b) > 1) {
   int c = (a + b) / 2;
    (sideOf(l[0], l[c], p) > 0 ? b : a) = c;
```

#### LineHullIntersection.h

return sgn(l[a].cross(l[b], p)) < r;

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 (i, i+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:  $\mathcal{O}(N + Q \log n)$ 

```
"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) {
    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) = m;
  return lo;
#define cmpL(i) sgn(line[0].cross(poly[i], line[1]))
array <int, 2> lineHull(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;</pre>
    (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 <arrav <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 i = it ->second:
       Pd = 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)
"Point.h"
                                                     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; }
 P pt; // if this is a leaf, the single point in it
 T x0 = INF, x1 = -INF, y0 = INF, y1 = -INF; // bounds
 Node \star first = 0, \star second = 0;
```

```
T distance (const P& p) { // min squared distance to a point
    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).dist2();
  Node(vector \langle P \rangle \& vp \rangle: 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 >= v1 - v0 ? 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 \rightarrow pt) return \{INF, P()\};
      return make pair((p - node->pt).dist2(), node->pt);
    Node *f = node \rightarrow first, *s = node \rightarrow 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"

typedef Point<ll> P; typedef struct Quad\* Q; typedef int128 t lll; // (can be ll if coords are < 2e4) P arb (LLONG MAX, LLONG MAX); // not equal to any other point struct Quad { bool mark; Qo, rot; Pp;

# PolyhedronVolume Point3D 3dHull sphericalDistance

sort(all(pts)); assert(unique(all(pts)) == pts.end());

```
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 > 0;
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) \le 3) {
    Q a = makeEdge(s[0], s[1]), b = makeEdge(s[1], s.back());
    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->r() : a, side < 0 ? c : b->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->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\rightarrow r(), e\rightarrow r()\rightarrow 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) {
```

```
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 { Q c = e; do { c->mark = 1; pts.push_back(c->p); \
 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.
                                                       1ec4d3, 6 lines
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
long long.
                                                       8058ae, 32 lines
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) {}
  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, pi]
  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+y*y),z); }
 P unit() const { return *this/(T) dist(); } //makes dist()=1
  //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(angle), c = cos(angle); P u = axis.unit();
    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"

```
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[i] - 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[i];
      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[i];
#define C(a, b, c) if (E(a,b).cnt() != 2) mf(f.a, f.b, i, f.c);
      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

c172e9, 49 lines

**Description:** Returns the shortest distance on the sphere with radius radius between the points with azimuthal angles (longitude) f1  $(\phi_1)$  and f2  $(\phi_2)$  from x axis and zenith angles (latitude) t1  $(\theta_1)$  and t2  $(\theta_2)$  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. 8 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:  $\mathcal{O}(n)$ 

d4375c, 16 lines

```
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) {
  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:  $\mathcal{O}(N)$ 4bd552, 8 lines

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); break;}
  if (s[a+k] > s[b+k]) \{ a = b; break; \}
return a;
```

#### Suffix Array.h

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 = v, iota(all(sa), 0);
    for (int j = 0, p = 0; p < n; j = max(1, j * 2), lim = p) {
      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++;
    rep(i,1,n) rank[sa[i]] = i;
    for (int i = 0, j; i < n - 1; lcp[rank[i++]] = k)
     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:  $\mathcal{O}(26N)$ 

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, size of t);
    fill(t[1],t[1]+ALPHA,0);
    s[0] = 1; l[0] = l[1] = -1; r[0] = r[1] = p[0] = p[1] = 0;
    rep(i,0,sz(a)) ukkadd(i, toi(a[i]));
  // example: find longest common substring (uses ALPHA = 28)
  int lcs(int node, int i1, int i2, int olen) {
    if (|[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]) : 0;
    rep(c,0,ALPHA) if (t[node][c] != -1)
      mask |= lcs(t[node][c], i1, i2, len);
    if (mask == 3)
      best = \max(\text{best}, \{\text{len}, r[\text{node}] - \text{len}\}):
    return mask;
  static pii LCS(string s, string t) {
    SuffixTree st(s + (char)('z' + 1) + t + (char)('z' + 2));
    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 2^64).
// "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 + ~o.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 = (ll)1e11+3; // (order ~ 3e9; random also ok)
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)

# AhoCorasick DivideAndConquerDP MosAlgo

```
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.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  $\mathcal{O}(NM) \cdot_{716ac4, 67 \text{ lines}}$ 

```
struct AhoCorasick {
  enum {alpha = 26, first = 'A'};
  struct Node {
    // (nmatches is optional)
    int back, next[alpha], start = -1, end = -1, nmatches = 0;
    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].back = y;
          (N[ed].end == -1 ? N[ed].end : backp[N[ed].start])
           = N[y].end;
         N[ed].nmatches += N[v].nmatches;
          q.push(ed);
  vi find (string word) {
    int n = 0;
    vi res; // 11 count = 0;
    trav(c, word) {
     n = N[n].next[c - first];
```

```
res.push_back(N[n].end);
    // count += N[n].nmatches;
}
return res;
}
vector<vi> findAll(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;
}
```

# 10.1.1 Knuth

When doing DP on intervals:

when doing BI of the value  $a[i][j] = \min_{i < k < j} (a[i][k] + a[k][j]) + f(i,j)$ , where the (minimal) optimal k increases with both i and j, one can solve intervals in increasing order of length, and search k = p[i][j] for a[i][j] only between p[i][j-1] and p[i+1][j]. This is known as Knuth DP. Sufficient criteria for this are Monotonicity:  $f(b,c) \le f(a,d)$  and Quadrangle inequality:  $f(a,c) + f(b,d) \le f(a,d) + f(b,c)$  for all  $a \le b \le c \le d$ . 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) | 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

**Description:** Mo's on trees with path queries to print the smallest nonnegative integer on simple path  $a_i$  and  $b_i$ .

```
idx += (idx\&-idx);
int qry()
    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--;
   }
    return idx;
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:
    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)
    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++)</pre>
        scanf("%d%d%d", &u, &v, &x);
        u--, v--;
        graph[u].push back(\{v, x\}), graph[v].push back(\{u, x\});
    dfs0(0, -1, inf);
    //for(auto it: dis) cout << it << " ";
    //cout << "\n";
    for(int i = 0; i < q; i++)</pre>
        scanf("%d%d", &u, &v);
        u--. v--:
        if(start[u] > start[v]) swap(u, v);
        if(start[u] <= start[v] && start[v] <= en[u]) Q[i] = {{</pre>
             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 . second
             << " " << Q[i].second <math><< "\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 \leftarrow 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 < T1 >() ( p . first ) ^ hash < T2 >() ( p . second
     ) ;
unordered map < pair < int , int > , int , pair hash < int ,
     int > > M
```

# SIMD.h

Description: Cheat sheet of SSE/AVX intrinsics, for doing arithmetic on several numbers at once. Can provide a constant factor improvement of about 4, orthogonal to loop unrolling. Operations follow the pattern "\_mm(256)?\_name\_(si(128|256)|epi(8|16|32|64)|pd|ps)". Not all are described here; grep for \_mm\_ in /usr/lib/gcc/\*/4.9/include/ for more. If AVX is unsupported, try 128-bit operations, "emmintrin.h" and #define \_\_SSE\_\_ and \_\_MMX\_\_ before including it. For aligned memory use \_mm\_malloc(size, 32) or int buf[N] alignas(32), but prefer loadu/storeu

```
#pragma GCC target ("avx2") // or sse4.1
#include "immintrin.h"
typedef m256i mi;
#define \overline{L}(x) _mm256_loadu_si256((mi*)&(x))
// High-level/specific methods:
// load(u)? si256, store(u)? si256, setzero si256, mm malloc
// blendv (epi8|ps|pd) (z?y:x), movemask epi8 (hibits of bytes)
 // i32gather epi32(addr, x, 4): map addr lover 32-b parts of x
 / sad epu8: sum of absolute differences of u8, outputs 4xi64
  maddubs epi16: dot product of unsigned i7's, outputs 16xi15
  madd epi16: dot product of signed i16's, outputs 8xi32
  extractf128 si256(, i) (256->128), cvtsi128 si32 (128->lo32)
  permute2f128 si256(x,x,1) swaps 128-bit lanes
 shuffle epi3\overline{2}(x, 3*64+2*16+1*4+0) = x for each lane
// shuffle epi8(x, y) takes a vector instead of an imm
// Methods that work with most data types (append e.g. epi32):
// set1, blend (i8?x:y), add, adds (sat.), mullo, sub, and/or,
// andnot, abs, min, max, sign(1,x), cmp(gt|eq), unpack(lo|hi)
int sumi32(mi m) { union {int v[8]; mi m;} u; u.m = m;
 int ret = 0; rep(i,0,8) ret += u.v[i]; return ret; }
mi zero() { return mm256 setzero si256(); }
mi one() { return mm256 set1 epi32(-1); }
bool all zero (mi m) { return mm256 testz si256 (m, m); }
bool all one (mi m) { return mm256 testc si256(m, one()); }
ll example filteredDotProduct(int n, short* a, short* b) {
 int i = 0; ll r = 0;
 mi zero = mm256 setzero si256(), acc = zero;
 while (i + 16 <= n) {
   mi va = L(a[i]), vb = L(b[i]); i += 16;
   va = mm256 and si256( mm256 cmpgt epi16(vb, va), va);
   mi vp = mm256 madd epi16(va, vb);
    acc = mm256 add epi64( mm256 unpacklo epi32(vp, zero),
      mm256 add epi64(acc, mm256 unpackhi epi32(vp, zero)));
 union {ll v[4]; mi m;} u; u.m = acc; rep(i,0,4) r += u.v[i];
 for (; i < n; ++i) if (a[i] < b[i]) r += a[i] *b[i]; // <- equiv
 return r;
FastScanner.java
```

Description: Fast scanner class

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(); }

```
2
```

```
UCSD
                return st.nextToken();
       }
       int nextInt() { return Integer.parseInt(next()); }
long nextLong() { return Long.parseLong(next()); }
double nextDouble() { return Double.parseDouble(next()); }
```