

Birla Institute of Technology and Science, Pilani

SRS sus

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```
#include <bits/stdc++.h>
using namespace std;
#define rep(i, a, b) for(int i = a; i < (b); ++i)
#define all(x) begin(x), end(x)
#define sz(x) (int)(x).size()
#define endl '\n' //comment out for interactive problems
#define cout(x) (x?cout<<"YES"<<endl:cout<<"NO"<<endl)</pre>
#define arrPut(var) for (auto &inVar : var) {cin >> inVar;}
#define arrPrint(var) for (auto outVar : var) {cout << outVar
    << ' ';} cout << endl
// #define int long long
// #define double long double
const int INF =
#ifdef int
    LONG LONG MAX/2
#else
    INT_MAX/2
#endif
typedef long long 11;
typedef pair<int, int> pii;
typedef vector<int> vi;
signed main() {
  ios_base::sync_with_stdio(false);
    cin.tie(NULL); cout.tie(NULL);
    cout.precision(numeric_limits<double>::max_digits10);
    // freopen("input.txt","r",stdin);
    // freopen("output.txt","w", stdout);
```

.bashrc

alias c='q++ -Wall -Wconversion -Wfatal-errors -q -std=c++17 -fsanitize=undefined,address' xmodmap -e 'clear lock' -e 'keycode 66=less greater' $\#caps = \diamondsuit$

compileAndRun.txt

q++ -std=c++20 -DONLINE_JUDGE -O2 -x c++ {code} && ./a.out

Mathematics (2)

2.1 Equations

$$ax + by = e$$

$$cx + dy = f \Rightarrow x = \frac{ed - bf}{ad - bc}$$

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

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

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

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

2.2 Recurrences

If $a_n = c_1 a_{n-1} + \cdots + c_k a_{n-k}$, and r_1, \dots, 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. $a_n = (d_1 n + d_2)r^n.$

2.3 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

2.4.1 Triangles

Side lengths: a, b, c

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

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

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

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

Length of median (divides triangle into two equal-area triangles): $m_a = \frac{1}{2}\sqrt{2b^2 + 2c^2 - a^2}$

Length of bisector (divides angles in two):

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

Law of sines: $\frac{\sin \alpha}{a} = \frac{\sin \beta}{b} = \frac{\sin \gamma}{c} = \frac{1}{2R}$ Law of cosines: $a^2 = b^2 + c^2 - 2bc \cos \alpha$

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

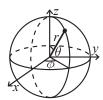
2.4.2 Quadrilaterals

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

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

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

2.4.3 Spherical coordinates



$$\begin{aligned} x &= r \sin \theta \cos \phi & r &= \sqrt{x^2 + y^2 + z^2} \\ y &= r \sin \theta \sin \phi & \theta &= \arccos(z/\sqrt{x^2 + y^2 + z^2}) \\ z &= r \cos \theta & \phi &= \operatorname{atan2}(y, x) \end{aligned}$$

Derivatives/Integrals

$$\frac{d}{dx}\arcsin x = \frac{1}{\sqrt{1-x^2}} \qquad \frac{d}{dx}\arccos x = -\frac{1}{\sqrt{1-x^2}}$$

$$\frac{d}{dx}\tan x = 1 + \tan^2 x \qquad \frac{d}{dx}\arctan x = \frac{1}{1+x^2}$$

$$\int \tan ax = -\frac{\ln|\cos ax|}{a} \qquad \int x\sin ax = \frac{\sin ax - ax\cos ax}{a^2}$$

$$\int e^{-x^2} = \frac{\sqrt{\pi}}{2}\operatorname{erf}(x) \qquad \int xe^{ax}dx = \frac{e^{ax}}{a^2}(ax-1)$$

Integration by parts:

$$\int_{a}^{b} f(x)g(x)dx = [F(x)g(x)]_{a}^{b} - \int_{a}^{b} F(x)g'(x)dx$$

2.6Sums

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

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

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

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

$$1^{4} + 2^{4} + 3^{4} + \dots + n^{4} = \frac{n(n+1)(2n+1)(3n^{2} + 3n - 1)}{20}$$

2.7 Series
$$e^{x} = 1 + x + \frac{x^{2}}{2!} + \frac{x^{3}}{3!} + \dots, (-\infty < x < \infty)$$

$$\ln(1+x) = x - \frac{x^{2}}{2} + \frac{x^{3}}{3} - \frac{x^{4}}{4} + \dots, (-1 < x \le 1)$$

$$\sqrt{1+x} = 1 + \frac{x}{2} - \frac{x^{2}}{8} + \frac{2x^{3}}{32} - \frac{5x^{4}}{128} + \dots, (-1 \le x \le 1)$$

$$\sin x = x - \frac{x^{3}}{3!} + \frac{x^{5}}{5!} - \frac{x^{7}}{7!} + \dots, (-\infty < x < \infty)$$

$$\cos x = 1 - \frac{x^{2}}{2!} + \frac{x^{4}}{4!} - \frac{x^{6}}{6!} + \dots, (-\infty < x < \infty)$$

Probability theory

2.8.1 Discrete distributions

Binomial distribution

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

$$p(k) = \binom{n}{k} p^k (1-p)^{n-k}$$
$$\mu = np, \ \sigma^2 = np(1-p)$$

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

First success distribution

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

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

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

Poisson distribution

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

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

2.8.2 Continuous distributions

Uniform distribution

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

$$f(x) = \begin{cases} \frac{1}{b-a} & a < x < b \\ 0 & \text{otherwise} \end{cases}$$
$$\mu = \frac{a+b}{2}, \ \sigma^2 = \frac{(b-a)^2}{12}$$

Exponential distribution

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

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

Normal distribution

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

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

If $X_1 \sim \mathcal{N}(\mu_1, \sigma_1^2)$ and $X_2 \sim \mathcal{N}(\mu_2, \sigma_2^2)$ then

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

2.9 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.

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

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

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

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

Data structures (3)

OrderStatisticTree.h

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

```
#include <bits/extc++.h>
using namespace gnu pbds;
template < class T>
using Tree = tree<T, null_type, less<T>, rb_tree_tag,
    tree_order_statistics_node_update>;
void example() {
 Tree<int> t, t2; t.insert(8);
  auto it = t.insert(10).first;
  assert(it == t.lower_bound(9));
  assert (t.order of key(10) == 1);
  assert(t.order_of_key(11) == 2);
  assert(*t.find by order(0) == 8);
 t.join(t2); // assuming T < T2 or T > T2, merge t2 into t
```

HashMap.h

Description: Hash map with mostly the same API as unordered_map, but ~3x faster. Uses 1.5x memory. Initial capacity must be a power of 2 (if provided). d77092, 7 lines

```
#include <bits/extc++.h>
// To use most bits rather than just the lowest ones:
struct chash { // large odd number for C
  const uint64 t C = 11(4e18 * acos(0)) | 71;
 11 operator()(11 x) const { return builtin bswap64(x*C); }
};
__gnu_pbds::gp_hash_table<11, int, chash> h({}, {}, {}, {}, {1<<16});
```

SegmentTree.h

Description: Zero-indexed max-tree. Bounds are inclusive to the left and exclusive to the right. Can be changed by modifying T, f and unit.

Time: $\mathcal{O}(\log N)$ 0f4bdb, 19 lines

```
struct Tree {
 typedef int T;
  static constexpr T unit = INT MIN;
 T f(T a, T b) { return max(a, b); } // (any associative fn)
  vector<T> s; int n;
  Tree(int n = 0, T def = unit) : s(2*n, def), n(n) {}
  void update(int pos, T val) {
    for (s[pos += n] = val; pos /= 2;)
      s[pos] = f(s[pos * 2], s[pos * 2 + 1]);
 T query (int b, int e) { // query [b, e)
    T ra = unit, rb = unit;
    for (b += n, e += n; b < e; b /= 2, e /= 2) {
      if (b % 2) ra = f(ra, s[b++]);
      if (e % 2) rb = f(s[--e], rb);
    return f(ra, rb);
};
```

RecursiveSegTree.h

```
Description: Uncomment for lazy.
                                                     db218d, 69 lines
template <typename T = int, typename V = T>
struct segtree{
   int n;
    vector<T> tree;
    // vector<V> lazy;
    T zero = 0; // Change according to QUERY operation
    // V lazy_zero = 0; // Change according to MODIFY operation
       n = sz;
        tree.clear(),tree.resize(2 * sz - 1, zero);;
        // lazy.clear(), lazy.resize(2 * sz - 1, lazy_zero);
   template <typename U>
    segtree(vector<U> &v) : segtree(v.size()){build(v);}
    T combine (T a, T b) { return a + b; } // Change according to
        QUERY operation
    template <typename U>
    void build(vector<U> &v, int id = 0, int seq1 = 0, int seqr
         = -1) {
        if (segr == -1)segr = n - 1;
       if (segl == segr) {
           tree[id] = v[seql]; // Change according to MODIFY
                 operation
            return:
        int mid = (segl + segr) / 2;
       build(v, id + 1, seql, mid);
       build(v, id + 2 * (mid - segl + 1), mid + 1, segr);
       tree[id] = combine(tree[id + 1], tree[id + 2 * (mid -
            segl + 1));
    /* void propagate(int id, int seql, int seqr){
        if(lazy / id) == lazy\_zero) return;
        if(seql != seqr)f
            int \ mid = (seql + seqr) / 2;
            array < int, 2 > children = {id + 1, id + 2 * (mid - 1)}
                 segl + 1);
            for(auto child : children) {
                tree[child] = lazy[id]; // Change according to
                      MODIFY operation
                lazy[child] = lazy[id]; // Change according to
                     MODIFY operation
        lazy[id] = lazy\_zero;
    template <typename U>
    void modify(U val, int index_l, int index_r, int id = 0,
        int segl = 0, int segr = -1) {
        if (segr == -1) segr = n - 1;
       if (index_l > index_r || index_l > seqr || seql >
            index r) {return; }
        // propagate(id, segl, segr);
       if (segl >= index_l && segr <= index_r){</pre>
            tree[id] = val; // Change according to MODIFY
            // lazy[id] = val; // Change according to MODIFY
                 operation
            return:
        int mid = (seql + seqr) / 2;
       modify(val, index_l, index_r, id + 1, segl, mid);
```

```
modify(val, index_1, index_r, id + 2 * (mid - seql + 1)
            , mid + 1, segr);
       tree[id] = combine(tree[id + 1], tree[id + 2 * (mid -
            segl + 1));
   T query(int index_l, int index_r, int id = 0, int segl = 0,
         int segr = -1) {
        if (segr == -1)segr = n - 1;
       if (index_l > index_r || index_l > seqr || seql >
             index_r) {return zero;}
        // propagate(id, segl, segr);
        if (segl >= index_1 && segr <= index_r) {return tree[id</pre>
            ];}
       int mid = (segl + segr) / 2;
       T leftVal = query(index_l, index_r, id + 1, seql, mid);
       T rightVal = query(index_l, index_r, id + 2 * (mid -
             segl + 1), mid + 1, segr);
       return combine (leftVal, rightVal);
};
PersistentDynamicLazySegTree.h
Description: Make the commented changes to remove persistence. O(q)
should be around 2e5 for persistent, seems like 1e6 gives memory problems
struct SegNode {
    long long freq = 0;
    int lazv = 0;
    int left = -1:
    int right = -1;
struct PersistentDynamicSegTree {
 vector<SegNode> tree;
 const int minl, maxr;
 int timer = 0;
    int lastroot = 0; // -
 long long comb(long long a, long long b) { return a + b; }
 void apply(int cur, int len, int val) {
       tree[cur].lazy = val;
       tree[cur].freq = (long long) len * val;
   // \ldots, int q = 0) .. if (q > 0) {tree.reserve(2 * q * _-lg(r))
         -l+1)); \}...
 PersistentDynamicSegTree(int 1, int r) : minl(l), maxr(r) {
      tree.push_back(SegNode());}
 void push_down(int cur, int 1, int r) {
    if (tree[cur].left == -1) {
     tree[cur].left = ++timer;
     tree.push_back(SegNode());
        else if (tree[cur].lazy != 0) {
           int newleft = ++timer;
            tree.push_back(tree[tree[cur].left]);
            tree[cur].left = newleft;
    if (tree[cur].right == -1) {
     tree[cur].right = ++timer;
      tree.push_back(SegNode());
        else if (tree[cur].lazy != 0) {
            int newright = ++timer;
            tree.push_back(tree[tree[cur].right]); // -
            tree[cur].right = newright;
```

```
if(tree[cur].lazy == 0) {return;}
    int m = 1 + (r - 1 >> 1);
    apply(tree[cur].left, m - 1 + 1, tree[cur].lazy);
    apply(tree[cur].right, r - m, tree[cur].lazy);
    tree[cur].lazy = 0;
    /\!/void\ modify\ \dots\ ,\ int\ cur = 0,\ \dots
  int modify(int val, int index_l, int index_r, int cur = -1,
       int 1 = 1, int r = -1) {
        if(cur == -1) cur = lastroot; // -
        if (1 == 1 \text{ and } r == -1) 1 = min1, r = maxr;
    if (index_r < 1 || index_l > r) { return cur; } // { return
        int newcur = cur; // -
    if (index_l <= l && r <= index_r) {</pre>
            newcur = ++timer;
            tree.push_back(tree[cur]); // -
      apply(newcur, r - 1 + 1, val); // replace newcur with cur
    } else {
      push_down(cur, 1, r);
      int m = 1 + (r - 1 >> 1);
            newcur = ++timer;
            tree.push_back(tree[cur]); // -
      tree[newcur].left = modify(val, index_1, index_r, tree[
           cur].left, 1, m);
                                    // remove lhs
      tree[newcur].right = modify(val, index_l, index_r, tree[
           cur].right, m + 1, r); //remove lhs
      tree[newcur].freq = // replace newcur with cur
          comb(tree[tree[newcur].left].freq, tree[tree[newcur].
               right].freq); // replace newcur with cur
        return lastroot = newcur;
    // .. int cur = 0, ..
  long long query (int index 1, int index r, int cur = -1, int 1
        = 1, int r = -1) {
        if(cur == -1) cur = lastroot; // -
        if (1 == 1 \text{ and } r == -1) 1 = minl, r = maxr;
    if (index_r < 1 || index_l > r) { return 0; }
    if (index 1 <= 1 && r <= index r) { return tree[cur].freq;</pre>
    push_down(cur, 1, r);
    int m = 1 + (r - 1 >> 1);
    return comb(query(index_1, index_r, tree[cur].left, 1, m),
                query(index 1, index r, tree[cur].right, m + 1,
    // remove function
    int load_root(int cur){
        lastroot = ++timer;
        tree.push_back(tree[cur]);
        return lastroot;
};
Segtree2d.h
Description: 2d segtree (params are inclusive)
                                                     7322a3, 106 lines
struct segtree2d
    vector<vector<int>>& ar;
    vector<vector<int>> tree;
    int n, m;
```

UnionFindRollback SubMatrix Matrix LineContainer

```
void buildcolumn (int idi, int idj, int segil, int segir,
    int segjl, int segjr)
    if(segjl == segjr)
        if(segil == segir)
            tree[idi][idj] = ar[seqil][seqjl];
        else
            tree[idi][idj] = tree[2*idi + 1][idj] + tree[2*idi + 1][idj]
                 idi + 2][idj];
    el se
        int segjmid = segjl + segjr >> 1;
        buildcolumn(idi,2*idj + 1, segil, segir, segil,
             segimid);
        buildcolumn(idi,2*idj + 2, segil, segir, segjmid +
             1, segjr);
        tree[idi][idj] = tree[idi][2*idj + 1] + tree[idi
             [2*idj + 2];
void buildrow(int idi, int seqil, int seqir)
    if(segil != segir)
        int segimid = segil + segir >> 1;
        buildrow(2*idi + 1, segil, segimid);
        buildrow(2*idi + 2, segimid + 1, segir);
   buildcolumn(idi,0,segil,segir,0,m-1);
segtree2d(vector<vector<int>>& ar) : ar(ar)
   n = ar.size();
   m = ar[0].size();
   tree.resize(4 * n, vector<int>(4 * m));
   buildrow (0,0,n-1);
int querycolumn(int jl, int jr, int idi, int idj, int seqjl
     , int segir)
    if (min(segjr, jr) < max(segjl, jl)) return 0;</pre>
   if(segil >= il and segir <= ir)return tree[idi][idi];</pre>
   int segjmid = segjl + segjr >> 1;
    return querycolumn(jl,jr,idi,2*idj + 1, segjl, segjmid)
          + querycolumn(jl,jr,idi,2*idj + 2, segjmid + 1,
         segjr);
int queryrow(int il, int ir, int jl, int jr, int idi = 0,
    int segil = 0, int segir = -1)
   if(segir == -1)segir += n;
   if (min (segir, ir) < max(segil, il)) return 0;</pre>
   if (segil >= il and segir <= ir) return querycolumn(jl,</pre>
         jr,idi,0,0,m-1);
    int segimid = segil + segir >> 1;
```

```
return queryrow(il,ir,jl,jr,2*idi + 1,segil,segimid) +
             queryrow(il,ir,jl,jr,2*idi + 2,segimid + 1,segir);
    void modifycolumn(int i, int j, int idi, int idj, int segil
         , int segir, int segjl, int segjr)
        if(min(segjr,j) < max(segjl,j))return;</pre>
        if(segjl == segjr)
            if(segil == segir)
                assert(i == segil and j == segjl);
                ar[i][j] ^= 1;
                tree[idi][idj] = ar[i][j];
            else
                tree[idi][idj] = tree[2*idi + 1][idj] + tree[2*idi + 1][idj]
                     idi + 2][idj];
        else
            int segjmid = segjl + segjr >> 1;
            modifycolumn(i,j,idi,2*idj + 1, segil,segir, segjl,
            modifycolumn(i,j,idi,2*idj + 2,segil,segir,segjmid
                  + 1, segjr);
            tree[idi][idj] = tree[idi][2*idj + 1] + tree[idi
                 ][2*idi + 2];
   }
   void modifyrow(int i, int j, int idi = 0, int seqil = 0,
         int segir = -1)
        if(segir == -1)segir += n;
        if (min(segir,i) < max(segil,i)) return;</pre>
        if(segil != segir)
            int segimid = segil + segir >> 1;
            modifyrow(i, j, 2*idi + 1, seqil, seqimid);
            modifyrow(i, j, 2*idi + 2, segimid + 1, segir);
        modifycolumn(i,j,idi,0,segil,segir,0,m-1);
};
UnionFindRollback.h
Description: Disjoint-set data structure with undo. If undo is not needed,
skip st, time() and rollback().
Usage: int t = uf.time(); ...; uf.rollback(t);
Time: \mathcal{O}(\log(N))
                                                      de4ad0, 21 lines
struct RollbackUF
 vi e; vector<pii> st;
 RollbackUF(int n) : e(n, -1) {}
 int size(int x) { return -e[find(x)]; }
 int find(int x) { return e[x] < 0 ? x : find(e[x]); }</pre>
 int time() { return sz(st); }
 void rollback(int t) {
    for (int i = time(); i --> t;)
     e[st[i].first] = st[i].second;
   st.resize(t);
 bool join(int a, int b) {
```

a = find(a), b = find(b);

```
if (a == b) return false;
    if (e[a] > e[b]) swap(a, b);
    st.push back({a, e[a]});
    st.push_back({b, e[b]});
    e[a] += e[b]; e[b] = a;
    return true;
};
SubMatrix.h
Description: Calculate submatrix sums quickly, given upper-left and lower-
right corners (half-open).
Usage: SubMatrix<int> m (matrix);
m.sum(0, 0, 2, 2); // top left 4 elements
Time: \mathcal{O}(N^2+Q)
                                                        c59ada, 13 lines
template<class T>
struct SubMatrix {
  vector<vector<T>> p:
  SubMatrix(vector<vector<T>>& v) {
    int R = sz(v), C = sz(v[0]);
    p.assign(R+1, vector<T>(C+1));
    rep(r, 0, R) rep(c, 0, C)
      p[r+1][c+1] = v[r][c] + p[r][c+1] + p[r+1][c] - p[r][c];
  T sum(int u, int 1, int d, int r) {
    return p[d][r] - p[d][l] - p[u][r] + p[u][l];
};
Matrix.h
Description: Basic operations on square matrices.
Usage: Matrix<int, 3> A;
A.d = \{\{\{\{1,2,3\}\}\}, \{\{4,5,6\}\}, \{\{7,8,9\}\}\}\};
array<int, 3 > \text{vec} = \{1, 2, 3\};
vec = (A^N) * vec;
                                                       6ab5db, 26 lines
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];
    return a;
  array<T, N> operator*(const array<T, N>& vec) const {
    array<T, N> ret{};
    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);
    M a, b(*this);
    rep(i, 0, N) \ a.d[i][i] = 1;
    while (p) {
      if (p&1) a = a*b;
      b = b*b;
      p >>= 1;
    return a;
};
```

LineContainer.h

Description: Container where you can add lines of the form kx+m, and query maximum values at points x. Useful for dynamic programming ("convex hull trick").

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

8ec1c7, 30 lines

Treap FenwickTree FenwickTree2d SparseTable

```
struct Line {
  mutable 11 k, m, p;
  bool operator<(const Line& o) const { return k < o.k; }</pre>
 bool operator<(11 x) const { return p < x; }</pre>
struct LineContainer : multiset<Line, less<>>> {
  // (for doubles, use inf = 1/.0, div(a,b) = a/b)
  static const 11 inf = LLONG MAX;
  ll div(ll a, ll b) { // floored division
    return a / b - ((a ^ b) < 0 && a % b); }
  bool isect(iterator x, iterator y) {
    if (y == end()) return x \rightarrow p = inf, 0;
    if (x->k == y->k) x->p = x->m > y->m ? inf : -inf;
    else x->p = div(y->m - x->m, x->k - y->k);
    return x->p >= y->p;
  void add(ll k, ll m) {
    auto z = insert(\{k, m, 0\}), y = z++, x = y;
    while (isect(y, z)) z = erase(z);
    if (x != begin() \&\& isect(--x, y)) isect(x, y = erase(y));
    while ((y = x) != begin() && (--x)->p >= y->p)
      isect(x, erase(y));
  11 query(ll x) {
    assert(!empty());
   auto 1 = *lower_bound(x);
   return 1.k * x + 1.m;
};
```

Treap.h

Description: A short self-balancing tree. It acts as a sequential container with log-time splits/joins, and is easy to augment with additional data. Time: $\mathcal{O}(\log N)$

53871e, 63 lines

```
const int M = 1e9 + 7;
typedef struct item * pitem;
struct item {
   int prior, cnt, value, b = 1, c = 0, sum = 0;
   pitem 1, r;
   bool rev = false;
   item(int val): value(val), prior(rand()), cnt(0), l(NULL),
        r(NULL) {}
int cnt(pitem it) { return it ? it->cnt : 0; }
int giveSum(pitem it) { return it ? it->sum : 0; }
void combineLazy(pitem t, int b, int c) {
    t->value = ((t->value * b) % M + c) % M;
    t->b = (t->b * b) % M;
   t->c = ((t->c * b) + c) % M;
void push(pitem it) {
    if(it && it->rev) {
       it->rev = false;
       swap(it->l, it->r);
       if(it->1) it->1->rev ^= 1;
       if(it->r) it->r->rev ^= 1;
   if(it) {
       if(it->1) combineLazy(it->1, it->b, it->c);
       if(it->r) combineLazy(it->r, it->b, it->c);
       it->sum = ((it->sum * it->b) % M + (it->cnt * it->c) %
            M) % M;
        it->b = 1, it->c = 0;
```

```
void pull(pitem it) {
  if (!it) { return; }
  push(it->1), push(it->r);
  it->cnt = cnt(it->1) + cnt(it->r) + 1;
  it->sum = (giveSum(it->1) + giveSum(it->r) + it->value) % M;
void merge(pitem &t, pitem 1, pitem r) {
    push(1), push(r);
    if(!1 || !r) t = 1 ? 1 : r;
    else if (1->prior > r->prior) merge (1->r, 1->r, r), t = 1;
    else merge (r->1, 1, r->1), t = r;
    pull(t);
void split(pitem t, pitem &1, pitem &r, int key, int add = 0) {
    if(!t) return void(1 = r = 0);
    push(t);
    int curr_key = add + cnt(t->1);
    if(key <= curr_key) split(t->1, 1, t->1, key, add), r = t;
    else split(t->r, t->r, r, key, curr_key + 1), 1 = t;
    pull(t);
void insert(pitem &t, pitem new_item, int pos) {
    pitem t1 = NULL, t2 = NULL;
    split(t, t1, t2, pos);
    merge(t1, t1, new_item);
    merge(t, t1, t2);
void remove(pitem &t, int pos) {
   pitem t1 = NULL, t2 = NULL, t3 = NULL;
    split(t, t1, t3, pos);
    split(t3, t2, t3, 1);
    merge(t, t1, t3);
```

FenwickTree.h

Description: Computes partial sums a[0] + a[1] + ... + a[pos - 1], and updates single elements a[i], taking the difference between the old and new

Time: Both operations are $\mathcal{O}(\log N)$.

```
e62fac, 22 lines
```

```
struct FT {
 vector<ll> s;
 FT(int n) : s(n) {}
 void update(int pos, ll dif) { // a[pos] \neq dif
    for (; pos < sz(s); pos |= pos + 1) s[pos] += dif;</pre>
 11 query (int pos) { // sum of values in [0, pos)
   11 \text{ res} = 0;
    for (; pos > 0; pos &= pos - 1) res += s[pos-1];
    return res;
 int lower_bound(ll sum) \{// min \ pos \ st \ sum \ of \ [0, \ pos] >= sum
    // Returns n if no sum is >= sum, or -1 if empty sum is.
    if (sum <= 0) return -1;
   int pos = 0;
    for (int pw = 1 << 25; pw; pw >>= 1) {
     if (pos + pw <= sz(s) && s[pos + pw-1] < sum)
        pos += pw, sum -= s[pos-1];
   return pos;
```

FenwickTree2d.h

Description: Computes sums a[i,j] for all i<I, j<J, and increases single elements 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"
                                                                               157f07, 22 lines
```

```
struct FT2 {
  vector<vi> vs; vector<FT> ft;
  FT2(int limx) : ys(limx) {}
  void fakeUpdate(int x, int y) {
    for (; x < sz(ys); x |= x + 1) ys[x].push_back(y);
  void init() {
    for (vi& v : ys) sort(all(v)), ft.emplace_back(sz(v));
  int ind(int x, int y) {
    return (int)(lower_bound(all(ys[x]), y) - ys[x].begin()); }
  void update(int x, int y, ll dif) {
    for (; x < sz(ys); x | = x + 1)
      ft[x].update(ind(x, y), dif);
  11 query(int x, int y) {
    11 \text{ sum} = 0;
    for (; x; x &= x - 1)
      sum += ft[x-1].query(ind(x-1, y));
    return sum;
};
SparseTable.h
Description: Min/Max queries on static array. both l,r is inclusive 1540b1, 38 lines
struct SparseTable
    vector<int>& a;
```

vector<vector<int>> table; bool compare (auto a, auto b) { return a < b; } // Change according to QUERY operation int identity = INF; // Change according to QUERY operation SparseTable(vector<int>& a) : a(a), n(a.size()) int lgN = (int)bit_width((unsigned int)n) - 1; table.resize(lqN + 1, vector<int>(n)); range(i, 0, lgN + 1) range(j, 0, n - (1 << i) + 1) **if**(i == 0) table[i][j] = j;else if(_compare (a[table[i-1][j]] , a[table[i-1][j + (1<<i-1)||)) table[i][j] = table[i-1][j];else table[i][j] = table[i-1][j + (1 << i-1)];int query_i(int 1, int r) assert($1 \le r$ and $1 \le n$ and $r \ge 0$); $1 = \max(1, (int)0), r = \min(r, n - 1);$ int lgN = (int)bit_width((unsigned int)(r - 1 + 1)) if (_compare (a[table[lgN][l]] , a[table[lgN][r - (1<<</pre> lgN) + 1]]))return table[lqN][l]; **return** table[lqN][r - (1<<lqN) + 1];

int query_val(int 1, int r) { return $(1 > r \mid | 1 >= n \mid | r$

< 0) ? identity : a[query_i(l,r)]; }

};

MoQueries.h

Description: Answer interval or tree path queries by finding an approximate TSP through the queries, and moving from one query to the next by adding/removing points at the ends. If values are on tree edges, change step to add/remove the edge (a, c) and remove the initial add call (but keep in). Time: $\mathcal{O}(N\sqrt{Q})$

```
void add(int ind, int end) { ... } // add a[ind] (end = 0 or 1)
void del(int ind, int end) { ... } // remove a [ind]
int calc() { ... } // compute current answer
vi mo(vector<pii> Q) {
 int L = 0, R = 0, blk = 350; // \sim N/sqrt(Q)
 vi s(sz(Q)), res = s;
#define K(x) pii(x.first/blk, x.second ^ -(x.first/blk & 1))
  iota(all(s), 0);
  sort(all(s), [\&](int s, int t){ return K(Q[s]) < K(Q[t]); });
  for (int qi : s) {
   pii q = Q[qi];
   while (L > q.first) add(--L, 0);
   while (R < q.second) add(R++, 1);</pre>
   while (L < q.first) del(L++, 0);
   while (R > q.second) del(--R, 1);
   res[qi] = calc();
 return res;
vi moTree(vector<array<int, 2>> Q, vector<vi>& ed, int root=0) {
 int N = sz(ed), pos[2] = {}, blk = 350; // \sim N/sqrt(Q)
  vi s(sz(Q)), res = s, I(N), L(N), R(N), in(N), par(N);
  add(0, 0), in[0] = 1;
  auto dfs = [&] (int x, int p, int dep, auto& f) -> void {
   par[x] = p;
   L[x] = N;
   if (dep) I[x] = N++;
   for (int y : ed[x]) if (y != p) f(y, x, !dep, f);
   if (!dep) I[x] = N++;
   R[x] = N;
  dfs(root, -1, 0, dfs);
#define K(x) pii(I[x[0]] / blk, I[x[1]] ^ -(I[x[0]] / blk & 1))
  iota(all(s), 0);
  sort(all(s), [&](int s, int t) { return K(Q[s]) < K(Q[t]); });
  for (int qi : s) rep(end, 0, 2) {
   int &a = pos[end], b = O[qi][end], i = 0;
#define step(c) { if (in[c]) { del(a, end); in[a] = 0; } \
                  else { add(c, end); in[c] = 1; } a = c; }
    while (!(L[b] <= L[a] && R[a] <= R[b]))
     I[i++] = b, b = par[b];
    while (a != b) step(par[a]);
   while (i--) step(I[i]);
   if (end) res[qi] = calc();
  return res:
```

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: polyRoots(\{\{2, -3, 1\}\}, -1e9, 1e9\}) // solve x^2-3x+2=0
Time: \mathcal{O}\left(n^2\log(1/\epsilon)\right)
"Polynomial.h"
vector<double> polyRoots(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 = polyRoots(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 ^{\circ} (p(h) > 0)) {
      rep(it, 0, 60) { // while (h - l > 1e-8)
        double m = (1 + h) / 2, f = p(m);
        if ((f \le 0) ^ sign) 1 = m;
        else h = m;
      ret.push_back((1 + h) / 2);
  return ret;
```

PolyInterpolate.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 bruteforcing 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"
vector<ll> berlekampMassey(vector<ll> s) {
 int n = sz(s), L = 0, m = 0;
 vector<ll> C(n), B(n), T;
 C[0] = B[0] = 1;
  11 b = 1;
  rep(i,0,n) \{ ++m;
    11 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());
  for (11& 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)
typedef vector<ll> Poly;
```

```
11 linearRec(Poly S, Poly tr, 11 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);
 rep(i, 0, n) res = (res + pol[i + 1] * S[i]) % mod;
 return res;
```

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 and no local maximum. The maximum error in the result is eps. Works equally well for maximization with a small change in the code. See Ternary-Search, 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_{8eeeaf, 14 lines}

```
typedef array<double, 2> P;

template<class F> pair<double, P> hillClimb(P start, F f) {
  pair<double, P> cur(f(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(f(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 $x*x + y*y + z*z < 1; });});});$

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

template <class F>
d rec(F& f, d a, d b, d eps, d S) {
    d c = (a + b) / 2;
    d S1 = S(a, c), S2 = S(c, b), T = S1 + S2;
```

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

Simplex.h

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

```
Usage: vvd A = \{\{1,-1\}, \{-1,1\}, \{-1,-2\}\}; vd b = \{1,1,-4\}, c = \{-1,-1\}, x; T val = LPSolver(A, b, c).solve(x);
```

const T eps = 1e-8, inf = 1/.0;

Time: $\mathcal{O}(NM * \#pivots)$, where a pivot may be e.g. an edge relaxation. $\mathcal{O}(2^n)$ in the general case.

```
typedef double T; // long double, Rational, double + mod<P>...
typedef vector<T> vd;
typedef vector<vd> vvd;
```

```
#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;</pre>
```

```
void pivot(int r, int s) {
  T \star a = D[r].data(), inv = 1 / a[s];
  rep(i, 0, m+2) if (i != r \&\& abs(D[i][s]) > eps) {
    T *b = D[i].data(), inv2 = b[s] * inv;
    rep(j,0,n+2) b[j] -= a[j] * inv2;
    b[s] = a[s] * inv2;
  rep(j, 0, n+2) if (j != s) D[r][j] *= inv;
  rep(i,0,m+2) if (i != r) D[i][s] \star = -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;
      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;
}
</pre>
```

4.3 Matrices

Determinant.h

Description: Calculates determinant of a matrix. Destroys the matrix. Time: $\mathcal{O}\left(N^3\right)$ 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 11 mod = 12345;
11 det(vector<vector<ll>>& a) {
  int n = sz(a); 11 ans = 1;
  rep(i,0,n) {
    rep(j,i+1,n) {
      while (a[j][i] != 0) { // gcd step}
        11 t = a[i][i] / a[j][i];
        if (t) rep(k,i,n)
          a[i][k] = (a[i][k] - a[j][k] * t) % mod;
        swap(a[i], a[j]);
        ans \star = -1;
    ans = ans * 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: $\mathcal{O}\left(n^2m\right)$

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];

SolveLinear2.h

Description: To get all uniquely determined values of x back from Solve-Linear, make the following changes:

return rank; // (multiple solutions if rank < m)

```
"SolveLinear.h"
                                                       08e495, 7 lines
rep(j,0,n) if (j != i) // instead of rep(j,i+1,n)
// ... then at the end:
x.assign(m, undefined);
rep(i,0,rank) {
 rep(j,rank,m) if (fabs(A[i][j]) > eps) goto fail;
 x[col[i]] = b[i] / A[i][i];
fail:; }
```

SolveLinearBinary.h

Description: Solves Ax = b over \mathbb{F}_2 . If there are multiple solutions, one is returned arbitrarily. Returns rank, or -1 if no solutions. Destroys A and b. Time: $\mathcal{O}\left(n^2m\right)$ fa2d7a, 34 lines

```
typedef bitset<1000> bs;
int solveLinear(vector<bs>& A, vi& b, bs& x, int m) {
 int n = sz(A), rank = 0, br;
  assert(m \le sz(x));
  vi col(m); iota(all(col), 0);
  rep(i,0,n) {
   for (br=i; br<n; ++br) if (A[br].any()) break;</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

return n;

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

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

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

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

This is useful for solving problems on the type

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

where a_0, a_{n+1}, b_i, c_i and d_i are known. a can then be obtained from

$$\{a_i\} = \operatorname{tridiagonal}(\{1, -1, -1, \dots, -1, 1\}, \{0, c_1, c_2, \dots, c_n\}, \\ \{b_1, b_2, \dots, b_n, 0\}, \{a_0, d_1, d_2, \dots, d_n, a_{n+1}\}).$$

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];</pre>
      diag[i+1] = sub[i]; tr[++i] = 1;
     else {
      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. N must be a power of 2. 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 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) {
```

```
int n = sz(a), L = 31 - __builtin_clz(n);
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, acos(-1.0L) / k);
```

```
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 << L;</pre>
  vector<C> in(n), out(n);
  copy(all(a), begin(in));
  rep(i,0,sz(b)) in[i].imag(b[i]);
  fft(in);
  for (C& x : in) x *= x;
  rep(i, 0, n) out[i] = in[-i & (n - 1)] - conj(in[i]);
  rep(i, 0, sz(res)) res[i] = imag(out[i]) / (4 * n);
  return res;
```

FastFourierTransformMod.h

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

Time: $\mathcal{O}(N \log N)$, where N = |A| + |B| (twice as slow as NTT or FFT) "FastFourierTransform.h" b82773, 22 lines

```
typedef vector<ll> v1;
template<int M> vl convMod(const vl &a, const vl &b) {
  if (a.empty() || b.empty()) return {};
  vl res(sz(a) + sz(b) - 1);
  int B=32-__builtin_clz(sz(res)), n=1<<B, cut=int(sqrt(M));</pre>
  vector<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)) {
    11 \text{ av} = 11(\text{real}(\text{outl}[i]) + .5), \text{ cv} = 11(\text{imag}(\text{outs}[i]) + .5);
    11 \text{ bv} = 11(\text{imag}(\text{outl}[i]) + .5) + 11(\text{real}(\text{outs}[i]) + .5);
    res[i] = ((av % M * cut + bv) % M * cut + cv) % M;
  return res:
```

NumberTheoreticTransform.h

Description: ntt(a) computes $\hat{f}(k) = \sum_{x} a[x]g^{xk}$ for all k, where $g = \sum_{x} a[x]g^{xk}$ $root^{(mod-1)/N}$. N must be a power of 2. Useful for convolution modulo specific nice primes of the form $2^a b + 1$, where the convolution result has size at most $2^{\hat{a}}$. For arbitrary modulo, see FFTMod. conv(a, b) = c, where $c[x] = \sum a[i]b[x-i]$. For manual convolution: NTT the inputs, multiply pointwise, divide by n, reverse(start+1, end), NTT back. Inputs must be in [0, mod).

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

```
"../number-theory/ModPow.h"
                                                                        ced03d, 35 lines
```

```
const 11 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> v1;
void ntt(vl &a) {
 int n = sz(a), L = 31 - __builtin_clz(n);
  static v1 rt(2, 1);
  for (static int k = 2, s = 2; k < n; k *= 2, s++) {
    rt.resize(n);
   ll z[] = \{1, modpow(root, mod >> s)\};
   rep(i,k,2*k) rt[i] = rt[i / 2] * z[i & 1] % mod;
 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]]);</pre>
  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] = ai - z + (z > ai ? mod : 0);
     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;
  int inv = modpow(n, mod - 2);
 vl L(a), R(b), out(n);
 L.resize(n), R.resize(n);
  ntt(L), ntt(R);
  rep(i,0,n)
    out[-i \& (n - 1)] = (l1)L[i] * R[i] % mod * inv % mod;
  return {out.begin(), out.begin() + s};
```

FastSubsetTransform.h

Description: Transform to a basis with fast convolutions of the form $c[z] = \sum_{z = x \oplus y} a[x] \cdot b[y], \text{ where } \oplus \text{ is one of AND, OR, XOR.}$ The size of a must be a power of two.

```
Time: \mathcal{O}(N \log N)
                                                      464cf3, 16 lines
void FST(vi& a, bool inv) {
 for (int n = sz(a), step = 1; step < n; step *= 2) {
   for (int i = 0; i < n; i += 2 * step) rep(i, i, i+step) {
      int \&u = a[j], \&v = a[j + step]; tie(u, v) =
        inv ? pii(v - u, u) : pii(v, u + v); // AND
        inv ? pii(v, u - v) : pii(u + v, u); // OR
                                               // XOR
       pii(u + v, u - v);
 if (inv) for (int& x : a) x /= sz(a); // XOR only
vi conv(vi a, vi b) {
 FST(a, 0); FST(b, 0);
 rep(i, 0, sz(a)) a[i] *= b[i];
 FST(a, 1); return a;
```

Number theory (5)

5.1 Modular arithmetic

Modular Arithmetic.h

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

```
"euclid.h"
                                                          35bfea, 18 lines
const 11 mod = 17; // change to something else
struct Mod {
```

```
11 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^(11 e) {
    if (!e) return Mod(1);
    Mod r = *this ^ (e / 2); r = r * r;
    return e&1 ? *this * r : r;
};
ModInverse.h
```

Description: Pre-computation of modular inverses. Assumes LIM < mod and that mod is a prime. 6f684f, 3 lines

```
const 11 mod = 1000000007, LIM = 200000:
11* inv = new 11[LIM] - 1; inv[1] = 1;
rep(i,2,LIM) inv[i] = mod - (mod / i) * inv[mod % i] % mod;
```

ModPow.h

b83e45, 8 lines

```
const 11 mod = 1000000007; // faster if const
11 modpow(11 b, 11 e) {
 11 \text{ ans} = 1;
 for (; e; b = b * b % mod, e /= 2)
   if (e & 1) ans = ans * b % mod;
```

ModLog.h

Description: Returns the smallest x > 0 s.t. $a^x = b \pmod{m}$, or -1 if no such x exists, $\operatorname{modLog}(a,1,m)$ can be used to calculate the order of a. Time: $\mathcal{O}\left(\sqrt{m}\right)$

```
11 modLog(ll a, ll b, ll m) {
 11 n = (11)   sqrt(m) + 1, e = 1, f = 1, j = 1;
 unordered_map<11, 11> A;
  while (j \le n \&\& (e = f = e * a % m) != b % m)
   A[e * b % m] = j++;
  if (e == b % m) return j;
 if (__gcd(m, e) == __gcd(m, b))
    rep(i,2,n+2) if (A.count(e = e * f % m))
      return n * i - A[e];
  return -1;
```

ModSum.h

Description: Sums of mod'ed arithmetic progressions.

modsum(to, c, k, m) = $\sum_{i=0}^{\text{to}-1} (ki+c)\%m$. divsum is similar but for floored division.

Time: $\log(m)$, with a large constant.

5c5bc5, 16 lines

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

```
11 modsum(ull to, 11 c, 11 k, 11 m) {
  c = ((c % m) + m) % m;
  k = ((k % m) + m) % m;
  return to * c + k * sumsq(to) - m * divsum(to, c, k, m);
}
```

ModMulLL.h

Description: Calculate $a \cdot b \mod c$ (or $a^b \mod c$) for $0 \le a, b \le c \le 7.2 \cdot 10^{18}$. **Time:** $\mathcal{O}(1)$ for modmul, $\mathcal{O}(\log b)$ for modpow

```
typedef unsigned long long ull;
ull modmul(ull a, ull b, ull M) {
    ll ret = a * b - M * ull(1.L / M * a * b);
    return ret + M * (ret < 0) - M * (ret >= (ll)M);
}
ull modpow(ull b, ull e, ull mod) {
    ull ans = 1;
    for (; e; b = modmul(b, b, mod), e /= 2)
        if (e & 1) ans = modmul(ans, b, mod);
    return ans;
}
```

ModSqrt.h

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

Time: $\mathcal{O}\left(\log^2 p\right)$ worst case, $\mathcal{O}\left(\log p\right)$ for most p

```
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} \text{ or } 2^{(n+3)/8} * 2^{(n-1)/4} \text{ works if } p \% 8 == 5
  11 s = p - 1, n = 2;
  int r = 0, m;
  while (s % 2 == 0)
   ++r, s /= 2;
  while (modpow(n, (p-1) / 2, p) != p-1) ++n;
  11 x = modpow(a, (s + 1) / 2, p);
 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;
   11 \text{ gs} = \text{modpow}(q, 1LL << (r - m - 1), p);
    q = qs * qs % p;
   x = x * gs % p;
   b = b * g % p;
```

5.2 Primality

FastEratosthenes.h

Description: Prime sieve for generating all primes smaller than LIM. **Time:** LIM=1e9 ≈ 1.5 s $_{6b2912,\ 20\ lines}$

```
const int LIM = 1e6;
bitset<LIM> isPrime;
vi eratosthenes() {
  const int S = (int)round(sqrt(LIM)), R = LIM / 2;
  vi pr = {2}, sieve(S+1); pr.reserve(int(LIM/log(LIM)*1.1));
  vector<pii> cp;
  for (int i = 3; i <= S; i += 2) if (!sieve[i]) {
    cp.push_back({i, i * i / 2});
  for (int j = i * i; j <= S; j += 2 * i) sieve[j] = 1;</pre>
```

```
for (int L = 1; L <= R; L += S) {
    array<bool, S> block{};
    for (auto & [p, idx] : cp)
        for (int i=idx; i < S+L; idx = (i+=p)) block[i-L] = 1;
    rep(i,0,min(S, R - L))
        if (!block[i]) pr.push_back((L + i) * 2 + 1);
}
for (int i : pr) isPrime[i] = 1;
return pr;</pre>
```

MillerRabin.h

Description: Deterministic Miller-Rabin primality test. Guaranteed to work for numbers up to $7\cdot 10^{18}$; for larger numbers, use Python and extend A randomly.

Time: 7 times the complexity of $a^b \mod c$.
"ModMullL.h"

```
"ModMulLL.h" 60dcd1, 12 lines
bool isPrime(ull n) {
   if (n < 2 | | n % 6 % 4 != 1) return (n | 1) == 3;
   ull A[] = {2, 325, 9375, 28178, 450775, 9780504, 1795265022},
        s = __builtin_ctzl1(n-1), d = n >> s;
   for (ull a : A) { // ^ count trailing zeroes
        ull p = modpow(a%n, d, n), i = s;
        while (p != 1 && p != n - 1 && a % n && i--)
        p = modmul(p, p, n);
   if (p != n-1 && i != s) return 0;
   }
   return 1;
}
```

Factor.h

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

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

```
"ModMulLL.h", "MillerRabin.h"
                                                     d8d98d, 18 lines
ull pollard(ull n) {
 ull x = 0, y = 0, t = 30, prd = 2, i = 1, q;
 auto f = [\&](ull x) \{ return modmul(x, x, n) + i; \};
 while (t++ % 40 || __gcd(prd, n) == 1) {
   if (x == y) x = ++i, y = f(x);
   if ((q = modmul(prd, max(x,y) - min(x,y), n))) prd = q;
   x = f(x), y = f(f(y));
 return __gcd(prd, n);
vector<ull> factor(ull n) {
 if (n == 1) return {};
 if (isPrime(n)) return {n};
 ull x = pollard(n);
 auto 1 = factor(x), r = factor(n / x);
 1.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 $a \pmod{b}$.

```
11 euclid(11 a, 11 b, 11 &x, 11 &y) {
   if (!b) return x = 1, y = 0, a;
   11 d = euclid(b, a % b, y, x);
   return y -= a/b * x, d;
}
```

CRT

Description: Chinese Remainder Theorem. crt (a, m, b, n) computes x such that $x \equiv a \pmod m$, $x \equiv b \pmod n$. If |a| < m and |b| < n, x will obey $0 \le x < \operatorname{lcm}(m,n)$. Assumes $mn < 2^{62}$. **Time:** $\log(n)$

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

Euler's thm: a, n coprime $\Rightarrow a^{\phi(n)} \equiv 1 \pmod{n}$.

Fermat's little thm: $p \text{ prime } \Rightarrow a^{p-1} \equiv 1 \pmod{p} \ \forall a.$

cf7d6d, 8 lines

```
const int LIM = 50000000;
int phi[LIM];

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

5.4 Fractions

ContinuedFractions.h

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

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

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

dd6c5e, 21 lines

```
typedef double d; // for N \sim 1e7; long double for N \sim 1e9 pair<11, 11> approximate(d x, 11 N) {
    11 LP = 0, LQ = 1, P = 1, Q = 0, inf = LLONG_MAX; d y = x; for (;;) {
    11 lim = min(P? (N-LP) / P: inf, Q? (N-LQ) / Q: inf), a = (11) floor(y), b = min(a, lim), NP = b*P + LP, NQ = b*Q + LQ; if (a > b) {
        // If b > a/2, we have a semi-convergent that gives us a // better approximation; if b = a/2, we *may* have one. // Return \{P, Q\} here for a more canonical approximation. return (abs(x - (d)NP / (d)NQ) < abs(x - (d)P / (d)Q))?
```

FracBinarySearch FastPrimeCounting Mobius

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

FracBinarySearch.h

Description: Given f and N, finds the smallest fraction $p/q \in [0,1]$ such that f(p/q) is true, and $p,q \leq N$. You may want to throw an exception from f if it finds an exact solution, in which case N can be removed.

Usage: fracBS([](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) {
   11 adv = 0, step = 1; // move hi if dir, else lo
    for (int si = 0; step; (step *= 2) >>= si) {
     Frac mid{lo.p * adv + hi.p, lo.q * adv + hi.q};
     if (abs(mid.p) > N || mid.q > N || dir == !f(mid)) {
       adv -= step; si = 2;
   hi.p += lo.p * adv;
   hi.q += lo.q * adv;
   dir = !dir;
    swap(lo, hi);
   A = B; B = !!adv;
  return dir ? hi : lo;
```

5.5 Pythagorean Triples

The Pythagorean triples are uniquely generated by

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

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

5.6 Primes

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

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

FastPrimeCounting.h

Description: count number of primes less than or equal to n fc1e92, 70 lines

```
struct _count_primes_struct_t_ {
    vector<int> primes, mnprimes;
    ll ans, y;
```

```
vector<pair<pli, int>, char>> queries;
11 count_primes(ll n) {
    y = pow(n, 0.64);
    if (n < 100) y = n;
    primes.clear();
    mnprimes.assign(y + 1, -1);
    ans = 0;
    for (int i = 2; i <= y; ++i) {</pre>
        if (mnprimes[i] == -1) {
            mnprimes[i] = primes.size();
            primes.push_back(i);
        } for (int k = 0; k < primes.size(); ++k) {</pre>
            int j = primes[k];
            if (i * j > y) break;
            mnprimes[i * j] = k;
            if (i % j == 0) break;
    if (n < 100) return primes.size();</pre>
    11 s = n / y;
    for (int p : primes) {
        if (p > s) break;
        ans++;
    int ssz = ans, ptr = primes.size() - 1;
    for (int i = ssz; i < primes.size(); ++i) {</pre>
        while (ptr >= i && (ll)primes[i] * primes[ptr] > n)
        if (ptr < i) break;</pre>
        ans -= ptr - i + 1;
    phi(n, ssz - 1);
    sort(queries.begin(), queries.end());
    int ind = 2, sz = primes.size();
    fenwick fw(sz);
    for (auto [na, sign] : queries) {
        auto [n, a] = na;
        while (ind <= n) fw.add(sz - 1 - mnprimes[ind++],</pre>
        ans += (fw.ask(sz - a - 2) + 1) * sign;
    queries.clear();
    return ans - 1;
void phi(ll n, int a, int sign = 1) {
    if (n == 0) return;
    if (a == -1) {
        ans += n * sign;
        return;
    } if (n <= v) {
        queries.emplace_back(make_pair(n, a), sign);
    phi(n, a - 1, sign);
    phi(n / primes[a], a - 1, -sign);
struct fenwick {
    vector<int> tree;
    fenwick(int n = 0) : n(n) { tree.assign(n, 0); }
    void add(int i, int k) { for (; i < n; i = (i | (i + 1)
        )) tree[i] += k; }
    int ask(int r) {
        int res = 0;
        for (; r \ge 0; r = (r & (r + 1)) - 1) res += tree[r
            ];
        return res;
};
```

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

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

Mobius Inversion:

$$g(n) = \sum_{d|n} f(d) \Leftrightarrow f(n) = \sum_{d|n} \mu(d)g(n/d)$$

Other useful formulas/forms:

$$\sum_{d|n} \mu(d) = [n = 1]$$
 (very useful)

$$g(n) = \sum_{n|d} f(d) \Leftrightarrow f(n) = \sum_{n|d} \mu(d/n)g(d)$$

$$g(n) = \sum_{1 \le m \le n} f(\left| \frac{n}{m} \right|) \Leftrightarrow f(n) = \sum_{1 \le m \le n} \mu(m) g(\left| \frac{n}{m} \right|)$$

Mobius.h

Description: Mobius function

726525, 8 lines

```
vector<int> mobius(VALMAX)
mobius[1] = -1;
for (int i = 1; i < VALMAX; i++) {
   if (mobius[i]) {
      mobius[i] = -mobius[i];
      for (int j = 2 * i; j < VALMAX; j += i) { mobius[j] +=
            mobius[i]; }
   }
}</pre>
```

5.9 Stirling Approximation

$$n! = \sqrt{2\pi n} \left(\frac{n}{e}\right)^n \left(1 + \frac{1}{12n} + \frac{1}{288n^2} - \frac{139}{51840n^3} + \mathcal{O}\left(\frac{1}{n^4}\right)\right)$$

Combinatorial (6)

6.1 Permutations

6.1.1 Factorial

| n | | _ | | • | 8 | - | 10 | |
|-----------------|-------|--------|--------|--------|---------|-----------|--------------|----|
| $\overline{n!}$ | 1 2 6 | 24 1 | 20 720 | 0 5040 | 40320 | 362880 | 3628800 | - |
| n | 11 | 12 | 13 | 14 | 15 | 5 16 | 17 | |
| n! | 4.0e7 | ′ 4.8e | 8 6.2e | 9 8.7e | 10 1.3e | 12 2.1el | 3 3.6e14 | |
| n | 20 | 25 | 30 | 40 | 50 1 | 00 15 | 171 | |
| $\overline{n!}$ | 2e18 | 2e25 | 3e32 | 8e47 3 | 664 9e | 157 6e20 | $62 > DBL_M$ | AX |

IntPerm multinomial BellmanFord

IntPerm.h

Description: Permutation -> integer conversion. (Not order preserving.) Integer -> permutation can use a lookup table.

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

044568, 6 lines

```
int permToInt(vi& v) {
  int use = 0, i = 0, r = 0;
  for(int x:v) r = r * ++i + \underline{\quad} builtin_popcount(use & -(1<<x)),
                                            // (note: minus, not \sim!)
  return r:
```

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 Derangements

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

$$D(n) = (n-1)(D(n-1) + D(n-2)) = nD(n-1) + (-1)^n = \left\lfloor \frac{n!}{e} \right\rfloor$$

6.1.4 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

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

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.

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

$$p(n) \sim 0.145/n \cdot \exp(2.56\sqrt{n})$$

6.2.2 Lucas' Theorem

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 $\binom{n}{m} \equiv \prod_{i=0}^{k} \binom{n_i}{m_i} \pmod{p}$.

6.2.3 Binomials

multinomial.h

General purpose numbers

6.3.1 Bernoulli numbers

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

Sums of powers:

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

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

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

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

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

6.3.4 Stirling numbers of the second kind

Partitions of n distinct elements into exactly k groups.

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

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

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

6.3.5 Bell numbers

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

$$B(p^m + n) \equiv mB(n) + B(n+1) \pmod{p}$$

6.3.6 Labeled unrooted trees

```
# 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

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

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

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

- 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.

Graph (7)

7.1 Fundamentals

BellmanFord.h

Description: Calculates shortest paths from s in a graph that might have negative edge weights. Unreachable nodes get dist = inf; nodes reachable through negative-weight cycles get dist = -inf. Assumes $V^2 \max |w_i| < 2^{63}$. Time: $\mathcal{O}(VE)$

```
const 11 inf = LLONG MAX;
struct Ed { int a, b, w, s() { return a < b ? a : -a; }};
struct Node { ll dist = inf; int prev = -1; };
void bellmanFord(vector<Node>& nodes, vector<Ed>& eds, int s) {
  nodes[s].dist = 0;
  sort(all(eds), [](Ed a, Ed b) { return a.s() < b.s(); });</pre>
  int lim = sz(nodes) / 2 + 2; // /3+100 with shuffled vertices
  rep(i,0,lim) for (Ed ed : eds) {
```

FloydWarshall TopoSort MinCostMaxFlow Dinic

```
Node cur = nodes[ed.a], &dest = nodes[ed.b];
if (abs(cur.dist) == inf) continue;
ll d = cur.dist + ed.w;
if (d < dest.dist) {
  dest.prev = ed.a;
  dest.dist = (i < lim-1 ? d : -inf);
}
rep(i,0,lim) for (Ed e : eds) {
  if (nodes[e.a].dist == -inf)
    nodes[e.b].dist = -inf;
}</pre>
```

FlovdWarshall.h

Description: Calculates all-pairs shortest path in a directed graph that might have negative edge weights. Input is an distance matrix m, where $m[i][j] = \inf$ if i and j are not adjacent. As output, m[i][j] is set to the shortest distance between i and j, inf if no path, or -inf if the path goes through a negative-weight cycle.

Time: $\mathcal{O}\left(N^3\right)$

531245, 12 lines

```
const 11 inf = 1LL << 62;
void floydWarshall(vector<vector<1l>>& m) {
   int n = sz(m);
   rep(i,0,n) m[i][i] = min(m[i][i], 0LL);
   rep(k,0,n) rep(i,0,n) rep(j,0,n)
   if (m[i][k] != inf && m[k][j] != inf) {
      auto newDist = max(m[i][k] + m[k][j], -inf);
      m[i][j] = min(m[i][j], newDist);
   }
   rep(k,0,n) if (m[k][k] < 0) rep(i,0,n) rep(j,0,n)
   if (m[i][k] != inf && m[k][j] != inf) m[i][j] = -inf;
}</pre>
```

TopoSort.h

Description: Topological sorting. Given is an oriented graph. Output is an ordering of vertices, such that there are edges only from left to right. If there are cycles, the returned list will have size smaller than n – nodes reachable from cycles will not be returned.

Time: $\mathcal{O}\left(|V| + |E|\right)$

V|+|E|) d678d8, 8 lines

```
vi topoSort(const vector<vi>& gr) {
  vi indeg(sz(gr)), q;
  for (auto& li : gr) for (int x : li) indeg[x]++;
  rep(i,0,sz(gr)) if (indeg[i] == 0) q.push_back(i);
  rep(j,0,sz(q)) for (int x : gr[q[j]])
   if (--indeg[x] == 0) q.push_back(x);
  return q;
}
```

7.2 Network flow

MinCostMaxFlow.h

Description: Min-cost max-flow. 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: $\mathcal{O}(FE\log(V))$ where F is max flow. $\mathcal{O}(VE)$ for setpi. _{58385b}, 79 lines

```
#include <bits/extc++.h>
const ll INF = numeric_limits<ll>::max() / 4;

struct MCMF {
    struct edge {
        int from, to, rev;
        ll cap, cost, flow;
    };
    int N;
    vector<vector<edge>> ed;
```

```
vi seen;
  vector<ll> dist, pi;
  vector<edge*> par;
  MCMF(int N) : N(N), ed(N), seen(N), dist(N), pi(N), par(N) {}
  void addEdge(int from, int to, ll cap, ll cost) {
    if (from == to) return;
    ed[from].push_back(edge{ from, to, sz(ed[to]), cap, cost, 0 });
    ed[to].push_back(edge{ to,from,sz(ed[from])-1,0,-cost,0 });
  void path(int s) {
   fill(all(seen), 0);
    fill(all(dist), INF);
    dist[s] = 0; ll di;
    __qnu_pbds::priority_queue<pair<ll, int>> q;
    vector<decltype(g)::point_iterator> its(N);
    q.push({ 0, s });
    while (!q.empty()) {
     s = q.top().second; q.pop();
      seen[s] = 1; di = dist[s] + pi[s];
      for (edge& e : ed[s]) if (!seen[e.to]) {
       11 val = di - pi[e.to] + e.cost;
       if (e.cap - e.flow > 0 && val < dist[e.to]) {</pre>
          dist[e.to] = val;
          par[e.to] = &e;
          if (its[e.to] == q.end())
            its[e.to] = q.push({ -dist[e.to], e.to });
            q.modify(its[e.to], { -dist[e.to], e.to });
    rep(i, 0, N) pi[i] = min(pi[i] + dist[i], INF);
  pair<11, 11> maxflow(int s, int t) {
    11 \text{ totflow} = 0, totcost = 0;
    while (path(s), seen[t]) {
      11 fl = INF;
      for (edge* x = par[t]; x; x = par[x->from])
        fl = min(fl, x->cap - x->flow);
      totflow += fl;
      for (edge* x = par[t]; x; x = par[x->from]) {
       x \rightarrow flow += fl;
        ed[x->to][x->rev].flow -= fl;
    rep(i,0,N) for(edge& e : ed[i]) totcost += e.cost * e.flow;
    return {totflow, totcost/2};
  // If some costs can be negative, call this before maxflow:
  void setpi(int s) { // (otherwise, leave this out)
    fill(all(pi), INF); pi[s] = 0;
    int it = N, ch = 1; ll v;
    while (ch-- && it--)
      rep(i,0,N) if (pi[i] != INF)
        for (edge& e : ed[i]) if (e.cap)
          if ((v = pi[i] + e.cost) < pi[e.to])
            pi[e.to] = v, ch = 1;
    assert(it >= 0); // negative cost cycle
};
```

```
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 match-
```

Dinic.h

```
struct Dinic {
  struct Edge {
    int to, rev;
    11 c, oc;
    11 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) {}
  void addEdge(int a, int b, ll c, ll rcap = 0) {
    adj[a].push_back({b, sz(adj[b]), c, c});
    adj[b].push_back({a, sz(adj[a]) - 1, rcap, rcap});
 11 dfs(int v, int t, 11 f) {
    if (v == t || !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;
  11 calc(int s, int t) {
    11 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++];
        for (Edge e : adi[v])
          if (!lvl[e.to] && e.c >> (30 - L))
            q[qe++] = e.to, lvl[e.to] = lvl[v] + 1;
      while (ll p = dfs(s, t, LLONG_MAX)) flow += p;
    } while (lvl[t]);
    return flow;
 bool leftOfMinCut(int a) { return lvl[a] != 0; }
```

7.2.1 Flows with Demands(Lower Bound on flows)

Finding an Arbitrary flow: We make the following changes in the network. We add a new source s' and a new sink t', a new edge from the source s' to every other vertex, a new edge for every vertex to the sink t', and one edge from t to s. Additionally we define the new capacity function c' as:

$$\begin{split} c'((s',v)) &= \sum_{u \in V} d((u,v)) \forall (s',v) \\ c'((v,t')) &= \sum_{w \in V} d((v,w)) \forall (v,t') \\ c'((u,v)) &= c((u,v)) - d((u,v)) \forall (u,v) \end{split}$$

in the old network

$$c'((t,s)) = INF$$

Minimal Flow: The edge (t,s) flows the entire flow of the corresponding old network, and if we keep the c'((t,s)) to be INF, it will have the same flow of old network, but if we reduce the capacity to a very small value, the demands will not meet. So we can do a binary search on capacity of this edge and try to reduce it as much as possible so that demands meet.

MinCut.h

Description: After running max-flow, the left side of a min-cut from s to t is given by all vertices reachable from s, only traversing edges with positive residual capacity.

GlobalMinCut.h

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

Time: $\mathcal{O}\left(V^3\right)$

8b0e19, 21 lines

```
pair<int, vi> globalMinCut(vector<vi> mat) {
  pair<int, vi> best = {INT_MAX, {}};
  int n = sz(mat);
  vector<vi> co(n);
 rep(i, 0, n) co[i] = {i};
  rep(ph,1,n) {
   vi w = mat[0];
    size_t s = 0, t = 0;
   rep(it,0,n-ph) { //O(V^2) \rightarrow O(E log V) with prio. queue
     w[t] = INT_MIN;
     s = t, t = max_element(all(w)) - w.begin();
     rep(i, 0, n) w[i] += mat[t][i];
   best = min(best, \{w[t] - mat[t][t], co[t]\});
    co[s].insert(co[s].end(), all(co[t]));
    rep(i,0,n) mat[s][i] += mat[t][i];
   rep(i, 0, n) mat[i][s] = mat[s][i];
   mat[0][t] = INT_MIN;
  return best;
```

GomoryHu.h

Description: Given a list of edges representing an undirected flow graph, returns edges of the Gomory-Hu tree. The max flow between any pair of vertices is given by minimum edge weight along the Gomory-Hu tree path. **Time:** $\mathcal{O}\left(V\right)$ Flow Computations

```
"PushRelabel.h" 0418b3, 13 lines
```

```
typedef array<11, 3> Edge;
vector<Edge> gomoryHu(int N, vector<Edge> ed) {
  vector<Edge> tree;
  vi par(N);
  rep(i,1,N) {
    PushRelabel D(N); // Dinic also works
    for (Edge t : ed) D.addEdge(t[0], t[1], t[2], t[2]);
    tree.push_back({i, par[i], D.calc(i, par[i])});
  rep(j,i+1,N)
    if (par[j] == par[i] && D.leftOfMinCut(j)) par[j] = i;
  }
  return tree;
}
```

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;
  for (int b : q[a]) if (B[b] == L + 1) {
   B[b] = 0;
    if (btoa[b] == -1 || dfs(btoa[b], L + 1, q, 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();
    for (int a : btoa) if (a != -1) A[a] = -1;
    rep(a, 0, sz(q)) if(A[a] == 0) cur.push_back(a);
    for (int lay = 1;; lay++) {
      bool islast = 0;
      next.clear();
      for (int a : cur) for (int b : q[a]) {
        if (btoa[b] == -1) {
          B[b] = lay;
          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;
      for (int a : next) A[a] = lay;
      cur.swap(next);
    rep(a, 0, sz(q))
      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(q, btoa);

```
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"
vi cover(vector<vi>& g, int n, int m) {
 vi match(m, -1);
 int res = dfsMatching(q, match);
  vector<bool> lfound(n, true), seen(m);
  for (int it : match) if (it != -1) lfound[it] = false;
  rep(i,0,n) if (lfound[i]) q.push_back(i);
  while (!q.emptv()) {
   int i = q.back(); q.pop_back();
    lfound[i] = 1;
    for (int 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: Given a weighted bipartite graph, matches every node on the left with a node on the right such that no nodes are in two matchings and the sum of the edge weights is minimal. Takes cost[N][M], where cost[i][j] = cost for L[i] to be matched with R[j] and returns (min cost, match), where L[i] is matched with R[match[i]]. Negate costs for max cost. Requires $N \leq M$. **Time:** $\mathcal{O}(N^2M)$

```
pair<int, vi> hungarian(const vector<vi> &a) {
   if (a.empty()) return {0, {}};
   int n = sz(a) + 1, m = sz(a[0]) + 1;
   vi u(n), v(m), p(m), ans(n - 1);
   rep(i,l,n) {
     p[0] = i;
   int j0 = 0; // add "dumnny" worker 0
```

```
vi dist(m, INT_MAX), pre(m, -1);
vector<bool> done(m + 1);
do { // dijkstra
  done[j0] = true;
  int i0 = p[j0], j1, delta = INT_MAX;
  rep(j,1,m) if (!done[j]) {
    auto cur = a[i0 - 1][j - 1] - u[i0] - v[j];
    if (cur < dist[j]) dist[j] = cur, pre[j] = j0;
    if (dist[j] < delta) delta = dist[j], j1 = j;</pre>
   if (done[j]) u[p[j]] += delta, v[j] -= delta;
    else dist[j] -= delta;
  j0 = j1;
} while (p[j0]);
while (j0) { // update alternating path
  int j1 = pre[j0];
  p[j0] = p[j1], j0 = j1;
```

```
rep(j,1,m) if (p[j]) ans[p[j] - 1] = j - 1;
return {-v[0], ans}; // min cost
```

GeneralMatchingBlossom.h

Description: Variant on Gabow's Impl of Edmond's Blossom Algorithm. General unweighted max matching with 1-based indexing. If white[v] = 0 after solve() returns, v is part of every max matching.

Time: $\mathcal{O}(NM)$, faster in practice

fd5cc7, 50 lines

```
struct MaxMatching {
  int N; V<vi> adj;
  V<int> mate, first; vb white; vpi label;
  void init(int _N) { N = _N; adj = V<vi>(N+1);
   mate = first = vi(N+1); label = vpi(N+1); white = vb(N+1);
  void ae(int u, int v) { adj.at(u).pb(v), adj.at(v).pb(u); }
  int group(int x) { if (white[first[x]]) first[x] = group(
       first[x]);
    return first[x]; }
  void match (int p, int b) {
    swap(b, mate[p]); if (mate[b] != p) return;
    if (!label[p].s) mate[b] = label[p].f, match(label[p].f,b);
          // vertex label
    else match(label[p].f,label[p].s), match(label[p].s,label[p
        ].f); // edge label
  bool augment(int st) { assert(st);
    white[st] = 1; first[st] = 0; label[st] = \{0,0\};
    queue<int> q; q.push(st);
    while (!q.empty()) {
     int a = q.ft; q.pop(); // outer vertex
      each(b,adj[a]) { assert(b);
        if (white[b]) { // two outer vertices, form blossom
          int x = group(a), y = group(b), lca = 0;
          while (x | | v) {
            if (y) swap(x,y);
            if (label[x] == pi{a,b}) { lca = x; break; }
            label[x] = \{a,b\}; x = group(label[mate[x]].first);
          for (int v: {group(a),group(b)}) while (v != lca) {
            assert(!white[v]); // make everything along path
            q.push(v); white[v] = true; first[v] = lca;
            v = group(label[mate[v]].first);
        } else if (!mate[b]) { // found augmenting path
          mate[b] = a; match(a,b); white = vb(N+1); // reset
          return true;
        } else if (!white[mate[b]]) {
          white[mate[b]] = true; first[mate[b]] = b;
          label[b] = \{0,0\}; label[mate[b]] = pi\{a,0\};
          q.push(mate[b]);
   return false:
  int solve() {
    int ans = 0:
   FOR(st,1,N+1) if (!mate[st]) ans += augment(st);
   FOR(st,1,N+1) if (!mate[st] && !white[st]) assert(!augment(
        st));
    return ans;
};
```

7.4 DFS algorithms

SCC.h

Description: Finds strongly connected components in a directed graph. If vertices u, v belong to the same component, we can reach u from v and vice

Usage: $scc(graph, [\&](vi\& v) \{ ... \})$ visits all components in reverse topological order. comp[i] holds the component index of a node (a component only has edges to components with lower index). ncomps will contain the number of components. Time: $\mathcal{O}(E+V)$

```
vi val, comp, z, cont;
int Time, ncomps;
template < class G, class F> int dfs (int j, G& g, F& f) {
 int low = val[j] = ++Time, x; z.push_back(j);
  for (auto e : q[j]) if (comp[e] < 0)</pre>
    low = min(low, val[e] ?: dfs(e,g,f));
  if (low == val[i]) {
    do {
     x = z.back(); z.pop_back();
     comp[x] = ncomps;
     cont.push back(x);
    } while (x != i);
    f(cont); cont.clear();
    ncomps++;
  return val[j] = low;
template < class G, class F > void scc(G& q, F f) {
 int n = sz(q);
 val.assign(n, 0); comp.assign(n, -1);
 Time = ncomps = 0;
 rep(i, 0, n) if (comp[i] < 0) dfs(i, g, f);
```

BiconnectedComponents.h

if (num[v] < me)</pre>

int si = sz(st);

if (up == me) {

} else {

st.push_back(e);

int up = dfs(y, e, f);

top = min(top, up);

st.push_back(e);

st.resize(si);

f(vi(st.begin() + si, st.end()));

Description: Finds all biconnected components in an undirected graph, and runs a callback for the edges in each. In a biconnected component there are at least two distinct paths between any two nodes. Note that a node can be in several components. An edge which is not in a component is a bridge, i.e., not part of any cycle.

```
Usage: int eid = 0; ed.resize(N);
for each edge (a,b) {
ed[a].emplace_back(b, eid);
ed[b].emplace_back(a, eid++); }
bicomps([&](const vi& edgelist) {...});
Time: \mathcal{O}(E+V)
                                                      c6b7c7, 32 lines
vi num, st;
vector<vector<pii>> ed;
int Time:
template<class F>
int dfs(int at, int par, F& f) {
 int me = num[at] = ++Time, top = me;
 for (auto [y, e] : ed[at]) if (e != par) {
    if (num[y]) {
      top = min(top, num[y]);
```

```
else if (up < me) st.push_back(e);</pre>
      else { /* e is a bridge */ }
 return top;
template<class F>
void bicomps(F f) {
 num.assign(sz(ed), 0);
 rep(i,0,sz(ed)) if (!num[i]) dfs(i, -1, f);
```

2sat.h

76b5c9, 24 lines

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

```
Usage: TwoSat ts (number of boolean variables);
ts.either(0, \sim3); // Var 0 is true or var 3 is false
ts.setValue(2); // Var 2 is true
ts.atMostOne(\{0, \sim 1, 2\}); // <= 1 of vars 0, \sim 1 and 2 are true
ts.solve(); // Returns true iff it is solvable
ts.values[0..N-1] holds the assigned values to the vars
```

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

```
5f9706, 56 lines
```

```
struct TwoSat {
  int N;
  vector<vi> gr:
 vi values; // 0 = false, 1 = true
  TwoSat(int n = 0) : N(n), qr(2*n) {}
  int addVar() { // (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 setValue(int x) { either(x, x); }
  void atMostOne(const vi& li) { // (optional)
    if (sz(li) <= 1) return;</pre>
    int cur = \simli[0];
    rep(i,2,sz(li)) {
      int next = addVar();
      either(cur, ~li[i]);
      either(cur, next);
      either(~li[i], next);
      cur = ~next;
    either(cur, ~li[1]);
 vi val, comp, z; int time = 0;
 int dfs(int i) {
    int low = val[i] = ++time, x; z.push_back(i);
    for(int e : qr[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;
};
```

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);
  for (int x : D) if (x < 0 \mid \mid sz(ret) != nedges+1) return {};
  return {ret.rbegin(), ret.rend()};
```

7.5 Coloring

EdgeColoring.h

Description: Given a simple, undirected graph with max degree D, computes a (D+1)-coloring of the edges such that no neighboring edges share a color. (D-coloring is NP-hard, but can be done for bipartite graphs by repeated matchings of max-degree nodes.)

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

e210e2, 31 lines

```
vi edgeColoring(int N, vector<pii> eds) {
 vi cc(N + 1), ret(sz(eds)), fan(N), free(N), loc;
 for (pii e : eds) ++cc[e.first], ++cc[e.second];
 int u, v, ncols = *max_element(all(cc)) + 1;
 vector<vi> adj(N, vi(ncols, -1));
  for (pii e : eds) {
   tie(u, v) = e;
   fan[0] = v;
   loc.assign(ncols, 0);
   int at = u, end = u, d, c = free[u], ind = 0, i = 0;
   while (d = free[v], !loc[d] && (v = adj[u][d]) != -1)
     loc[d] = ++ind, cc[ind] = d, fan[ind] = v;
   cc[loc[d]] = c;
   for (int cd = d; at != -1; cd ^= c ^ d, at = adj[at][cd])
     swap(adj[at][cd], adj[end = at][cd ^ c ^ d]);
    while (adj[fan[i]][d] != -1) {
     int left = fan[i], right = fan[++i], e = cc[i];
     adj[u][e] = left;
     adj[left][e] = u;
     adj[right][e] = -1;
     free[right] = e;
```

```
adj[u][d] = fan[i];
  adj[fan[i]][d] = u;
  for (int y : {fan[0], u, end})
    for (int& z = free[y] = 0; adj[y][z] != -1; z++);
rep(i, 0, sz(eds))
  for (tie(u, v) = eds[i]; adj[u][ret[i]] != v;) ++ret[i];
return ret;
```

7.6 Heuristics

MaximalCliques.h

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

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

b0d5b1, 12 lines

```
typedef bitset<128> B;
template<class F>
void cliques(vector<B>& eds, F f, B P = \simB(), 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. f7c0bc, 49 lines

```
typedef vector<br/>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) {
   for (auto& v : r) v.d = 0;
   for (auto& v : r) for (auto 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;</pre>
      q.push_back(R.back().i);
      vv T;
     for(auto v:R) if (e[R.back().i][v.i]) T.push_back({v.i});
      if (sz(T)) {
       if (S[lev]++ / ++pk < limit) init(T);</pre>
       int j = 0, mxk = 1, mnk = max(sz(qmax) - sz(q) + 1, 1);
       C[1].clear(), C[2].clear();
        for (auto 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++].i = v.i;
          C[k].push_back(v.i);
        if (j > 0) T[j - 1].d = 0;
        rep(k, mnk, mxk + 1) for (int i : C[k])
          T[j].i = i, T[j++].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});
};
```

MaximumIndependentSet.h

Description: To obtain a maximum independent set of a graph, find a max clique of the complement. If the graph is bipartite, see MinimumVertex-

7.7 Trees

BinaryLifting.h

Description: Calculate power of two jumps in a tree, to support fast upward jumps and LCAs. Assumes the root node points to itself.

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

bfce85, 25 lines

```
vector<vi> treeJump(vi& P) {
 int on = 1, d = 1;
  while (on < sz(P)) on *= 2, d++;
 vector<vi> jmp(d, P);
  rep(i,1,d) rep(j,0,sz(P))
   jmp[i][j] = jmp[i-1][jmp[i-1][j]];
  return jmp;
int jmp(vector<vi>& tbl, int nod, int steps){
 rep(i,0,sz(tbl))
    if(steps&(1<<i)) nod = tbl[i][nod];
  return nod;
int lca(vector<vi>& tbl, vi& depth, int a, int b) {
 if (depth[a] < depth[b]) swap(a, b);</pre>
  a = jmp(tbl, a, depth[a] - depth[b]);
 if (a == b) return a;
 for (int i = sz(tbl); i--;) {
    int c = tbl[i][a], d = tbl[i][b];
    if (c != d) a = c, b = d;
 return tbl[0][a];
```

LCA.h

Description: Lowest common ancestor in 0-rooted tree

"../data-structures/SparseTable.h" f6fdc3, 21 lines struct LCA { int T = 0;vi time, path, ret; SparseTable st; $LCA(vector < vi > \& C) : time(sz(C)), st((dfs(C,0,-1), ret)) {}$ void dfs(vector<vi>& C, int v, int par) { time[v] = T++;

CompressTree HLD DirectedMST Point

```
for (int y : C[v]) if (y != par) {
   path.push_back(v), ret.push_back(time[v]);
   dfs(C, y, v);
int lca(int a, int b) {
 if (a == b) return a;
 tie(a, b) = minmax(time[a], time[b]);
 return path[st.query_val(a, b - 1)];
//dist(a,b){return depth[a] + depth[b] - 2*depth[lca(a,b)];}
```

CompressTree.h

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

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

"LCA.h" 9775a0, 21 lines typedef vector<pair<int, int>> vpi; vpi compressTree(LCA& lca, const vi& subset) { static vi rev; rev.resize(sz(lca.time)); vi li = subset, &T = lca.time; auto cmp = [&](int a, int b) { return T[a] < T[b]; };</pre> sort(all(li), cmp); int m = sz(li)-1;rep(i,0,m) { int a = li[i], b = li[i+1]; li.push_back(lca.lca(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.lca(a, b)], b); return ret;

Description: Decomposes a tree into vertex disjoint heavy paths and light edges such that the path from any leaf to the root contains at most log(n) light edges. Code does additive modifications and max queries, but can support commutative segtree modifications/queries on paths and subtrees. Takes as input the full adjacency list. VALS_EDGES being true means that values are stored in the edges, as opposed to the nodes. All values initialized to the segtree default. Root must be 0.

```
Time: \mathcal{O}\left((\log N)^2\right)
"../data-structures/LazySegmentTree.h"
                                                         9547af, 46 lines
template <bool VALS EDGES> struct HLD {
 int N, tim = 0;
 vector<vi> adj;
 vi par, siz, rt, pos;
 Node *tree;
  HLD(vector<vi> adj_)
    : N(sz(adj_)), adj(adj_), par(N, -1), siz(N, 1),
      rt(N),pos(N),tree(new Node(0, N)) { dfsSz(0); dfsHld(0); }
  void dfsSz(int v) {
    for (int& u : adj[v]) {
      adj[u].erase(find(all(adj[u]), v));
      par[u] = v;
      dfsSz(u);
      siz[v] += siz[u];
      if (siz[u] > siz[adj[v][0]]) swap(u, adj[v][0]);
```

```
void dfsHld(int v) {
   pos[v] = tim++;
   for (int u : adj[v]) {
     rt[u] = (u == adj[v][0] ? rt[v] : u);
     dfsHld(u);
 template <class B> void process(int u, int v, B op) {
   for (;; v = par[rt[v]]) {
     if (pos[u] > pos[v]) swap(u, v);
     if (rt[u] == rt[v]) break;
     op(pos[rt[v]], pos[v] + 1);
   op(pos[u] + VALS_EDGES, pos[v] + 1);
 void modifyPath(int u, int v, int val) {
   process(u, v, [&](int l, int r) { tree->add(l, r, val); });
 int queryPath(int u, int v) { // Modify depending on problem
   int res = -1e9;
   process(u, v, [&](int l, int r) {
       res = max(res, tree->query(1, r));
   });
   return res;
 int querySubtree(int v) { // modifySubtree is similar
   return tree->query(pos[v] + VALS_EDGES, pos[v] + siz[v]);
};
```

DirectedMST.h

Description: Finds a minimum spanning tree/arborescence of a directed graph, given a root node. If no MST exists, returns -1.

Time: $\mathcal{O}(E \log V)$

"../data-structures/UnionFindRollback.h"

```
struct Edge { int a, b; ll w; };
struct Node {
 Edge key;
 Node *1, *r;
  11 delta;
  void prop() {
    kev.w += delta;
    if (1) 1->delta += delta;
    if (r) r->delta += delta;
    delta = 0:
  Edge top() { prop(); return key; }
Node *merge(Node *a, Node *b) {
  if (!a || !b) return a ?: b;
  a->prop(), b->prop();
  if (a->key.w > b->key.w) swap(a, b);
  swap(a->1, (a->r = merge(b, a->r)));
  return a:
void pop(Node*& a) { a->prop(); a = merge(a->1, a->r); }
pair<ll, vi> dmst(int n, int r, vector<Edge>& g) {
  RollbackUF uf(n);
  vector<Node*> heap(n);
  for (Edge e : q) heap[e.b] = merge(heap[e.b], new Node{e});
  11 \text{ res} = 0;
  vi seen(n, -1), path(n), par(n);
  seen[r] = r;
  vector<Edge> Q(n), in(n, \{-1,-1\}), comp;
  deque<tuple<int, int, vector<Edge>>> cycs;
  rep(s, 0, n) {
```

```
int u = s, qi = 0, w;
  while (seen[u] < 0) {
    if (!heap[u]) return {-1,{}};
    Edge e = heap[u] \rightarrow top();
    heap[u]->delta -= e.w, pop(heap[u]);
    Q[qi] = e, path[qi++] = u, seen[u] = s;
    res += e.w, u = uf.find(e.a);
    if (seen[u] == s) {
      Node * cvc = 0;
      int end = qi, time = uf.time();
      do cyc = merge(cyc, heap[w = path[--qi]]);
      while (uf.join(u, w));
      u = uf.find(u), heap[u] = cyc, seen[u] = -1;
      cycs.push_front({u, time, {&Q[qi], &Q[end]}});
  rep(i, 0, qi) in[uf.find(Q[i].b)] = Q[i];
for (auto& [u,t,comp] : cycs) { // restore sol (optional)
  uf.rollback(t);
  Edge inEdge = in[u];
  for (auto& e : comp) in[uf.find(e.b)] = e;
  in[uf.find(inEdge.b)] = inEdge;
rep(i,0,n) par[i] = in[i].a;
return {res, par};
```

7.8 Math

39e620, 60 lines

7.8.1 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]--, mat[a][a]++if G is undirected). Remove the *i*th row and column and take the determinant; this yields the number of directed spanning trees rooted at i (if G is undirected, remove any row/column).

7.8.2 Erdős–Gallai theorem

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

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

Geometry (8)

8.1 Geometric primitives

Point.h

```
Description: Class to handle points in the plane. T can be e.g. double or
long long. (Avoid int.)
```

```
template \langle class T \rangle int sgn(T x) \{ return (x > 0) - (x < 0); \}
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); }</pre>
 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) {</pre>
 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 /S on the result of the cross product.



f6bf6b, 4 lines

```
template<class P>
double lineDist(const P& a, const P& b, const P& p) {
 return (double) (b-a).cross(p-a)/(b-a).dist();
```

SegmentDistance.h

Description:

Returns the shortest distance between point p and the line segment from point s to e.

Usage: Point < double > a, b(2,2), p(1,1); bool onSegment = segDist(a,b,p) < 1e-10;



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<|l> and the intersection point does not have integer coordinates. Products of three coordinates are used in intermediate steps so watch out for overflow if using int or long long.



```
Usage: vector<P> inter = segInter(s1,e1,s2,e2);
if (sz(inter) == 1)
cout << "segments intersect at " << inter[0] << endl;
"Point.h", "OnSegment.h"
                                                       9d57f2, 13 lines
```

```
template < class P > vector < P > segInter (P a, P b, P c, P d) {
 auto oa = c.cross(d, a), ob = c.cross(d, b),
       oc = a.cross(b, c), od = a.cross(b, d);
```

```
// Checks if intersection is single non-endpoint point.
if (sgn(oa) * sgn(ob) < 0 && sgn(oc) * sgn(od) < 0)
 return { (a * ob - b * oa) / (ob - oa) };
set<P> s:
if (onSegment(c, d, a)) s.insert(a);
if (onSegment(c, d, b)) s.insert(b);
if (onSegment(a, b, c)) s.insert(c);
if (onSegment(a, b, d)) s.insert(d);
return {all(s)};
```

lineIntersection.h

Description:

If a unique intersection point of the lines going through s1,e1 and s2,e2 exists {1, point} is returned. If no intersection point exists $\{0, (0,0)\}\$ is returned and if infinitely many exists $\{-1,$ (0,0)} is returned. The wrong position will be returned if P is Point<|l> and the intersection point does not have integer coordinates. Products of three coordinates are used in inter- \(\sigma 1 \) mediate steps so watch out for overflow if using int or ll.

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

sideOf.h

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

```
"Point.h"
                                                       3af81c, 9 lines
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).cross(p-s);
 double 1 = (e-s).dist()*eps;
 return (a > 1) - (a < -1);
```

Usage: bool left = sideOf(p1,p2,q)==1;

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) <= 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"
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();
```

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, v;
  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 || v);
    return v < 0 || (v == 0 && x < 0);
  Angle t90() const { return \{-y, x, t + (half() \&\& x >= 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) <</pre>
         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> segmentAngles(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;</pre>
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)};
```

8.2 Circles

CircleIntersection.h

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

```
"Point.h"
                                                      84d6d3, 11 lines
typedef Point<double> P;
bool circleInter(P a, P b, double r1, double r2, pair < P, P >* out) {
  if (a == b) { assert(r1 != r2); return false; }
  P \text{ vec} = b - a;
  double d2 = 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);
  *out = {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" b0153d, 13 lines

```
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.dist2(), h2 = d2 - dr * dr;
  if (d2 == 0 || h2 < 0) return {};</pre>
  vector<pair<P, P>> out;
  for (double sign : {-1, 1}) {
   P v = (d * dr + d.perp() * sqrt(h2) * sign) / d2;
   out.push_back(\{c1 + v * r1, c2 + v * r2\});
 if (h2 == 0) out.pop_back();
 return out;
```

CirclePolygonIntersection.h

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

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

```
"../../content/geometry/Point.h"
                                                       alee63, 19 lines
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;
   P d = q - p;
    auto a = d.dot(p)/d.dist2(), b = (p.dist2()-r*r)/d.dist2();
   auto det = a * a - b;
   if (det <= 0) return arg(p, q) * r2;</pre>
    auto s = max(0., -a-sqrt(det)), t = min(1., -a+sqrt(det));
   if (t < 0 || 1 <= s) return arg(p, g) * r2;</pre>
   P u = p + d * s, v = p + d * t;
   return arg(p,u) * r2 + u.cross(v)/2 + arg(v,q) * r2;
```

circumcircle.h

return sum;

auto sum = 0.0;

rep(i, 0, sz(ps))

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.

sum += tri(ps[i] - c, ps[(i + 1) % sz(ps)] - c);



1caa3a, 9 lines "Point.h"

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

MinimumEnclosingCircle.h

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

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

8.3 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! f12300, 6 lines

```
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

"Point.h"

Description: Returns the center of mass for a polygon. Time: $\mathcal{O}(n)$

```
9706dc, 9 lines
typedef Point<double> P;
P polygonCenter(const vector<P>& v) {
 P res(0, 0); double A = 0;
 for (int i = 0, j = sz(v) - 1; i < sz(v); j = i++) {
   res = res + (v[i] + v[j]) * v[j].cross(v[i]);
   A += v[j].cross(v[i]);
 return res / A / 3;
```

PolygonCut.h

Description:

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

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



```
"Point.h", "lineIntersection.h"
                                                       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;</pre>
    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:

Returns a vector of the points of the convex hull in counterclockwise order. Points on the edge of the hull between two other points are not considered part of the hull.



Time: $\mathcal{O}(n \log n)$ "Point.h"

```
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)))
    for (P p : pts) {
      while (t \ge s + 2 \&\& h[t-2].cross(h[t-1], p) \le 0) t--;
  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/collinear points). Time: $\mathcal{O}(n)$

```
"Point.h"
                                                              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(\text{res}, \{(S[i] - S[j]).dist2(), \{S[i], S[j]\}\});
      if ((S[(j+1) % n] - S[j]).cross(S[i+1] - S[i]) >= 0)
        break:
  return res.second;
```

PointInsideHull.h

Description: Determine whether a point t lies inside a convex hull (CCW order, with no collinear 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(l) - 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;
  }
  return sgn(l[a].cross(l[b], p)) < r;
}</pre>
```

LineHullIntersection.h

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

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

```
"Point.h"
                                                     7cf45b, 39 lines
#define cmp(i, j) sqn(dir.perp().cross(poly[(i)%n]-poly[(j)%n]))
#define extr(i) cmp(i + 1, i) >= 0 && cmp(i, i - 1 + n) < 0
template <class P> 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 1s = cmp(1o + 1, 1o), ms = cmp(m + 1, m);
   (ls < ms \mid | (ls == ms \&\& ls == cmp(lo, m)) ? hi : lo) = m;
 return lo;
#define cmpL(i) sgn(a.cross(poly[i], b))
template <class P>
array<int, 2> lineHull(P a, P b, vector<P>& poly) {
  int endA = extrVertex(poly, (a - b).perp());
  int endB = extrVertex(poly, (b - a).perp());
  if (cmpL(endA) < 0 \mid \mid 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:
```

8.4 Misc. Point Set Problems

```
ClosestPair.h
Description: Finds the closest pair of points.
Time: \mathcal{O}(n \log n)
"Point.h"
                                                       ac41a6, 17 lines
typedef Point<ll> P;
pair<P, P> closest (vector<P> v) {
 assert (sz(v) > 1);
 set<P> S;
 sort(all(v), [](P a, P b) { return a.y < b.y; });
  pair<ll, pair<P, P>> ret{LLONG MAX, {P(), P()}};
 int j = 0;
 for (P p : v) {
   P d{1 + (ll)sqrt(ret.first), 0};
    while (v[j].y \le p.y - d.x) S.erase(v[j++]);
    auto lo = S.lower_bound(p - d), hi = S.upper_bound(p + d);
    for (; lo != hi; ++lo)
      ret = min(ret, {(*lo - p).dist2(), {*lo, p}});
    S.insert(p);
  return ret.second;
```

8.5 3D

PolyhedronVolume.h

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

3058c3, 6 lines

```
template<class V, class L>
double signedPolyVolume(const V& p, const L& trilist) {
  double v = 0;
  for (auto 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.

```
template < class T > struct Point 3D {
 typedef Point3D P:
 typedef const P& R;
 T x, v, 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;
};
```

3dHull.h

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"
                                                     5b45fc, 49 lines
typedef Point3D<double> P3;
struct PR {
  void ins(int x) { (a == -1 ? a : b) = x; }
  void rem(int x) { (a == x ? a : b) = -1; }
  int cnt() { return (a !=-1) + (b !=-1); }
  int a, b;
struct F { P3 q; int a, b, c; };
vector<F> hull3d(const vector<P3>& A) {
  assert(sz(A) >= 4);
  vector<vector<PR>>> E(sz(A), vector<PR>(sz(A), {-1, -1}));
#define E(x, y) E[f.x][f.y]
  vector<F> FS:
  auto mf = [\&] (int i, int j, int k, int l) {
    P3 q = (A[j] - A[i]).cross((A[k] - A[i]));
    if (q.dot(A[1]) > q.dot(A[i]))
      q = q * -1;
    F f{q, i, j, k};
    E(a,b).ins(k); E(a,c).ins(j); E(b,c).ins(i);
    FS.push_back(f);
  rep(i,0,4) rep(j,i+1,4) rep(k,j+1,4)
    mf(i, j, k, 6 - i - j - k);
  rep(i,4,sz(A)) {
    rep(j,0,sz(FS)) {
      F f = FS[j];
      if(f.q.dot(A[i]) > f.q.dot(A[f.a])) {
        E(a,b).rem(f.c);
        E(a,c).rem(f.b);
        E(b,c).rem(f.a);
        swap(FS[j--], FS.back());
        FS.pop_back();
    int nw = sz(FS);
    rep(j,0,nw) {
     F f = FS[j];
#define C(a, b, c) if (E(a,b).cnt() != 2) mf(f.a, f.b, i, f.c);
      C(a, b, c); C(a, c, b); C(b, c, a);
  for (F& it : FS) if ((A[it.b] - A[it.a]).cross(
    A[it.c] - A[it.a]).dot(it.q) <= 0) swap(it.c, it.b);
  return FS:
```

sphericalDistance.h

Description: Returns the shortest distance on the sphere with radius radius between the points with azimuthal angles (longitude) f1 (ϕ_1) and f2 (ϕ_2) from x axis and zenith angles (latitude) t1 (θ_1) and t2 (θ_2) from z axis (0 = north pole). 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.

```
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: $\mathcal{O}\left(n\right)$ 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[i] computes the length of the longest common prefix of s[i:] and s, except z[0] = 0. (abacaba -> 0010301) **Time:** $\mathcal{O}(n)$

```
vi Z(const string& S) {
```

```
vi Z(const 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).

```
array < vi, 2 > p = {vi(n+1), vi(n)};
 rep(z,0,2) for (int i=0,1=0,r=0; i < n; i++) {
    int t = r-i+!z;
    if (i<r) p[z][i] = min(t, p[z][l+t]);</pre>
    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()+minRotation(v), v.end());
Time: \mathcal{O}(N)
int minRotation(string s) {
 int a=0, N=sz(s); s += s;
 rep(b, 0, N) rep(k, 0, N) {
    if (a+k == b \mid | s[a+k] < s[b+k]) {b += max(0, k-1); break;}
    if (s[a+k] > s[b+k]) { a = b; break; }
 return a;
SuffixArray.h
Description: Suffix Array
"../data-structures/SparseTable.h"
                                                        6e50f5, 74 lines
struct SuffixArrav
    vector<int> p,c,_lcp;
    SparseTable* st:
    int n;
    void count_sort(vector<int>& p, vector<int>&c)
        int n = p.size();
        vector<int> cnt(n);
        for(auto x:c)cnt[x]++;
        vector<int> pos(n);
        vector<int> p_new(n);
        pos[0] = 0;
        for(int i=1;i<n;i++)pos[i] = pos[i-1] + cnt[i-1];</pre>
        for(auto x:p)p_new[pos[c[x]]++] = x;
        p = p_new;
    SuffixArray(string s) : n(s.length() + 1)
        p.assign(n,0),c.assign(n,0),_lcp.assign(n,0);
        s += (char) 31;
        vector<pair<int,int>> a(n);
        for(int i=0;i<n;i++)a[i] = {s[i],i};</pre>
        sort(all(a));
        for(int i=0;i<n;i++) p[i] = a[i].second;</pre>
        c[p[0]] = 0;
        for (int i=1; i<n; i++)</pre>
            if(a[i].first == a[i-1].first)c[p[i]] = c[p[i-1]];
            else c[p[i]] = c[p[i-1]] + 1;
        for (int k=0; 1 << k < n; k++)
             for (int i=0; i<n; i++)</pre>
```

p[i] = (p[i] - (1 << k) + n) %n;

```
count_sort(p,c);
                                      vector<int> c new(n);
                                      c_{new[p[0]]} = 0;
                                      for (int i=1; i<n; i++)</pre>
                                                  pair < int, int > now = {c[p[i]], c[(p[i] + (1 << k))%}
                                                  pair < int, int > prev = {c[p[i-1]], c[(p[i-1]] + prev = {c[p[i-1]], c[(p[i-1]] + prev = {c[p[i-1]], c[(p[i-1]] + prev = {c[p[i-1]], c[(p[i-1]), c[(
                                                                   (1<<k))%n]};
                                                  if (now == prev)c_new[p[i]] = c_new[p[i-1]];
                                                   else c_new[p[i]] = c_new[p[i-1]] + 1;
                                      c = c_new;
                         for (int i=0, k=0; i<n-1; i++)</pre>
                                      int j = p[c[i]-1];
                                      while (s[j + k] == s[i + k])k++;
                                      _{lcp[c[i] - 1] = k;}
                                      k&&k--;
                         st = new SparseTable(_lcp);
            int lcp(int i, int j)
                         if(i == j) return n - 1 - i;
                         int 1 = c[i];
                         int r = c[j];
                         if(1 > r)swap(1,r);
                         return st->query_val(1,r - 1);
};
SuffixAutomaton.h
Description: sa st 0 is first state. Transition from i using sa st i next c<sub>ines</sub>
struct SAState
            int len, link;
            map<char, int> next;
};
struct SuffixAutomaton
             vector<SAState> st:
            int sz, last;
             SuffixAutomaton()
                         sz = 1;
                         st.resize(sz);
                         st[0].len = 0;
                         st[0].link = -1;
                         last = 0;
             void sa_extend(char c)
                         int cur = sz++;
                         st.resize(sz);
                         st[cur].len = st[last].len + 1;
                         int p = last;
                         while (p != -1 && !st[p].next.count(c))
                                      st[p].next[c] = cur;
```

p = st[p].link;

st[cur].link = 0;

int q = st[p].next[c];

st[cur].link = q;

int clone = sz++;

st.resize(sz);

if (st[p].len + 1 == st[q].len)

st[clone].len = st[p].len + 1;

st[p].next[c] = clone; p = st[p].link;

st[q].link = st[cur].link = clone;

while (p != -1 && st[p].next[c] == q)

st[clone].next = st[q].next;

st[clone].link = st[q].link;

if (p == -1)

else

else

Hashing AhoCorasick IntervalContainer IntervalCover

```
last = cur;
};
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.
typedef uint64_t ull;
struct H {
  ull x; H(ull x=0) : x(x) \{ \}
  H operator+(H o) { return x + o.x + (x + o.x < x); }
  H operator-(H o) { return *this + ~o.x; }
  H operator*(H o) { auto m = (__uint128_t)x * o.x;
   return H((ull)m) + (ull)(m >> 64); }
  ull get() const { return x + !~x; }
  bool operator==(H o) const { return get() == o.get(); }
  bool operator<(H o) const { return get() < o.get(); }</pre>
static const H C = (11)1e11+3; // (order \sim 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)
   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{}}; for(char c:s) h=h*C+c;return h;}
```

AhoCorasick.h

Description: Aho-Corasick automaton, used for multiple pattern matching. Initialize with AhoCorasick ac(patterns); the automaton start node will be at index 0. find(word) returns for each position the index of the longest word that ends there, or -1 if none. findAll(-, 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. For large alphabets, split each symbol into chunks, with sentinel bits for symbol boundaries.

Time: construction takes $\mathcal{O}(26N)$, where N = sum of length of patterns. find(x) is $\mathcal{O}(N)$, where N = length of x. findAll is $\mathcal{O}(NM)$.

```
struct AhoCorasick {
 enum {alpha = 26, first = 'A'}; // change this!
 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;
 vi backp;
 void insert(string& s, int j) {
   assert(!s.emptv());
   int n = 0;
   for (char c : s) {
     int& m = N[n].next[c - first];
     if (m == -1) { n = m = sz(N); N.emplace_back(-1); }
     else n = m;
   if (N[n].end == -1) N[n].start = j;
   backp.push_back(N[n].end);
   N[n].end = j;
   N[n].nmatches++;
 AhoCorasick(vector<string>& pat) : N(1, -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[y].nmatches;
         q.push(ed);
 vi find(string word) {
   int n = 0;
   vi res; // ll count = 0;
```

```
for (char c : word) {
    n = N[n].next[c - first];
    res.push_back(N[n].end);
    // count \neq = 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;
```

Various (10)

10.1 Intervals

IntervalContainer.h

Description: Add and remove intervals from a set of disjoint intervals. Will merge the added interval with any overlapping intervals in the set when adding. Intervals are [inclusive, exclusive).

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

```
edce47, 23 lines
set<pii>::iterator addInterval(set<pii>& is, int L, int R) {
 if (L == R) return is.end();
  auto it = is.lower bound({L, R}), before = it;
  while (it != is.end() && it->first <= R) {</pre>
    R = max(R, it->second);
    before = it = is.erase(it);
 if (it != is.begin() && (--it)->second >= L) {
   L = min(L, it->first);
    R = max(R, it->second);
    is.erase(it);
 return is.insert(before, {L,R});
void removeInterval(set<pii>& is, int L, int R) {
 if (L == R) return;
 auto it = addInterval(is, L, R);
 auto r2 = it->second;
 if (it->first == L) is.erase(it);
 else (int&)it->second = L;
 if (R != r2) is.emplace(R, r2);
```

IntervalCover.h

Description: Compute indices of smallest set of intervals covering another interval. Intervals should be [inclusive, exclusive). To support [inclusive, inclusive, change (A) to add | R.empty(). Returns empty set on failure (or if G is empty). Time: $\mathcal{O}(N \log N)$

9e9d8d, 19 lines

```
template<class T>
vi cover(pair<T, T> G, vector<pair<T, T>> I) {
 vi S(sz(I)), R;
 iota(all(S), 0);
 sort(all(S), [&](int a, int b) { return I[a] < I[b]; });</pre>
 T cur = G.first;
```

```
int at = 0;
while (cur < G.second) { // (A)
  pair<T, int> mx = make_pair(cur, -1);
  while (at < sz(I) && I[S[at]].first <= cur) {
    mx = max(mx, make_pair(I[S[at]].second, S[at]));
    at++;
  }
  if (mx.second == -1) return {};
  cur = mx.first;
  R.push_back(mx.second);
}
return R;</pre>
```

ConstantIntervals.h

Description: Split a monotone function on [from, to) into a minimal set of half-open intervals on which it has the same value. Runs a callback g for each such interval.

```
 \begin{array}{ll} \textbf{Usage:} & \text{constantIntervals}(0, \text{sz(v), [\&](int x)}\{\text{return v[x];}\}, \\ \text{[\&](int lo, int hi, T val)}\{\ldots\}); \\ \textbf{Time:} & \mathcal{O}\left(k\log\frac{n}{k}\right) \end{array}
```

```
template < class F, class G, class T>
void rec(int from, int to, F& f, G& q, int& i, T& p, T q) {
 if (p == q) return;
  if (from == to) {
   g(i, to, p);
    i = to; p = q;
  } else {
   int mid = (from + to) >> 1;
   rec(from, mid, f, g, i, p, f(mid));
   rec(mid+1, to, f, g, i, p, q);
template < class F, class G>
void constantIntervals(int from, int to, F f, G g) {
 if (to <= from) return;</pre>
  int i = from; auto p = f(i), q = f(to-1);
 rec(from, to-1, f, g, i, p, q);
 g(i, to, q);
```

10.2 Misc. algorithms

TernarySearch.h

Description: Find the smallest i in [a,b] that maximizes f(i), assuming that $f(a) < \ldots < f(i) \ge \cdots \ge f(b)$. To reverse which of the sides allows non-strict inequalities, change the < marked with (A) to <=, and reverse the loop at (B). To minimize f, change it to >, also at (B).

Usage: int ind = ternSearch(0,n-1,[&](int i){return a[i];}); Time: $\mathcal{O}(\log(b-a))$

```
template < class F>
int ternSearch(int a, int b, F f) {
   assert(a <= b);
   while (b - a >= 5) {
      int mid = (a + b) / 2;
      if (f(mid) < f(mid+1)) a = mid; // (A)
      else b = mid+1;
   }
   rep(i,a+1,b+1) if (f(a) < f(i)) a = i; // (B)
   return a;
}</pre>
```

FastKnapsack.h

Description: Given N non-negative integer weights w and a non-negative target t, computes the maximum $S \le t$ such that S is the sum of some subset of the weights.

```
Time: \mathcal{O}(N \max(w_i)) b20ccc, 16 lines
```

```
int knapsack(vi w, int t) {
   int a = 0, b = 0, x;
   while (b < sz(w) && a + w[b] <= t) a += w[b++];
   if (b == sz(w)) return a;
   int m = *max_element(all(w));
   vi u, v(2*m, -1);
   v[a+m-t] = b;
   rep(i,b,sz(w)) {
      u = v;
      rep(x,0,m) v[x+w[i]] = max(v[x+w[i]], u[x]);
      for (x = 2*m; --x > m;) rep(j, max(0,u[x]), v[x])
      v[x-w[j]] = max(v[x-w[j]], j);
   }
   for (a = t; v[a+m-t] < 0; a--);
   return a;
}</pre>
```

10.3 Dynamic programming

KnuthDF

Description: When doing DP on intervals: $a[i][j] = \min_{i < k < j} (a[i][k] + a[k][j]) + f(i,j)$, 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 if $f(b,c) \le f(a,d)$ and $f(a,c) + f(b,d) \le f(a,d) + f(b,c)$ for all $a \le b \le c \le d$. Consider also: LineContainer (ch. Data structures), monotone queues, ternary search. **Time:** $\mathcal{O}\left(N^2\right)$

DivideAndConquerDP.h

Description: Given $a[i] = \min_{lo(i) \le k < hi(i)} (f(i, k))$ where the (minimal) optimal k increases with i, computes a[i] for i = L..R - 1.

```
Time: \mathcal{O}((N + (hi - lo)) \log N)
                                                      d38d2b, 18 lines
struct DP { // Modify at will:
 int lo(int ind) { return 0; }
 int hi(int ind) { return ind; }
 11 f(int ind, int k) { return dp[ind][k]; }
 void store(int ind, int k, ll v) { res[ind] = pii(k, v); }
 void rec(int L, int R, int LO, int HI) {
   if (L >= R) return;
   int mid = (L + R) >> 1;
   pair<11, int> best (LLONG_MAX, LO);
    rep(k, max(LO, lo(mid)), min(HI, hi(mid)))
     best = min(best, make_pair(f(mid, k), k));
    store(mid, best.second, best.first);
    rec(L, mid, LO, best.second+1);
    rec(mid+1, R, best.second, HI);
 void solve(int L, int R) { rec(L, R, INT_MIN, INT_MAX); }
```

10.4 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).

| 10.5 Optimization tricks

__builtin_ia32_ldmxcsr(40896); disables denormals (which make floats 20x slower near their minimum value).

10.5.1 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.

10.5.2 Pragmas

- #pragma GCC optimize ("ofast") will make GCC auto-vectorize loops and optimizes floating points better.
- #pragma GCC target ("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).

FastMod.h

Description: Compute a%b about 5 times faster than usual, where b is constant but not known at compile time. Returns a value congruent to $a \pmod{b}$ in the range [0,2b).

```
typedef unsigned long long ull;
struct FastMod {
  ull b, m;
  FastMod(ull b) : b(b), m(-1ULL / b) {}
  ull reduce(ull a) { // a % b + (0 or b)
    return a - (ull)((_uint128_t(m) * a) >> 64) * b;
  }
};
```

Techniques (A)

techniques.txt

Combinatorics

159 lines

Recursion Divide and conquer Finding interesting points in N log N Algorithm analysis Master theorem Amortized time complexity Greedy algorithm Scheduling Max contiquous subvector sum Invariants Huffman encoding Graph theory Dynamic graphs (extra book-keeping) Breadth first search Depth first search * Normal trees / DFS trees Dijkstra's algorithm MST: Prim's algorithm Bellman-Ford Konig's theorem and vertex cover Min-cost max flow Lovasz toggle Matrix tree theorem Maximal matching, general graphs Hopcroft-Karp Hall's marriage theorem Graphical sequences Floyd-Warshall Euler cycles Flow networks * Augmenting paths * Edmonds-Karp Bipartite matching Min. path cover Topological sorting Strongly connected components Cut vertices, cut-edges and biconnected components Edge coloring * Trees Vertex coloring * Bipartite graphs (=> trees) * 3^n (special case of set cover) Diameter and centroid K'th shortest path Shortest cycle Dynamic programming Knapsack Coin change Longest common subsequence Longest increasing subsequence Number of paths in a dag Shortest path in a dag Dynprog over intervals Dynprog over subsets Dynprog over probabilities Dynprog over trees 3^n set cover Divide and conquer Knuth optimization Convex hull optimizations RMQ (sparse table a.k.a 2^k-jumps) Bitonic cycle Log partitioning (loop over most restricted)

Computation of binomial coefficients Pigeon-hole principle Inclusion/exclusion Catalan number Pick's theorem Number theory Integer parts Divisibility Euclidean algorithm Modular arithmetic * Modular multiplication * Modular inverses * Modular exponentiation by squaring Chinese remainder theorem Fermat's little theorem Euler's theorem Phi function Frobenius number Ouadratic reciprocity Pollard-Rho Miller-Rabin Hensel lifting Vieta root jumping Game theory Combinatorial games Game trees Mini-max Nim Games on graphs Games on graphs with loops Grundy numbers Bipartite games without repetition General games without repetition Alpha-beta pruning Probability theory Optimization Binary search Ternary search Unimodality and convex functions Binary search on derivative Numerical methods Numeric integration Newton's method Root-finding with binary/ternary search Golden section search Matrices Gaussian elimination Exponentiation by squaring Sorting Radix sort Geometry Coordinates and vectors * Cross product * Scalar product Convex hull Polygon cut Closest pair Coordinate-compression Ouadtrees KD-trees All segment-segment intersection Sweeping Discretization (convert to events and sweep) Angle sweeping Line sweeping Discrete second derivatives Strings Longest common substring Palindrome subsequences

Knuth-Morris-Pratt Tries Rolling polynomial hashes Suffix array Suffix tree Aho-Corasick Manacher's algorithm Letter position lists Combinatorial search Meet in the middle Brute-force with pruning Best-first (A*) Bidirectional search Iterative deepening DFS / A* Data structures LCA (2^k-jumps in trees in general) Pull/push-technique on trees Heavy-light decomposition Centroid decomposition Lazy propagation Self-balancing trees Convex hull trick (wcipeg.com/wiki/Convex_hull_trick) Monotone queues / monotone stacks / sliding queues Sliding queue using 2 stacks Persistent segment tree

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