

University of Central Florida

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UCF Locals

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Contest (1)	
template.cpp	ines
<pre>#include <bits stdc++.h=""> using namespace std; #define rep(i, a, b) for (ll i = a; i < (ll)(b); ++i) #define all(x) (x).begin(), (x).end() #define sz(x) (ll) x.size() typedef long long ll; typedef vector<ll> vll;</ll></bits></pre>	
int main() { cip tic(0) > cupc with ctdic(0).	
<pre>cin.tie(0)->sync_with_stdio(0); cin.exceptions(cin.failbit); }</pre>	
.bashre	ines
alias c='g++ -Wall -Wconversion -Wfatal-errors -g -std=c++17 -fsanitize=undefined,address' xmodmap -e 'clear lock' -e 'keycode 66=less greater' $\#caps = 1$	
hash.sh	ines
# Hashes a file, ignoring all whitespace and comments. Use for the verifying that code was correctly typed. cpp -dD -P -fpreprocessed tr -d '[:space:]' md5sum cut -d	or
troubleshoot.txt 57 1	ines
Pre-submit:	
Write a few test cases INCLUDE MAX CASES.	
Is memory fine? Overflow?	
Correct file?	
Wrong answer:	

Print your solution + Print debug Clearing all structures between test cases? Can your code handle the whole input range? Read the problem statement again. Do you handle all corner cases correctly? Have you understood the problem correctly? Uninitialized variables? Overflows? Confusing N and M, i and j, etc.? Are you sure your algorithm works? Any very special cases? Do you know how the STL functions you use work? Add some assertions, maybe resubmit. Create some testcases Go through the algorithm for a simple case. Go through this list again. Explain your algorithm to a teammate. Ask the teammate to look at your code. Go for a small walk, e.g. to the toilet. Is your output format correct? (whitespace)

Runtime error:

Have you tested all corner cases locally?
Any uninitialized variables?
Are you reading or writing outside of range?
Consider increasing the size of your vectors
Any assertions that might fail?
Any possible division by 0? (mod 0 for example)
Any possible infinite recursion?
Invalidated pointers or iterators?
Are you using too much memory?

You/Teammate rewrite solution from the start

Time limit exceeded:

Do you have any possible infinite loops? What is the complexity of your algorithm? Are you copying a lot of unnecessary data? How big is the input and output? Avoid vector, map. (use arrays/unordered_map) What do your teammates think about your algorithm?

Memory limit exceeded:

Max amount of memory your code should need?
Clearing all data structures between test cases?

Mathematics (2)

2.1 Equations

$$ax^2 + bx + c = 0 \Rightarrow x = \frac{-b \pm \sqrt{b^2 - 4ac}}{2a}$$

The extremum is given by x = -b/2a.

$$ax + by = e$$

$$cx + dy = f$$

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

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

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

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

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

2.2 Recurrences

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

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

Non-distinct roots r become polynomial factors, e.g. $a_n = (d_1n + d_2)r^n$.

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

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):

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

template .bashrc hash troubleshoot

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

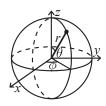
2.4.2 Quadrilaterals

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

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

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



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

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

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

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:

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

Sums 2.6

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

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

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

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

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

2.7

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

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

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

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

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

Probability theory

Let X be a discrete random variable with probability $p_X(x)$ of assuming the value x. It will then have an expected value (mean) $\mu = \mathbb{E}(X) = \sum_{x} x p_X(x)$ and variance $\sigma^2 = V(X) = \mathbb{E}(X^2) - (\mathbb{E}(X))^2 = \sum_x (x - \mathbb{E}(X))^2 p_X(x)$ where σ is the standard deviation. If X is instead continuous it will have a probability density function $f_X(x)$ and the sums above will instead be integrals with $p_X(x)$ replaced by $f_X(x)$.

Expectation is linear:

$$\mathbb{E}(aX + bY) = a\mathbb{E}(X) + b\mathbb{E}(Y)$$

For independent X and Y,

$$V(aX + bY) = a^2V(X) + b^2V(Y).$$

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), $0 \le p \le 1$.

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

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

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 b and 0 elsewhere it is U(a, b), a < b.

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

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

Exponential distribution

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

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

Normal distribution

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

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

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

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

Data structures (3)

Note: if there are no updates, consider merges

BitVector.h

Description: Given vector of bits, counts number of 0's in [0, r). Use with WaveletTree.h by using modifications in comments in that file and replacing bv[h][x] with bv[h].cnt0(x) **Time:** $\mathcal{O}(1)$ time

struct BitVector {
 vector
yector
struct BitVector {
 vector
yector
split
yector
yety
yety
split
yety
yety
yety
split
yety
yety
yety
yety
split
yety

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

- __builtin_popcountll(x & ((1ULL << (r & 63)) - 1));

HashMap.h

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

```
#include <bits/extc++.h>
// To use most bits rather than just the lowest ones:
struct chash { // large odd number for C
   const uint64_t C = l1(4e18 * acos(0)) | 71;
   l1 operator()(1l x) const { return _builtin_bswap64(x*C); }
};
__gnu_pbds::gp_hash_table<11,int,chash> h({},{},{},{},{},{1<<16});</pre>
```

KactlSegmentTree.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:** $\mathcal{O}(\log N)$

struct Tree {
 typedef 11 T;
 static constexpr T unit = LLONG_MIN;
 T f(T a, T b) { return max(a, b); }
 vector<T> s; 11 n;
 Tree(11 n = 0, T def = unit) : s(2*n, def), n(n) {}

```
void update(ll pos, T val) {
   for (s[pos += n] = val; pos /= 2;)
      s[pos] = f(s[pos * 2], s[pos * 2 + 1]);
}
T query(ll b, ll 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);
}
</pre>
```

int lo, hi, mset = inf, madd = 0, val = -inf;

LazySegmentTree.h

Node *1 = 0, *r = 0;

Description: Segment tree with ability to add or set values of large intervals, and compute max of intervals. Can be changed to other things. Use with a bump allocator for better performance, and SmallPtr or implicit indices to save memory.

```
Usage: Node* tr = new Node(v, 0, sz(v));
Time: \mathcal{O}(\log N).
```

const int inf = 1e9; struct Node {

```
Node (int lo, int hi):lo(lo), hi(hi) {} // Large interval of -inf
Node(vll& v, int lo, int hi) : lo(lo), hi(hi) {
  if (lo + 1 < hi) {
    int mid = 10 + (hi - 10)/2;
   1 = new Node(v, lo, mid); r = new Node(v, mid, hi);
    val = max(1->val, r->val);
  else val = v[lo];
int query(int L, int R) {
 if (R <= lo || hi <= L) return -inf;
  if (L <= lo && hi <= R) return val;
  push();
  return max(l->query(L, R), r->query(L, R));
void set(int L, int R, int x) {
  if (R <= lo || hi <= L) return;
  if (L <= lo && hi <= R) mset = val = x, madd = 0;
    push(), l->set(L, R, x), r->set(L, R, x);
    val = max(1->val, r->val);
void add(int L, int R, int x) {
  if (R <= lo || hi <= L) return;
  if (L <= lo && hi <= R) {
    if (mset != inf) mset += x;
    else madd += x:
    val += x;
    push(), 1->add(L, R, x), r->add(L, R, x);
    val = max(1->val, r->val);
void push() {
  if (!1) {
    int mid = lo + (hi - lo)/2;
   1 = new Node(lo, mid); r = new Node(mid, hi);
  if (mset != inf)
   1->set(lo,hi,mset), r->set(lo,hi,mset), mset = inf;
  else if (madd)
```

NodeBasedSegmentTree.h

Description: Simple node based segment tree for quick typing on more unique problems

9aa4ab, 23 lines

```
struct node {
   pll val = {LLONG_MAX, LLONG_MAX};
    ll lo, hi, mid;
   node *left = nullptr, *right = nullptr;
    node (11 1, 11 h) : lo(1), hi(h) {
       mid = (lo + hi) / 2;
       if (lo + 1 == hi) return;
       left = new node(lo, mid);
       right = new node (mid, hi);
   void merge() { val = min(left->val, right->val); }
    void set(ll i, ll x) {
       if (lo + 1 == hi) { val = {x, i}; return; }
       if (i < mid) left->set(i, x);
       else right->set(i, x);
       merge();
   pll query(ll 1, ll r) {
       if (r <= lo || l >= hi) return { LLONG_MAX, LLONG_MAX }
       if (lo >= 1 && hi <= r) return val;
        return min(left->query(l, r), right->query(l, r));
};
```

DynamicSegmentTree.h

Description: Segment Tree that only create nodes for used values. Consider using offline value compression before using a DST

248be1, 27 lines

```
struct Node {
    11 lo, hi;
    11 \text{ val} = 0;
    Node *left = nullptr, *right = nullptr;
    Node(11 lo, 11 hi) : lo(lo), hi(hi) {}
    void extend() {
        if (!left && lo + 1 < hi) {
            11 t = (10 + hi) / 2;
            left = new Node(lo, t);
            right = new Node(t, hi);
    void add(ll k, ll x) {
        extend();
        val += x;
       if (left) {
            if (k < left->hi) left->add(k, x);
            else right->add(k, x);
    11 get_sum(ll lq, ll rq) {
        if (lq <= lo && hi <= rq) return val;
        if (max(lo, lq) >= min(hi, rq)) return 0;
        return left->get_sum(lq, rq) + right->get_sum(lq, rq);
};
```

56286c, 21 lines

c59ada, 13 lines

```
SegmentTree.h
```

Description: General Segment Tree with functionality for common prob-

```
Usage: set(i, v, 0, 0, SIZE)
                                                     ff3d11, 55 lines
typedef struct Tree {
   ll size;
   vll vals;
   11 NEUTRAL_ELEMENT = 0;
   11 f(11 a, 11 b) { return a + b; }
   void init(ll n) {
       size = 1:
       while (size < n) size \star= 2;
       vals.assign(2 * size, OLL);
   void build(vll &a, ll x, ll lx, ll rx) {
       if (1x + 1 == rx) {
           if (lx < (ll)a.size()) vals[x] = a[lx];</pre>
       11 m = (1x + rx) / 2;
       build(a, 2 * x + 1, 1x, m);
       build(a, 2 * x + 2, m, rx);
       vals[x] = f(vals[2 * x + 1], vals[2 * x + 2]);
   void set(ll i, ll v, ll x, ll lx, ll rx) {
       if (1x + 1 == rx) {
           vals[x] = v;
           return:
       11 m = (1x + rx) / 2;
       if (i < m) set(i, v, 2 * x + 1, 1x, m);
       else set(i, v, 2 * x + 2, m, rx);
       vals[x] = f(vals[2 * x + 1], vals[2 * x + 2]);
   11 calc(11 1, 11 r, 11 x, 11 1x, 11 rx) {
       if (lx >= r || l >= rx) return NEUTRAL_ELEMENT;
       if (lx >= 1 && rx <= r) return vals[x];</pre>
       11 m = (1x + rx) / 2;
       11 s1 = calc(1, r, 2 * x + 1, 1x, m);
       11 \ s2 = calc(1, r, 2 * x + 2, m, rx);
       return f(s1, s2);
   ll findKth(ll k, ll x, ll lx, ll rx) {
       if (lx + 1 == rx) return lx;
       11 m = (1x + rx) / 2;
       ll sl = vals[2*x + 1];
       if (k < s1) return findKth(k, 2*x + 1, 1x, m);
       else return findKth(k-sl, 2*x + 2, m, rx);
   11 first_above(ll v, ll l, ll x, ll lx, ll rx) {
       if (vals[x] < v) return -1;
       if (rx <= 1) return -1;
       if (rx - lx == 1) return lx;
       11 m = (1x + rx) / 2;
       ll res = first_above(v, 1, 2*x + 1, 1x, m);
       if (res == -1) res = first_above(v, 1, 2*x + 2, m, rx);
        return res;
```

} Tree; PST.h

Description: Persistent segment tree with laziness **Time:** $\mathcal{O}(\log N)$ per query, $\mathcal{O}((n+q)\log n)$ memory

6e8af5, 39 lines

```
struct PST {
 PST *1 = 0, *r = 0;
  int lo, hi;
 11 \text{ val} = 0, 1\text{zadd} = 0;
  PST(vl& v, int lo, int hi) : lo(lo), hi(hi) {
```

```
if (lo + 1 < hi) {
      int mid = lo + (hi - lo)/2;
      l = new PST(v, lo, mid); r = new PST(v, mid, hi);
     val = 1->val + r->val;
    else val = v[lo];
 11 query(int L, int R) {
   if (R <= lo || hi <= L) return 0; // idempotent
    if (L <= lo && hi <= R) return val;
    return 1->query(L, R) + r->query(L, R);
 PST* add(int L, int R, ll v) {
   if (R <= lo || hi <= L) return this;
   PST *n = new PST(*this);
   if (L <= lo && hi <= R) {
     n\rightarrow val += v * (hi - lo);
     n->1zadd += v;
   } else {
     n->push();
     n->1 = n->1->add(L, R, v);
     n->r = n->r->add(L, R, v);
     n->val = n->l->val + n->r->val;
    return n;
 void push() {
   if(lzadd == 0) return;
   1 = 1 - > add(10, hi, lzadd);
   r = r->add(lo, hi, lzadd);
   lzadd = 0;
};
```

SqrtDecomp.h

Description: decomposes (l, r) range into pair of (list of fully covered blocks, list of partially covered blocks)

```
Time: \mathcal{O}\left(B+(r-l)/B\right)
```

```
799a5b, 16 lines
template<int B>
pair<vi, vector<array<int, 3>>> decomp(int 1, int r) {
 if (1/B == (r-1)/B) return \{\{\}, \{\{1/B, 1, r\}\}\};
   vi full;
  vector<array<int, 3>> subs;
  if (1%B != 0) {
    subs.push back(\{1/B, 1, (1/B+1)*B\});
    1 = subs.back()[2];
 if (r%B != 0) {
   subs.push_back({r/B, r/B*B, r});
   r = subs.back()[1];
 rep(i, 1/B, r/B) full.push_back(i);
 return {full, subs};
```

LiChao.h

Description: Creates a segment tree style data structure that supports adding a function to the set and query the min value at a given x. For any two added functions, they must intersect at most once. If queries can be floating point, consider line container instead.

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

```
bdebe2, 27 lines
struct line {
 11 m, b;
 line(ll m = 0, ll b = LLONG_MAX): m(m), b(b) {}
 11 operator() (11 x) { return m * x + b; }
```

```
struct node {
 int lo, md, hi;
 line f;
 node *left, *right;
 node(int L, int R): lo(L), md((L+R)>>1), hi(R) {
   if(lo == hi) return;
   left = new node(lo, md);
   right = new node(md+1, hi);
 void update(line q) {
   if(g(md) < f(md)) swap(f, g);
   if(lo == hi) return;
   if(f(lo) <= g(lo) && f(hi) <= g(hi)) return;
   if(f(lo) > g(lo)) left->update(g);
    else right->update(q);
 ll query(ll x) {
   if(lo == hi) return f(x);
    return min(f(x), (x \le md ? left : right) \rightarrow query(x));
};
```

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))
```

```
struct RollbackUF {
 vll e; vector<pll> 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]); }
 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 lowerright corners (half-open).

```
Usage: SubMatrix<int> m (matrix);
m.sum(0, 0, 2, 2); // top left 4 elements
Time: \mathcal{O}(N^2+Q)
```

```
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 l, 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> 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 {
    rep(i,0,N) rep(j,0,N)
     rep(k, 0, N) \ a.d[i][j] += d[i][k]*m.d[k][j];
  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
struct Line {
  mutable ll k, m, p;
 bool operator<(const Line& o) const { return k < o.k; }</pre>
  bool operator<(ll x) const { return p < x; }</pre>
struct LineContainer : multiset<Line, less<>>> {
  // (for doubles, use inf = 1/.0, div(a,b) = a/b)
  static const ll 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 -> 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(11 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:** $\mathcal{O}(\log N)$

1754b4, 53 lines

```
struct Node {
 Node *1 = 0, *r = 0;
 int val, y, c = 1;
 Node(int val) : val(val), y(rand()) {}
 void recalc();
int cnt(Node* n) { return n ? n->c : 0; }
void Node::recalc() { c = cnt(1) + cnt(r) + 1; }
template < class F > void each (Node * n, F f) {
 if (n) { each (n->1, f); f(n->val); each (n->r, f); }
pair<Node*, Node*> split(Node* n, int k) {
 if (!n) return {};
 if (cnt(n->1) >= k) \{ // "n \rightarrow val >= k" for lower_bound(k) \}
    auto [L,R] = split(n->1, k);
    n->1 = R;
    n->recalc();
    return {L, n};
    auto [L,R] = split(n->r, k - cnt(n->1) - 1); // and just "k"
    n->r = L;
    n->recalc();
    return {n, R};
Node* merge(Node* 1, Node* r) {
 if (!1) return r:
 if (!r) return 1;
  if (1->y > r->y) {
   1->r = merge(1->r, r);
    return 1->recalc(), 1;
  } else {
    r->1 = merge(1, r->1);
    return r->recalc(), r;
Node* ins(Node* t, Node* n, int pos) {
 auto [1,r] = split(t, pos);
  return merge(merge(l, n), r);
// Example application: move the range [l, r) to index k
void move(Node*& t, int 1, int r, int k) {
  Node *a, *b, *c;
  tie(a,b) = split(t, 1); tie(b,c) = split(b, r - 1);
 if (k \le 1) t = merge(ins(a, b, k), c);
 else t = merge(a, ins(c, b, k - r));
```

FenwickTree.

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 value.

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

be8dea, 22 lii

510c32, 16 lines

```
struct FT {
 vll s:
 FT(int n) : s(n) {}
 void update(int pos, 11 dif) { // a[pos] \neq = dif
    for (; pos < sz(s); pos |= pos + 1) s[pos] += dif;
 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 \geq 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 \le sz(s) \&\& s[pos + pw-1] \le 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"
struct FT2 {
  vector<vll> ys; vector<FT> ft;
  FT2(int limx) : ys(limx) {}
  void fakeUpdate(int x, int y) {
    for (; x < sz(ys); x |= x + 1) ys[x].push_back(y);
  void init() {
    for (vll& 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;
};
```

RMQ.h

Description: Range Minimum Queries on an array. Returns $\min(V[a], V[a+1], \dots V[b-1])$ in constant time.

Usage: RMQ rmq(values);

rmq.query(inclusive, exclusive); Time: $\mathcal{O}(|V|\log|V|+Q)$

template<class T>
struct RMQ {
 vector<vector<T>> jmp;
 RMQ(const vector<T>& V) : jmp(1, V) {

MoQueries PQUpdate WaveletTree Polynomial PolyRoots

```
for (int pw = 1, k = 1; pw * 2 <= sz(V); pw *= 2, ++k) {
    jmp.emplace_back(sz(V) - pw * 2 + 1);
    rep(j,0,sz(jmp[k]))
        jmp[k][j] = min(jmp[k - 1][j], jmp[k - 1][j + pw]);
}

T query(int a, int b) {
    assert(a < b); // or return inf if a == b
    int dep = 31 - _builtin_clz(b - a);
    return min(jmp[dep][a], jmp[dep][b - (1 << dep)]);
};</pre>
```

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))
  sort(all(s), [\&](int s, int t) \{ return K(Q[s]) < K(Q[t]); \});
  for (int qi : s) {
   pii q = O[qi];
    while (L > q.first) add(--L, 0);
   while (R < q.second) add(R++, 1);
    while (L < q.first) del(L++, 0);
    while (R > q.second) del(--R, 1);
    res[gi] = calc();
  return res:
vi moTree(vector<array<int, 2>> 0, 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 = Q[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] \le L[a] \&\& R[a] \le 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;
```

PQUpdate.h

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

Description: T: value/update type. DS: Stores T. Same semantics as std::priority_queue.

```
template<class T, class DS, class Compare = less<T>>
struct POUpdate {
 DS inner:
 multimap<T, int, Compare> rev_upd;
 using iter = decltype(rev_upd)::iterator;
 POUpdate(DS inner, Compare comp={}):
   inner(inner), rev_upd(comp) {}
 bool empty() { return st.empty(); }
 const T& top() { return rev_upd.rbegin()->first; }
 void push(T value) {
   inner.push(value);
   st.push back(rev_upd.insert({value, sz(st)}));
 void pop() {
   vector<iter> extra;
   iter curr = rev upd.end();
   int min ind = sz(st);
     extra.push back (--curr);
     min ind = min(min ind, curr->second);
   } while (2*sz(extra) < sz(st) - min ind);</pre>
   while (sz(st) > min ind) {
     if (rev_upd.value_comp()(*st.back(), *curr))
       extra.push back(st.back());
     inner.pop(); st.pop_back();
   rev_upd.erase(extra[0]);
   for (auto it : extra | views::drop(1) | views::reverse) {
     it->second = sz(st);
     inner.push(it->first);
     st.push_back(it);
```

WaveletTree.h

};

Description: kth: finds k+1th smallest number in [l,r), count: rank of k (how many < k) in [l,r). Doesn't support negative numbers, and requires a[i] <= maxval. Use BitVector to make 1.6x faster and 4x less memory.

```
Time: \mathcal{O}(\log MAX)
                                                      11aee1, 38 lines
struct WaveletTree {
 int n; vvi bv; // vector<BitVector> bv;
 WaveletTree(vl a, ll max_val):
   n(sz(a)), bv(1+__lg(max_val), \{\{\}\}) {
   v1 nxt(n):
    for (int h = sz(bv); h--;) {
      vector<bool> b(n);
      rep(i, 0, n) b[i] = ((a[i] >> h) & 1);
     bv[h] = vi(n+1); // bv[h] = b;
      rep(i, 0, n) bv[h][i+1] = bv[h][i] + !b[i]; // delete
      array it{begin(nxt), begin(nxt) + bv[h][n]};
      rep(i, 0, n) *it[b[i]]++ = a[i];
      swap(a, nxt);
 11 kth(int 1, int r, int k) {
   11 \text{ res} = 0;
   for (int h = sz(bv); h--;) {
     int 10 = bv[h][1], r0 = bv[h][r];
     if (k < r0 - 10) 1 = 10, r = r0;
```

k -= r0 - 10, res |= 1ULL << h,

```
1 += bv[h][n] - 10, r += bv[h][n] - r0;
}
return res;
}
int count(int 1, int r, 11 ub) {
  int res = 0;
  for (int h = sz(bv); h--;) {
    int 10 = bv[h][1], r0 = bv[h][r];
    if ((~ub >> h) & 1) 1 = 10, r = r0;
    else
      res += r0 - 10, 1 += bv[h][n] - 10,
      r += bv[h][n] - r0;
}
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: $\mathcal{O}(n^2 \log(1/\epsilon))$

```
"Polynomial.h"
                                                     b00bfe, 23 lines
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_i
  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 ^ (p(h) > 0)) {
      rep(it, 0, 60) { // while (h - l > 1e-8)
       double m = (1 + h) / 2, f = p(m);
       if ((f <= 0) ^ sign) 1 = m;
        else h = m;
      ret.push_back((1 + h) / 2);
 return ret;
```

PolyInterpolate.h

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

08bf48, 13 lines

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

BerlekampMassey.h

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

Usage: berlekampMassey($\{0, 1, 1, 3, 5, \overline{11}\}$) // $\{1, 2\}$ Time: $\mathcal{O}(N^2)$

```
53af5a, 20 lines
"../number-theory/ModPow.h"
vll berlekampMassev(vll s) {
 int n = sz(s), L = 0, m = 0;
  vll 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 $S[i] = \sum_j S[i-j-1]tr[j]$, given $S[0... \ge n-1]$ and tr[0...n-1]. Faster than matrix multiplication. Useful together with Berlekamp-Massey. Usage: linearRec($\{0, 1\}, \{1, 1\}, k$) // k'th Fibonacci number

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

```
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 * 2 + 1);
   rep(i,0,n+1) rep(j,0,n+1)
   res[i + j] = (res[i + j] + a[i] * b[j]) % mod;
   for (int i = 2 * n; i > n; --i) rep(j,0,n)
      res[i - 1 - j] = (res[i - 1 - j] + res[i] * tr[j]) % mod;
```

```
res.resize(n + 1);
  return res;
};

Poly pol(n + 1), e(pol);
pol[0] = e[1] = 1;

for (++k; k; k /= 2) {
   if (k % 2) pol = combine(pol, e);
    e = combine(e, e);
}

ll res = 0;
rep(i,0,n) res = (res + pol[i + 1] * S[i]) % mod;
return res;
}
```

4.2 Optimization

GoldenSectionSearch.h

Description: Finds the argument minimizing the function f in the interval [a,b] assuming f is unimodal on the interval, i.e. has only one local minimum 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; }
double xmin = gss(-1000,1000,func);
```

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

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

HillClimbing.h

Description: Poor man's optimization for unimodal functions_{Seceeaf, 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 = le9; jmp > le-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

f4e444, 26 lines

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

4756fc, 7 lines

```
template<class F> double quad(double a, double b, F f, const int n = 1000) {
```

```
double h = (b - a) / 2 / n, v = f(a) + f(b);
rep(i,1,n*2)
  v += f(a + i*h) * (i&1 ? 4 : 2);
return v * h / 3;
}
```

IntegrateAdaptive.h

```
Description: Fast integration using an adaptive Simpson's rule.
Usage: double sphereVolume = quad(-1, 1, [](double x) {
return quad(-1, 1, [&] (double y) {
return quad(-1, 1, [&] (double z)
return x*x + y*y + z*z < 1; {);};};
                                                     92dd79, 15 lines
typedef double d;
#define S(a,b) (f(a) + 4*f((a+b) / 2) + f(b)) * (b-a) / 6
template <class F>
d rec(F& f, da, db, deps, dS) {
  dc = (a + b) / 2;
  d S1 = S(a, c), S2 = S(c, b), T = S1 + S2;
  if (abs(T - S) \le 15 * eps | | b - a < 1e-10)
    return T + (T - S) / 15;
  return rec(f, a, c, eps / 2, S1) + rec(f, c, b, eps / 2, S2);
template<class F>
d quad(d a, d b, F f, d eps = 1e-8) {
 return rec(f, a, b, eps, S(a, b));
```

RungeKutta.h

Description: Numerically approximates the solution to a system of Differential Equations

```
template<class F, class T>
T solveSystem(F f, T x, double time, int iters) {
  double h = time / iters;
  for(int iter = 0; iter < iters; iter++) {
    T k1 = f(x);
    A k2 = f(x + 0.5 * h * k1);
    A k3 = f(x + 0.5 * h * k2);
    A k4 = f(x + h * k3);
    x = x + h / 6.0 * (k1 + 2.0 * k2 + 2.0 * k3 + k4);
  }
  return x;
}</pre>
```

Simplex.h

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

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

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

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

const T eps = le-8, inf = 1/.0;
#define MP make_pair
#define ltj(X) if(s == -1 || MP(X[j],N[j]) < MP(X[s],N[s])) s=j

struct LPSolver {
  int m, n;</pre>
```

```
vi N, B;
vvd D:
LPSolver(const vvd& A, const vd& b, const vd& c) :
  m(sz(b)), n(sz(c)), N(n+1), B(m), D(m+2), vd(n+2)) {
    rep(i, 0, m) rep(j, 0, n) D[i][j] = A[i][j];
    rep(i,0,m) { B[i] = n+i; D[i][n] = -1; D[i][n+1] = b[i];}
    rep(j,0,n) \{ N[j] = j; D[m][j] = -c[j]; \}
    N[n] = -1; D[m+1][n] = 1;
void pivot(int r, int s) {
  T *a = D[r].data(), inv = 1 / a[s];
  rep(i, 0, m+2) if (i != r \&\& abs(D[i][s]) > eps) {
   T *b = D[i].data(), inv2 = b[s] * inv;
    rep(j, 0, n+2) b[j] -= a[j] * inv2;
   b[s] = a[s] * inv2;
  rep(j, 0, n+2) if (j != s) D[r][j] *= inv;
  rep(i, 0, m+2) if (i != r) D[i][s] *= -inv;
  D[r][s] = inv;
  swap(B[r], N[s]);
bool simplex(int phase) {
  int x = m + phase - 1;
  for (;;) {
   int s = -1;
    rep(j,0,n+1) if (N[j] != -phase) ltj(D[x]);
    if (D[x][s] >= -eps) return true;
    int r = -1;
    rep(i,0,m) {
     if (D[i][s] <= eps) continue;</pre>
      if (r == -1 \mid | 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;
```

4.3 Matrices

Determinant.h

Description: Calculates determinant using modular arithmetics. Modulos can also be removed to get a pure-integer version.

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

```
3313dc, 18 lines
const 11 mod = 12345;
11 det(vector<vector<11>>& a) {
  int n = sz(a); ll ans = 1;
  rep(i,0,n) {
```

```
rep(j,i+1,n) {
    while (a[j][i] != 0) { // gcd step
      11 t = a[i][i] / a[j][i];
      if (t) rep(k,i,n)
        a[i][k] = (a[i][k] - a[j][k] * t) % mod;
      swap(a[i], a[j]);
      ans \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: $\mathcal{O}\left(n^2m\right)$ 44c9ab, 38 lines

```
typedef vector<double> vd;
const double eps = 1e-12;
int solveLinear(vector<vd>& A, vd& b, vd& x) {
 int n = sz(A), m = sz(x), rank = 0, br, bc;
 if (n) assert(sz(A[0]) == m);
 vi col(m); iota(all(col), 0);
 rep(i,0,n) {
   double v, bv = 0;
   rep(r,i,n) rep(c,i,m)
     if ((v = fabs(A[r][c])) > bv)
       br = r, bc = c, bv = v;
   if (bv <= eps) {
     rep(j,i,n) if (fabs(b[j]) > eps) return -1;
    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[i][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:; }
```

SolveLinearBinarv.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: $\mathcal{O}\left(n^2m\right)$

```
typedef bitset<1000> bs;
int solveLinear(vector<bs>& A, vi& b, bs& x, int m) {
 int n = sz(A), rank = 0, br;
 assert(m \le sz(x));
 vi col(m); iota(all(col), 0);
 rep(i,0,n) {
   for (br=i; br<n; ++br) if (A[br].any()) break;
   if (br == n) {
     rep(j,i,n) if(b[j]) return -1;
    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-mod.h

Description: Invert matrix A modulo a prime. Returns rank; result is stored in A unless singular (rank < n). 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: $\mathcal{O}\left(n^3\right)$

```
"../number-theory/ModPow.h"
                                                     0b7b13, 37 lines
int matInv(vector<vector<11>>& A) {
 int n = sz(A); vi col(n);
 vector<vector<ll>> tmp(n, vector<ll>(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 (A[j][k]) {
     r = j; c = k; goto found;
    return i;
found:
    A[i].swap(A[r]); tmp[i].swap(tmp[r]);
      swap(A[j][i], A[j][c]), swap(tmp[j][i], tmp[j][c]);
    swap(col[i], col[c]);
    11 v = modpow(A[i][i], mod - 2);
    rep(j,i+1,n) {
     11 f = A[j][i] * v % mod;
```

```
A[j][i] = 0;
   rep(k, i+1, n) A[j][k] = (A[j][k] - f*A[i][k]) % mod;
   rep(k, 0, n) tmp[j][k] = (tmp[j][k] - f*tmp[i][k]) % mod;
  rep(j, i+1, n) A[i][j] = A[i][j] * v % mod;
 rep(j, 0, n) tmp[i][j] = tmp[i][j] * v % mod;
 A[i][i] = 1;
for (int i = n-1; i > 0; --i) rep(j, 0, i) {
 ll v = A[j][i];
 rep(k,0,n) tmp[j][k] = (tmp[j][k] - v*tmp[i][k]) % mod;
rep(i,0,n) rep(j,0,n)
 A[col[i]][col[j]] = tmp[i][j] % mod + (tmp[i][j] < 0) *mod;
return n;
```

Tridiagonal.h

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

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

This is useful for solving problems on the type

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

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

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

Fails if the solution is not unique.

If $|d_i| > |p_i| + |q_{i-1}|$ for all i, or $|d_i| > |p_{i-1}| + |q_i|$, or the matrix is positive definite, the algorithm is numerically stable and neither tr nor the check for diag[i] == 0 is needed.

```
Time: \mathcal{O}(N)
                                                      8f9fa8, 26 lines
typedef double T:
vector<T> tridiagonal(vector<T> diag, const vector<T>& super,
    const vector<T>& sub, vector<T> b) {
  int n = sz(b); vi tr(n);
  rep(i, 0, n-1) {
    if (abs(diag[i]) < 1e-9 * abs(super[i])) { // diag[i] == 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;
```

JacobianMatrix.h

```
Description: Makes Jacobian Matrix using finite differences 75dc90, 15 lines
template<class F, class T>
vector<vector<T>> makeJacobian(F &f, vector<T> &x) {
 int n = sz(x);
  vector<vector<T>> J(n, vector<T>(n));
  vector < T > fX0 = f(x);
  rep(i, 0, n) {
    x[i] += eps;
    vector<T> fX1 = f(x);
    rep(j, 0, n){
      J[j][i] = (fX1[j] - fX0[j]) / eps;
    x[i] -= eps;
  return J;
```

NewtonsMethod.h

Description: Solves a system on non-linear equations

```
jacobianMatrix.h
                                                       6af945, 10 lines
template<class F. class T>
void solveNonlinear(F f, vector<T> &x){
 int n = sz(x);
 rep(iter, 0, 100) {
   vector<vector<T>> J = makeJacobian(f, x);
   matInv(J);
   vector < T > dx = J * f(x);
   x = x - dx;
```

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})
                                                                         00ced6, 35 lines
```

```
typedef complex<double> C:
typedef vector<double> vd;
void fft(vector<C>& a) {
 int n = sz(a), L = 31 - _builtin_clz(n);
  static vector<complex<long double>> R(2, 1);
  static vector<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 - \underline{\quad} builtin_clz(sz(res)), n = 1 << L;
  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 \star = x;
rep(i, 0, n) out[i] = in[-i & (n - 1)] - conj(in[i]);
rep(i, 0, sz(res)) res[i] = imag(out[i]) / (4 * n);
return res;
```

FastFourierTransformMod.h

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

```
Time: \mathcal{O}(N \log N), where N = |A| + |B| (twice as slow as NTT or FFT)
"FastFourierTransform.h"
typedef vector<ll> vl;
template<int M> vl convMod(const vl &a, const vl &b) {
 if (a.empty() || b.empty()) return {};
  vl res(sz(a) + sz(b) - 1);
  int B=32-__builtin_clz(sz(res)), n=1<<B, cut=int(sqrt(M));</pre>
  vector<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 bv = 11(imag(out1[i])+.5) + 11(real(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^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"
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> vl;
void ntt(vl &a) {
 int n = sz(a), L = 31 - _builtin_clz(n);
 static vl rt(2, 1);
 for (static int k = 2, s = 2; k < n; k \neq 2, s++) {
   rt.resize(n);
    11 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]]);
 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;
   ntt(out);
   return {out.begin(), out.begin() + s};
}</pre>
```

FastSubsetTransform.h

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

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

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(j,i,i+step) {
      int &u = a[j], &v = a[j + step]; tie(u, v) =
         inv ? pii(v - u, u) : pii(v, u + v); // AND
      inv ? pii(v, u - v) : pii(u + v, u); // OR
      pii(u + v, u - v);
    }
    if (inv) 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;
}</pre>
```

Number theory (5)

5.1 Modular arithmetic

ModInverse.h

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

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

ModPow.h

b83e45, 8 lines

```
const 11 mod = 1000000007; // faster if const

11 modpow(11 b, 11 e) {
    11 ans = 1;
    for (; e; b = b * b % mod, e /= 2)
        if (e & 1) ans = ans * b % mod;
    return ans;
}
```

ModLog.h

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

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

```
11 modLog(ll a, ll b, ll m) {
    ll n = (ll) sqrt(m) + 1, e = 1, f = 1, j = 1;
    unordered_map<ll, ll> A;
    while (j <= 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;
}</pre>
```

ModSum.h

 $\textbf{Description:} \ \ \text{Sums of mod'ed arithmetic progressions.}$

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

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

5c5bc5, 14 lines

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

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

```
"ModPow.h"
                                                      19a793, 24 lines
ll sqrt(ll a, ll p) {
 a = p; if (a < 0) a += p;
 if (a == 0) return 0;
 assert (modpow(a, (p-1)/2, p) == 1); // else no solution
 if (p % 4 == 3) return modpow(a, (p+1)/4, p);
  // a^{(n+3)/8} \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);
 11 b = modpow(a, s, p), g = modpow(n, s, p);
 for (;; r = m) {
   11 t = b;
   for (m = 0; m < r && t != 1; ++m)
```

```
t = t * t % p;
if (m == 0) return x;
ll gs = modpow(g, 1LL << (r - m - 1), p);
g = gs * gs % p;
x = x * gs % p;
b = b * g % p;
}</pre>
```

5.2 Primality

Eratosthenes.h

Description: Prime sieve for generating all primes up to a certain limit. isprime [i] is true iff i is a prime.

Time: lim=100'000'000 ≈ 0.8 s. Runs 30% faster if only odd indices are stored.

```
const int MAX_PR = 5e6;
bitset<MAX_PR> isprime;
vll eratosthenesSieve(int lim) {
  isprime.set(); isprime[0] = isprime[1] = 0;
  for (int i = 4; i < lim; i += 2) isprime[i] = 0;
  for (int i = 3; i*i < lim; i += 2) if (isprime[i])
    for (int j = i*i; j < lim; j += i*2) isprime[j] = 0;
  vll pr;
  rep(i,2,lim) if (isprime[i]) pr.push_back(i);
  return pr;
}
```

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" 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_ctzll(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;
}
```

FindPrimeFactors.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.

```
"ModMultL.h", "MilferRabin.h"

d8d98d, 19 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 l = factor(x), r = factor(n / x);
1.insert(1.end(), all(r));
return 1;
```

GetFactors.h

Description: Gets all factors of a number N given the prime factorization of the number.

Time: $\mathcal{O}\left(\sqrt[3]{N}\right)$

18bfb0, 5 lines void getFactors (auto &primes, auto &pows, auto &factors, int i $= 0, 11 n = 1) {$ if(i == sz(primes)) return void(factors.push_back(n)); for (11 j = 0, pow = 1; $j \le pows[i]$; j++, pow *= primes[i]) getFactors(primes, pows, factors, i+1, n * pow);

Divisibility

euclid.h

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

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

CRT.h

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: $\log(n)$

```
"euclid.h"
                                                      04d93a, 7 lines
ll crt(ll a, ll m, ll b, ll n) {
 if (n > m) swap(a, b), swap(m, n);
  ll x, y, q = euclid(m, n, x, y);
  assert((a - b) % g == 0); // else no solution
 x = (b - a) % n * x % n / g * m + a;
  return x < 0 ? x + m*n/g : x;
```

5.3.1 Bézout's identity

For $a \neq b \neq 0$, then d = qcd(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 $< n \text{ that are coprime with } n. \ \phi(1) = 1, \ p \text{ prime} \Rightarrow \phi(p^k) = (p-1)p^{k-1},$ $m, n \text{ coprime } \Rightarrow \phi(mn) = \phi(m)\phi(n).$ If $n = p_1^{k_1} p_2^{k_2} \dots p_r^{k_r}$ then $\phi(n) =$ $(p_1-1)p_1^{k_1-1}...(p_r-1)p_r^{k_r-1}. \ \phi(n)=n\cdot \prod_{p|n} (1-1/p).$ $\sum_{d|n} \phi(d) = n, \ \sum_{1 \le k \le n, \gcd(k,n)=1} k = n\phi(n)/2, n > 1$

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

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

```
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;
```

5.4 Fractions

ContinuedFractions.h

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

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

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

```
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;
   11 \lim = \min(P ? (N-LP) / P : \inf, Q ? (N-LQ) / Q : \inf),
       a = (11) floor(v), b = min(a, lim),
       NP = b*P + LP, NO = b*O + LO;
    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)) ?
        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

cf7d6d, 8 lines

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, 11 N) {
 bool dir = 1, A = 1, B = 1;
 Frac lo{0, 1}, hi{1, 1}; // Set hi to 1/0 to search (0, N)
 if (f(lo)) return lo;
 assert(f(hi));
 while (A | | B) {
   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}_{2a}^{\times} is instead isomorphic to $\mathbb{Z}_2 \times \mathbb{Z}_{2a-2}$.

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:

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

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

$$g(n) = \sum_{1 \le m \le n} f(\lfloor \frac{n}{m} \rfloor) \Leftrightarrow f(n) = \sum_{1 \le m \le n} \mu(m)g(\lfloor \frac{n}{m} \rfloor)$$

Combinatorial (6)

6.1 Permutations

6.1.1 Factorial

n	1 2 3	_		•	_		U	10
n!	1 2 6	24 1	20 72	0 504	0 403	20 36	$2880\ 3$	628800
n	11	12	13	1	4	15	16	17
$\overline{n!}$	4.0e7	′ 4.8e	8 6.2	e9 8.7	e10 1.	.3e12	2.1e13	3.6e14
n	20	25	30	40	50	100	150	171
n!	2e18	2e25	3e32	8e47	3e64	9e157	6e262	>DBL_MAX

IntPerm derangements multinomial

IntPerm.h

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

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

044568, 6 lir

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.

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

derangements.h

Description: Finds number of derangements

664116, 2 lines

$$d[0] = 1; d[1] = 0;$$

rep (i, 2, n+1) $d[i] = (i-1)*(d[i-1]+d[i-2]);$

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

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

6.2 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

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

11 multinomial(vi& v) {
 11 c = 1, m = v.empty() ? 1 : v[0];
 rep(i,1,sz(v)) rep(j,0,v[i]) c = c * ++m / (j+1);
 return c;

6.2.4 Stars and Bars Formula

The number of ways to distribute n identical objects into k distinct bins

$$\binom{n+k-1}{n} = \binom{n+k-1}{k-1}.$$

6.3 General purpose numbers

6.3.1 Bernoulli numbers

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

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

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

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

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

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, 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 : $n_1 n_2 \cdots n_k n^{k-2}$ # 3.74 departed by (number $(d_1-1)! \cdots (d_n-1)!$)

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

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

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

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

Graphs (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 $V^2 \max |w_i| < \sim 2^{63}$. **Time:** $\mathcal{O}(VE)$

830a8f, 23 lines

```
const ll 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) {
   Node cur = nodes[ed.a], &dest = nodes[ed.b];
   if (abs(cur.dist) == inf) continue;
   11 d = cur.dist + ed.w;
   if (d < dest.dist) {</pre>
     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;
```

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:** $\mathcal{O}(|V| + |E|)$

```
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);
    }
}
```

7.2 Network flow

Dinic.h

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

```
struct Dinic {
    struct Edge {
        int to, rev;
        ll c, oc;
        ll flow() { return max(oc - c, OLL); } // if you need flows
    };
    vi lvl, ptr, q;
    vector<vector<Edge>> adj;
    Dinic(int n) : lvl(n), ptr(n), q(n), adj(n) {}
    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});
    }
    ll dfs(int v, int t, ll f) {
        if (v == t || !f) return f;
        for (int& i = ptr[v]; i < sz(adj[v]); i++) {
            Edge& e = adj[v][i];
        }
    }
}</pre>
```

```
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 : adj[v])
         if (!lvl[e.to] && e.c >> (30 - L))
           q[qe++] = e.to, lvl[e.to] = lvl[v] + 1;
     while (ll p = dfs(s, t, LLONG_MAX)) flow += p;
   } while (lvl[t]);
   return flow;
 bool leftOfMinCut(int a) { return lvl[a] != 0; }
};
```

MinCostMaxFlow.h

Description: Min-cost max-flow. Negative cost cycles not supported. To obtain the actual flow, look at positive values only.

Time: Approximately $\mathcal{O}\left(E^2\right)$, actually $\mathcal{O}\left(FS\right)$ where S is the time complexity of the SSSP alg used in find path (in this case SPFA) $_{27\text{fafb},\ 55\ \text{lines}}$

```
struct mcmf {
 const 11 inf = LLONG MAX >> 2;
 struct edge {
   int v;
   11 cap, flow, cost;
 };
 int n;
 vector<edge> edges;
 vvi adj; vii par; vi in_q;
 vector<ll> dist, pi;
 mcmf(int n): n(n), adj(n), par(n), in_q(n), dist(n), pi(n) {}
 void add_edge(int u, int v, ll cap, ll cost) {
   int idx = sz(edges);
   edges.push_back({v, cap, 0, cost});
   edges.push_back({u, cap, cap, -cost});
   adj[u].push_back(idx);
   adj[v].push_back(idx ^ 1);
 bool find_path(int s, int t) {
   fill(all(dist), inf);
   fill(all(in_q), 0);
   queue<int> q; q.push(s);
   dist[s] = 0, in_q[s] = 1;
   while(!q.empty()) {
     int cur = q.front(); q.pop();
     in_q[cur] = 0;
     for(int idx: adj[cur]) {
       auto [nxt, cap, fl, wt] = edges[idx];
       11 nxtD = dist[cur] + wt;
       if(fl >= cap || nxtD >= dist[nxt]) continue;
       dist[nxt] = nxtD;
       par[nxt] = {cur, idx};
       if(in_q[nxt]) continue;
       q.push(nxt); in_q[nxt] = 1;
```

```
return dist[t] < inf;
}
pair<11, 11> calc(int s, int t) {
    11 flow = 0, cost = 0;
    while(find_path(s, t)) {
        11 f = inf;
        for(int i, u, v = t; tie(u, i) = par[v], v != s; v = u)
            f = min(f, edges[i].cap - edges[i].flow);
        flow += f;
        for(int i, u, v = t; tie(u, i) = par[v], v != s; v = u)
            edges[i].flow += f, edges[i^1].flow -= f;
    }
    rep(i, 0, sz(edges)>>1)
        cost += edges[i<<1].cost * edges[i<<1].flow;
    return {flow, cost};
}
</pre>
```

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

8b0a10 21 1

```
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:** $\mathcal{O}(V)$ Flow Computations

```
return tree:
```

UCF

MatroidIntersection.h

"../data-structures/UnionFind.h"

Description: Given two matroids, finds the largest common independent set. For the color and graph matroids, this would be the largest forest where no two edges are the same color. A matroid has 3 functions

- check(int x): returns if current matroid can add x without becoming dependent
- add(int x): adds an element to the matroid (guaranteed to never make it dependent)
- clear(): sets the matroid to the empty matroid

The matroid is given an int representing the element, and is expected to convert it (e.g. the color or the endpoints) Pass the matroid with more expensive add/clear operations to M1.

Time: $R^2N(M2.add+M1.check+M2.check)+R^3M1.add+R^2M1.clear+$ RNM2.clear

```
struct ColorMat {
  vi cnt, clr;
  ColorMat(int n, vector<int> clr) : cnt(n), clr(clr) {}
  bool check(int x) { return !cnt[clr[x]]; }
  void add(int x) { cnt[clr[x]]++; }
  void clear() { fill(all(cnt), 0); }
struct GraphMat {
 UF uf;
  vector<array<int, 2>> e;
  GraphMat(int n, vector<array<int, 2>> e) : uf(n), e(e) {}
  bool check(int x) { return !uf.sameSet(e[x][0], e[x][1]); }
  void add(int x) { uf.join(e[x][0], e[x][1]); }
  void clear() { uf = UF(sz(uf.e)); }
template <class M1, class M2> struct MatroidIsect {
 int n:
  vector<char> iset;
 M1 m1; M2 m2;
  MatroidIsect(M1 m1, M2 m2, int n) : n(n), iset(n + 1), m1(m1)
      , m2(m2) {}
  vi solve() {
    rep(i,0,n) if (m1.check(i) && m2.check(i))
     iset[i] = true, m1.add(i), m2.add(i);
    while (augment());
    vi ans;
    rep(i,0,n) if (iset[i]) ans.push_back(i);
    return ans;
  bool augment() {
    vector<int> frm(n, -1);
    queue<int> q({n}); // starts at dummy node
    auto fwdE = [&](int a) {
     vi ans;
     m1.clear();
     rep(v, 0, n) if (iset[v] \&\& v != a) m1.add(v);
     rep(b, 0, n) if (!iset[b] && frm[b] == -1 && m1.check(b))
       ans.push_back(b), frm[b] = a;
      return ans;
    auto backE = [&](int b) {
     m2.clear();
      rep(cas, 0, 2) rep(v, 0, n)
       if ((v == b \mid | iset[v]) && (frm[v] == -1) == cas) {
          if (!m2.check(v))
            return cas ? q.push(v), frm[v] = b, v : -1;
          m2.add(v);
      return n;
```

```
while (!q.empty()) {
 int a = q.front(), c; q.pop();
 for (int b : fwdE(a))
   while((c = backE(b)) >= 0) if (c == n) {
     while (b != n) iset[b] ^= 1, b = frm[b];
     return true;
return false;
```

7.3 Matching

hopcroftKarp.h

9812a7, 60 lines

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(q, btoa);

```
Time: \mathcal{O}\left(\sqrt{V}E\right)
                                                                      f612e4, 41 lines
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) {
   if (btoa[b] == -1 \mid | dfs(btoa[b], L + 1, g, btoa, A, B))
     return btoa[b] = a, 1;
 return 0;
int hopcroftKarp(vector<vi>& g, vi& btoa) {
 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 lav = 1;; lav++) {
     bool islast = 0;
     next.clear();
     for (int a : cur) for (int b : g[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: Maximum 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);
Time: \mathcal{O}(VE)
```

```
522b98, 20 lines
bool find(int j, vector<vi>& g, vi& btoa, vi& vis) {
 if (btoa[j] == -1) return 1;
 vis[j] = 1; int di = btoa[j];
 for (int e : q[di])
   if (!vis[e] && find(e, g, btoa, vis)) {
      btoa[e] = di; return 1;
 return 0;
int dfsMatching(vector<vi>& g, vi& btoa) {
 rep(i, 0, sz(q)) {
    vis.assign(sz(btoa), 0);
    for (int j : q[i])
      if (find(j, q, 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"
                                                     da4196, 20 lines
vi cover(vector<vi>& g, int n, int m) {
 vi match(m, -1);
 int res = dfsMatching(g, match);
 vector<bool> lfound(n, true), seen(m);
  for (int it : match) if (it != -1) lfound[it] = false;
 vi q, cover;
  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] = costfor 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,1,n) {
```

```
p[0] = i;
  int j0 = 0; // add "dummy" 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;</pre>
     if (dist[j] < delta) delta = dist[j], j1 = j;</pre>
    rep(j,0,m) {
     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
```

GeneralMatching.h

Description: Matching for general graphs. Fails with probability N/mod. Time: $\mathcal{O}\left(N^3\right)$

```
"../numerical/MatrixInverse-mod.h"
                                                     cb1912, 40 lines
vector<pii> generalMatching(int N, vector<pii>& ed) {
  vector<vector<ll>> mat(N, vector<ll>(N)), A;
  for (pii pa : ed) {
   int a = pa.first, b = pa.second, r = rand() % mod;
   mat[a][b] = r, mat[b][a] = (mod - r) % mod;
 int r = matInv(A = mat), M = 2*N - r, fi, f;
  assert(r % 2 == 0);
  if (M != N) do {
   mat.resize(M, vector<ll>(M));
   rep(i,0,N) {
     mat[i].resize(M);
     rep(j,N,M) {
       int r = rand() % mod;
       mat[i][j] = r, mat[j][i] = (mod - r) % mod;
  } while (matInv(A = mat) != M);
  vi has(M, 1); vector<pii> ret;
  rep(it,0,M/2) {
   rep(i,0,M) if (has[i])
     rep(j,i+1,M) if (A[i][j] && mat[i][j]) {
       fi = i; fj = j; goto done;
    } assert(0); done:
    if (fj < N) ret.emplace_back(fi, fj);</pre>
   has[fi] = has[fj] = 0;
    rep(sw,0,2) {
     11 a = modpow(A[fi][fi], mod-2);
     rep(i,0,M) if (has[i] && A[i][fj]) {
       ll b = A[i][fi] * a % mod;
        rep(j, 0, M) A[i][j] = (A[i][j] - A[fi][j] * b) % mod;
     swap(fi,fj);
```

```
return ret;
```

7.4 DFS algorithms

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 versa.

Usage: $sc(graph, [\&](vi\& v) \{ ... \})$ 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)
   low = min(low, val[e] ?: dfs(e,g,f));
 if (low == val[j]) {
     x = z.back(); z.pop_back();
     comp[x] = ncomps;
     cont.push back(x);
    } while (x != j);
    f(cont); cont.clear();
   ncomps++;
 return val[j] = low;
template < class G, class F> void scc(G& g, 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);
```

BCCs.h

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)
```

(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]);
      if (num[y] < me)
        st.push_back(e);
    } else {
      int si = sz(st);
      int up = dfs(y, e, f);
      top = min(top, up);
      if (up == me) {
        st.push_back(e);
```

```
f(vi(st.begin() + si, st.end()));
    st.resize(si);
}
else if (up < me) st.push_back(e);
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);
```

ArticulationPoints.h

Description: Finds the articulation points in a graph **Time:** $\mathcal{O}(N+M)$

e665c6, 31 lines

15

```
vector<vector<int>> adj;
vector<bool> visited;
vector<int> tin, low;
int timer;
void dfs(int v, int p = -1) {
    visited[v] = true;
    tin[v] = low[v] = timer++;
    int children = 0;
    for (int to : adj[v]) {
        if (to == p) continue;
        if (visited[to]) {
            low[v] = min(low[v], tin[to]);
        } else {
            dfs(to, v);
            low[v] = min(low[v], low[to]);
            if (low[to] >= tin[v] && p!=-1)
                // V IS A CUTPOINT:
            ++children;
    if (p == -1 && children > 1) // V IS A CUTPOINT;
void findAP() {
    timer = 0;
    visited.assign(n, false);
    tin.assign(n, -1);
    low.assign(n, -1);
    for (int i = 0; i < n; ++i)
        if (!visited[i]) dfs (i);
```

2sat.h

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

```
Usage: TwoSat ts(number of boolean variables); ts.either(0, \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, \sim1 and 2 are true ts.solve(); // Returns true iff it is solvable ts.values[0..N-1] holds the assigned values to the vars
```

Time: $\mathcal{O}\left(N+E\right)$, where N is the number of boolean variables, and E is the number of clauses. 5f9706, 50 lines

```
struct TwoSat {
  int N;
  vector<vi> gr;
```

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

EulerWalk.h

Description: Eulerian undirected/directed path/cycle algorithm. (visits every edge once) 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)
                                                      a07095, 15 lines
vi eulerWalk(vector<vector<pii>>>& gr, int nEdges, int src) {
  int n = sz(qr);
  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()};
```

DominatorTree.h

Description: Builds a dominator tree on a directed graph. Output tree is a parent array with src as the root.

```
Time: \mathcal{O}(V+E)
                                                     1d35d2, 46 lines
vi getDomTree(vvi &adj, int src) {
 int n = sz(adj), t = 0;
  vvi revAdj(n), child(n), sdomChild(n);
  vi label(n, -1), revLabel(n), sdom(n), idom(n), par(n), best(
  auto dfs = [&] (int cur, auto &dfs) -> void {
    label[cur] = t, revLabel[t] = cur;
    sdom[t] = par[t] = best[t] = t; t++;
    for(int nxt: adj[cur]) {
      if(label[nxt] == -1) {
        dfs(nxt, dfs);
        child[label[cur]].push_back(label[nxt]);
      revAdj[label[nxt]].push_back(label[cur]);
  };
  dfs(src, dfs);
  auto get = [&] (int x, auto &get) -> int {
    if(par[x] != x) {
      int t = get(par[x], get);
      par[x] = par[par[x]];
      if(sdom[t] < sdom[best[x]]) best[x] = t;</pre>
    return best[x];
  };
  for (int i = t-1; i >= 0; i--) {
    for(int j: revAdj[i]) sdom[i] = min(sdom[i], sdom[get(j,
    if(i > 0) sdomChild[sdom[i]].push_back(i);
    for(int j: sdomChild[i]) {
      int k = get(j, get);
      if(sdom[j] == sdom[k]) idom[j] = sdom[j];
      else idom[j] = k;
    for(int j: child[i]) par[j] = i;
  vi dom(n);
    if(idom[i] != sdom[i]) idom[i] = idom[idom[i]];
    dom[revLabel[i]] = revLabel[idom[i]];
```

7.5 Coloring

EdgeColoring.h

return dom:

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> adi(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 \in A eds, F f, B P = A \in A (), 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];
```

d2787e, 32 lines

```
sort(all(r), [](auto a, auto b) { return a.d > b.d; });
    int mxD = r[0].d;
   rep(i, 0, sz(r)) r[i].d = min(i, mxD) + 1;
  void expand(vv& R, int lev = 1) {
   S[lev] += S[lev - 1] - old[lev];
   old[lev] = S[lev - 1];
    while (sz(R)) {
     if (sz(q) + R.back().d <= sz(qmax)) return;
     q.push_back(R.back().i);
      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-Cover.

Trees

7.7.1 Number of Spanning Trees

Create an $N \times N$ matrix mat, and for each edge $a \to b \in G$, do mat[a][b]--, mat[b][b]++ (and mat[b][a]--, 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

$7.7.2 \frac{\text{row/column}}{\text{Erdos-Gallai theorem}}$

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 $k = 1 \dots n$,

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

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

```
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];</pre>
 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];
```

KthPath.h

Description: kth on path, goes in tree_lift

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

```
080dbc, 9 lines
int kth_path(int u, int v, int k) {
 int lca_d = t[lca(u, v)].d;
 int u lca = t[u].d - lca d;
 int v_lca = t[v].d - lca_d;
 if (k <= u_lca) return kth_par(u, k);</pre>
 if (k <= u lca + v lca)</pre>
   return kth_par(v, u_lca + v_lca - k);
 return -1:
```

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

```
"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;
```

CentroidDecomp.h

Description: Calls callback function on undirected forest for each centroid Usage: centroid(adj, [&](const vector<vector<int>>& adj, int cent) { ... }); Time: $\mathcal{O}(n \log n)$

```
template <class F, class G> struct centroid {
 G adj;
 F f;
 vi sub_sz, par;
  centroid(const G& adj, F f)
    : adj(adj), f(f), sub\_sz(sz(adj), -1), par(sz(adj), -1) {
    rep(i, 0, sz(adj))
      if (sub\_sz[i] == -1) dfs(i);
 void calc_sz(int u, int p) {
    sub_sz[u] = 1;
    for (int v : adj[u])
     if (v != p) calc_sz(v, u), sub_sz[u] += sub_sz[v];
 int dfs(int u) {
    calc_sz(u, -1);
    for (int p = -1, sz\_root = sub\_sz[u];;) {
     auto big_ch = find_if(all(adj[u]), [&](int v) {
        return v != p && 2 * sub_sz[v] > sz_root;
      if (big_ch == end(adj[u])) break;
      p = u, u = *big_ch;
    f(adj, u);
    for (int v : adj[u]) {
     iter_swap(find(all(adj[v]), u), rbegin(adj[v]));
      adj[v].pop_back();
      par[dfs(v)] = u;
    return u;
};
```

EdgeCD.h

Description: Recursively splits a tree into two edge sets that share a centroid. Consider all paths that pass through the centroid and use at least one edge from each set. A node can be a centroid multiple times. Consider all length 1 paths separately. Callback takes the graph, centroid, and split, where edges [0, split) from adj[centroid] are in the first set and the rest are in the second set.

Usage: edge_cd(adj, [&](const vector<vector<int>>& adj, int cent, int split) $\{ \ldots \}$); Time: $\mathcal{O}(n \log n)$

```
436f41, 34 lines
template <class F> struct edge_cd {
 vvi adj;
 F f;
 vi sub sz;
 edge_cd(const vvi& adj, F f) : adj(adj), f(f),
   sub_sz(sz(adj)) {
   dfs(0, sz(adj) - 1);
 int find_cent(int u, int p, int siz) {
   sub\_sz[u] = 1;
   for (int v : adj[u]) if (v != p) {
     int cent = find_cent(v, u, siz);
     if (cent != -1) return cent;
     sub_sz[u] += sub_sz[v];
```

HLD LinkCutTree DirectedMST

```
return 2 * sub sz[u] > siz ?
     p >= 0 \&\& (sub sz[p] = siz + 1 - sub sz[u]), u : -1;
  void dfs(int u, int siz) {
   if (siz < 2) return;
   u = find_cent(u, -1, siz);
   int sum = 0;
    auto it = partition(all(adj[u]), [&](int v) {
     11 x = sum + sub_sz[v];
     return x * x < siz * (siz - x)? sum += sub_sz[v], 1 : 0;
    f(adj, u, it - begin(adj[u]));
   vi oth(it, end(adj[u]));
   adj[u].erase(it, end(adj[u]));
   dfs(u, sum);
   swap(adj[u], oth);
   dfs(u, siz - sum);
};
```

HLD.h

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.

Usage: Maximum Values on path from any two arbitrary nodes For node to root, use Euler Tour

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 1, int r) { tree->add(1, 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]);
};
```

LinkCutTree.h

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

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

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

```
node[u].pp = &node[v];
void cut(int u, int v) { // remove an edge (u, v)
  Node *x = &node[u], *top = &node[v];
  makeRoot(top); x->splay();
  assert(top == (x->pp ?: x->c[0]));
  if (x->pp) x->pp = 0;
  else {
    x->c[0] = top->p = 0;
    x->fix():
bool connected(int u, int v) { // are u, v in the same tree?
  Node* nu = access(&node[u]) -> first();
  return nu == access(&node[v])->first();
void makeRoot(Node* u) {
  access(u);
  u->splay();
  if(u->c[0]) {
    u -> c[0] -> p = 0;
    u - c[0] - flip ^= 1;
    u - c[0] - pp = u;
    u - > c[0] = 0;
    u->fix();
Node* access(Node* u) {
  u->splay();
  while (Node* pp = u->pp) {
    pp->splay(); u->pp = 0;
    if (pp->c[1]) {
      pp->c[1]->p = 0; pp->c[1]->pp = pp; }
    pp - c[1] = u; pp - fix(); u = pp;
  return u;
```

DirectedMST.h

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

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

```
"../data-structures/UnionFindRollback.h"
                                                      39e620, 60 lines
struct Edge { int a, b; ll w; };
struct Node {
  Edge kev;
  Node *1, *r;
  11 delta:
  void prop() {
    key.w += delta;
   if (1) 1->delta += delta;
   if (r) r->delta += delta;
    delta = 0:
  Edge top() { prop(); return key; }
Node *merge(Node *a, Node *b) {
 if (!a || !b) return a ?: b;
  a->prop(), b->prop();
  if (a->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<11, vi> dmst(int n, int r, vector<Edge>& g) {
  RollbackUF uf(n);
  vector<Node*> heap(n);
```

```
for (Edge e : g) 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) {</pre>
   if (!heap[u]) return {-1,{}};
   Edge e = heap[u]->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 * cyc = 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};
```

Geometry (8)

8.1 Geometric primitives

Point.h

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

```
0c250f, 25 lines
template <class T> int sqn(T x) \{ return (x > 0) - (x < 0); \}
template<class T>
struct P {
 T x, y;
  explicit P(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; }
  ld dist() const { return sqrt((ld)dist2()); }
  // angle to x-axis in interval [-pi, pi]
  ld 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(ld a) const { return P(x*cos(a)-y*sin(a),x*sin(a)+y*
      cos(a)); }
```

lineDistance.h

${\bf Description:}$

Returns the signed distance between point p and the line containing points a and b. Positive value on left side and negative on right as seen from a towards b. a==b gives nan. P is supposed to be Point<T> or Point3D<T> where T is e.g. double or long long. It uses products in intermediate steps so watch out for overflow if using int or long long. Using Point3D will always give a non-negative distance. For Point3D, call dist on the result of the cross product.



"Point.h" 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();
}

LineProjectionReflection.h

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

```
"Point.h" b5562d, 5 lines
template<class P>
P lineProj(P a, P b, P p, bool refl=false) {
  P v = b - a;
  return p - v.perp()*(1+refl)*v.cross(p-a)/v.dist2();
}
```

inAngle.h

Description: Returns true if p is in the angle between b and c. The angle starts at point a (b and c are endpoints). The angle is assumed to be at most 180 and the points are not colinear. Modify equals sign for exclusive inAngle check.

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.

```
struct Angle {
  int x, y;
  int t;
  Angle(int x, int y, int t=0) : x(x), y(y), t(t) {}
  Angle operator-(Angle b) const { return {x-b.x, y-b.y, t}; }
  int half() const {
    assert(x || y);
    return y < 0 || (y == 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;
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)\};
```

polarSort.h

Description: Sorts vectors radially based on angles [0, 2pi). For parallel vectors, larger magnitude come later.

```
The point of the polar sort of the polar const point of the polar const polar const
```

lineSort.h

Description: Sorts points on a line s->e (or from their projections / parallel intersections) s and e represent the line we are comparing to

8.2 Lines and Segments

sideOf.h

Description: Returns where p is as seen from s towards e. $1/0/-1 \Leftrightarrow left/on line/right$. 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.

```
auto a = (e-s).cross(p-s);
double l = (e-s).dist()*eps;
return (a > 1) - (a < -1);</pre>
```

OnSegment.h

Description: Returns true iff p lies on the line segment from s to e. Use $(segDist(s,e,p) \le psilon)$ instead when using Point double.

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



Usage: auto res = lineInter(s1,e1,s2,e2);
if (res.first == 1)

```
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};
}
```

linearTransformation.h Description:

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



SegmentDistance.h

Description:

Returns the shortest distance between point p and the line segment from point s to e. For SEGMENT-SEGMENT DISTANCE, check for intersection. If there is no intersection, the closest distance is between one of the four points and the other segment.



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<11> and the intersection point does not have integer coordinates. Products of three coordinates are used in intermediate steps so watch out for overflow if using int or long long.



```
Usage: vector<P> inter = segInter(s1,e1,s2,e2);
if (sz(inter) == 1)
cout << "segments intersect at " << inter[0] << endl;</pre>
"Point.h", "OnSegment.h"
template<class P> vector<P> segInter(P a, P b, P c, P d) {
 auto oa = c.cross(d, a), ob = c.cross(d, b),
       oc = a.cross(b, c), od = a.cross(b, d);
  // Checks if intersection is single non-endpoint point.
  if (sqn(oa) * sqn(ob) < 0 \&\& sqn(oc) * sqn(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)};
```

8.3 Circles

Circle-Circle-Intersection.h

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

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.

```
return out;
}
```

CircleLine.h

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

CirclePolygonIntersection.h

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

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

```
"../../content/geometry/Point.h"
typedef Point < double > P;
#define arg(p, g) atan2(p.cross(g), p.dot(g))
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, g) * r2;
    auto s = max(0., -a-sqrt(det)), t = min(1., -a+sqrt(det));
    if (t < 0 || 1 <= s) return arg(p, q) * r2;
    P u = p + d * s, v = q + d * (t-1);
    return arg(p,u) * r2 + u.cross(v)/2 + arg(v,q) * r2;
  auto sum = 0.0;
  rep(i, 0, sz(ps))
   sum += tri(ps[i] - c, ps[(i + 1) % sz(ps)] - c);
  return sum;
```

circumcircle.h

Description:

"Point.h"

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



```
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};
```

8.4 Polygons

InsidePolygon.h

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

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

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

```
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;
```

Polygon Area, 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 a = v.back().cross(v[0]);
 rep(i, 0, sz(v)-1) a += v[i].cross(v[i+1]);
 return a:
```

triangleArea.h

Description: Finds the area of a triangle

f20dcb, 3 lines

```
ld triangleArea(P a, P b, P c) {
    return abs((b-a).cross(c-a)) / (ld)(2);
```

PolygonCenter.h

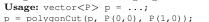
Description: Returns the center of mass for a polygon.

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

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





"Point.h" d07181, 13 lines typedef Point < double > P; vector<P> polygonCut(const vector<P>& poly, P s, P e) { vector<P> res; rep(i,0,sz(polv)) { P cur = poly[i], prev = i ? poly[i-1] : poly.back(); auto a = s.cross(e, cur), b = s.cross(e, prev); if ((a < 0) != (b < 0))res.push_back(cur + (prev - cur) * (a / (a - b))); if (a < 0)res.push back(cur); return res;

PolygonUnion.h

return ret / 2;

Description: Calculates the area of the union of n polygons (not necessarily convex). The points within each polygon must be given in CCW order. (Epsilon checks may optionally be added to sideOf/sgn, but shouldn't be

Time: $\mathcal{O}(N^2)$, where N is the total number of points

```
"Point.h", "sideOf.h"
                                                     3931c6, 33 lines
typedef Point < double > P;
double rat(P a, P b) { return sqn(b.x) ? a.x/b.x : a.y/b.y; }
double polyUnion(vector<vector<P>>& poly) {
  double ret = 0;
  rep(i, 0, sz(poly)) rep(v, 0, sz(poly[i])) {
    P A = poly[i][v], B = poly[i][(v + 1) % sz(poly[i])];
    vector<pair<double, int>> segs = {{0, 0}, {1, 0}};
    rep(j,0,sz(poly)) if (i != j) {
      rep(u, 0, sz(poly[j])) {
       P C = poly[j][u], D = poly[j][(u + 1) % sz(poly[j])];
        int sc = sideOf(A, B, C), sd = sideOf(A, B, D);
        if (sc != sd) {
          double sa = C.cross(D, A), sb = C.cross(D, B);
          if (\min(sc, sd) < 0)
            segs.emplace_back(sa / (sa - sb), sgn(sc - sd));
        } else if (!sc && !sd && j<i && sqn((B-A).dot(D-C))>0){
          segs.emplace_back(rat(C - A, B - A), 1);
          segs.emplace_back(rat(D - A, B - A), -1);
    sort (all (segs));
    for (auto\& s : segs) s.first = min(max(s.first, 0.0), 1.0);
    double sum = 0:
    int cnt = segs[0].second;
    rep(j,1,sz(segs)) {
     if (!cnt) sum += segs[j].first - segs[j - 1].first;
     cnt += segs[j].second;
    ret += A.cross(B) * sum;
```

HalfplaneIntersection.h

Description: Returns the intersection of halfplanes as a polygon Time: $\mathcal{O}(n \log n)$

```
const double eps = 1e-8;
typedef Point < double > P;
struct HalfPlane {
 P s, e, d;
  HalfPlane(P s = P(), P e = P()): s(s), e(e), d(e - s) {}
 bool contains(P p) { return d.cross(p - s) > -eps; }
  bool side() { return d.x < -eps | | (abs(d.x) < =eps && d.y > 0); }
  bool operator<(HalfPlane hp) {</pre>
    if(side() != hp.side()) return side();
    return d.cross(hp.d) > 0;
 P inter(HalfPlane hp) {
    auto p = hp.s.cross(e, hp.e), q = hp.s.cross(hp.e, s);
    return (s * p + e * q) / d.cross(hp.d);
};
vector<P> hpIntersection(vector<HalfPlane> hps) {
 sort(all(hps));
 int n = sz(hps), 1 = 1, r = 0;
 vector<HalfPlane> dq(n+1);
 rep(i, 0, n) {
    while(l<r && !hps[i].contains(dq[r].inter(dq[r-1]))) r--;</pre>
    while(l<r && !hps[i].contains(dg[l].inter(dg[l+1]))) l++;</pre>
    da[++r] = hps[i];
    if(1 < r \&\& abs(dq[r].d.cross(dq[r-1].d)) < eps) {
      if(dq[r].d.dot(dq[r-1].d) < 0) return {};
      if(dq[--r].contains(hps[i].s)) dq[r] = hps[i];
  while (1 < r-1 \& \& !dq[1].contains(dq[r].inter(dq[r-1]))) r--;
  while (1 < r-1 \& \& !dq[r].contains(dq[l].inter(dq[l+1]))) l++;
  if(1 > r-2) return {};
  vector<P> poly;
  rep(i, 1, r) poly.push_back(dq[i].inter(dq[i+1]));
 poly.push_back(dq[r].inter(dq[l]));
 return poly;
```

ConvexHull.h

Description:

"Point.h"

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



e89ff2, 13 lines

21

typedef Point<11> P; vector<P> convexHull(vector<P> pts) { if (sz(pts) <= 1) return pts;</pre> sort(all(pts)); vector<P> h(2*sz(pts)); // modified for cses hack

```
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--;
    h[t++] = p;
return \{h.begin(), h.begin() + t - (t == 2 && h[0] == h[1])\};
```

HullDiameter.h

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

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

```
"Point.h" c571b8, 12 lines

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

PointInsideHull.h

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

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

```
"Point.h", "sideOf.h", "OnSegment.h"

typedef Point<11> P;

bool inHull(const vector<P>& h, P p, bool strict = true) {
  int a = 1, b = sz(h) - 1, r = !strict;
  if (sz(h) < 3) return r && onSegment(h[0], h.back(), p);
  if (sideOf(h[0], h[a], h[b]) > 0) swap(a, b);
  if (sideOf(h[0], h[a], p) >= r || sideOf(h[0], h[b], p) <= -r)
    return false;

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

ExtremeVertex.h

Description: returns the point of a hull with the max projection onto a line.

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

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) sgn(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 ls = cmp(lo + 1, lo), ms = cmp(m + 1, m);
    (1s < ms \mid | (1s == ms \&\& 1s == cmp(1o, m)) ? hi : 1o) = 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 | cmpL(endB) > 0)
   return {-1, -1};
 array<int, 2> res;
 rep(i, 0, 2) {
   int lo = endB, hi = endA, n = sz(poly);
   while ((lo + 1) % n != hi) {
     int m = ((lo + hi + (lo < hi ? 0 : n)) / 2) % n;
      (cmpL(m) == cmpL(endB) ? lo : hi) = m;
    res[i] = (lo + !cmpL(hi)) % n;
    swap (endA, endB);
 if (res[0] == res[1]) return {res[0], -1};
 if (!cmpL(res[0]) && !cmpL(res[1]))
   switch ((res[0] - res[1] + sz(poly) + 1) % sz(poly)) {
     case 0: return {res[0], res[0]};
     case 2: return {res[1], res[1]};
 return res;
```

HullTangents.h

Description: Finds the left and right, respectively, tangent points on convex hull from a point. If the point is colinear to side(s) of the polygon, the point further away is returned. Requires ccw, $n \geq 3$, and the point be on or outside the polygon. Can be used to check if a point is inside of a convex hull. Will return -1 if it is strictly inside. If the point is on the hull, the two adjacent points will be returned

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

MinkowskiSum.h

Description: Returns the minkowski sum of a set of convex polygons **Time:** $\mathcal{O}(n \log n)$

```
#define side(p) (p.x > 0 | | (p.x == 0 && p.y > 0))
template<class P>
vector<P> convolve(vector<vector<P>> &polvs){
 P init; vector<P> dir;
 for(auto poly: polys) {
   int n = sz(poly);
   if(n) init = init + poly[0];
   if(n < 2) continue;
    rep(i, 0, n) dir.push_back(poly[(i+1)%n] - poly[i]);
 if(size(dir) == 0) return { init };
 stable_sort(all(dir), [&](P a, P b)->bool {
   if(side(a) != side(b)) return side(a);
   return a.cross(b) > 0;
 vector<P> sum; P cur = init;
 rep(i, 0, sz(dir))
   sum.push_back(cur), cur = cur + dir[i];
 return sum;
```

lineTranslations.h

Description: Translates a line using a given translation vector u or given a distance dist

```
void vectorTransaltion (P &s, P &e, P u) {
    s += u; e+=u;
}

void shiftLeft (P &s, P &e, ld dist) {
    P v = e-s, n = P(-v.y, v.x);
    n = n * (dist / abs(n));
    s += u; e+=u;
}
```

8.5 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; });</pre> pair<ll, pair<P, P>> ret{LLONG_MAX, {P(), P()}}; int j = 0;for (P p : v) { P d{1 + (11) 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;

ManhattanMST.h

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

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

```
"Point.h"
                                                     df6f59, 23 lines
typedef Point<int> P;
vector<array<int, 3>> manhattanMST(vector<P> ps) {
 vi id(sz(ps));
 iota(all(id), 0);
  vector<array<int, 3>> edges;
  rep(k,0,4) {
    sort(all(id), [&](int i, int j) {
         return (ps[i]-ps[j]).x < (ps[j]-ps[i]).y;});
   map<int, int> sweep;
    for (int i : id) {
      for (auto it = sweep.lower_bound(-ps[i].y);
                it != sweep.end(); sweep.erase(it++)) {
       int j = it->second;
       P d = ps[i] - ps[j];
       if (d.y > d.x) break;
        edges.push_back({d.y + d.x, i, j});
     sweep[-ps[i].y] = i;
    for (P\& p : ps) if (k \& 1) p.x = -p.x; else swap(p.x, p.y);
  return edges;
```

kdTree.h

Description: KD-tree (2d, can be extended to 3d)

```
bac5b0, 63 lines
typedef long long T;
typedef Point<T> P;
const T INF = numeric limits<T>::max();
bool on x(const P& a, const P& b) { return a.x < b.x; }
bool on_y(const P& a, const P& b) { return a.y < b.y; }
struct Node {
 P pt; // if this is a leaf, the single point in it
 T x0 = INF, x1 = -INF, y0 = INF, y1 = -INF; // bounds
 Node *first = 0, *second = 0;
 T distance (const P& p) { // min squared distance to a point
   T x = (p.x < x0 ? x0 : p.x > x1 ? x1 : p.x);
   T y = (p.y < y0 ? y0 : p.y > y1 ? y1 : p.y);
    return (P(x,y) - p).dist2();
 Node(vector<P>&& vp) : pt(vp[0]) {
    for (P p : vp) {
     x0 = min(x0, p.x); x1 = max(x1, p.x);
     y0 = min(y0, p.y); y1 = max(y1, p.y);
    if (vp.size() > 1) {
     // split on x if width >= height (not ideal...)
     sort(all(vp), x1 - x0 >= y1 - y0 ? on_x : on_y);
      // divide by taking half the array for each child (not
      // best performance with many duplicates in the middle)
     int half = sz(vp)/2;
     first = new Node({vp.begin(), vp.begin() + half});
      second = new Node({vp.begin() + half, vp.end()});
struct KDTree {
 Node* root;
  KDTree(const vector<P>& vp) : root(new Node({all(vp)})) {}
```

```
pair<T, P> search(Node *node, const P& p) {
   if (!node->first) {
      // uncomment if we should not find the point itself:
      // if (p = node \rightarrow pt) return {INF, P()};
     return make_pair((p - node->pt).dist2(), node->pt);
   Node *f = node->first, *s = node->second;
   T bfirst = f->distance(p), bsec = s->distance(p);
   if (bfirst > bsec) swap(bsec, bfirst), swap(f, s);
    // search closest side first, other side if needed
    auto best = search(f, p);
   if (bsec < best.first)</pre>
     best = min(best, search(s, p));
    return best;
 // find nearest point to a point, and its squared distance
 // (requires an arbitrary operator< for Point)
 pair<T, P> nearest(const P& p) {
   return search (root, p);
};
```

DelaunayTriangulation.h

Description: Computes the Delaunay triangulation of a set of points. Each circumcircle contains none of the input points. If any three points are collinear or any four are on the same circle, behavior is undefined. **Time:** $\mathcal{O}\left(n^2\right)$

PlanarFaceExtraction.h

Description: Given a planar graph and where the points are, extract the set of faces that the graph makes. The inner faces will be returned in counter-clockwise order, and the outermost face will be returned in clockwise order. Time: $\mathcal{O}\left(ElogE\right)$

```
63f230, 39 lines
template<class P>
vector<vector<P>> extract_faces(vvi adj, vector<P> pts) {
 int n = sz(pts);
 #define cmp(i) [&](int pi, int qi) -> bool { \
   P p = pts[pi] - pts[i], q = pts[qi] - pts[i]; \
   bool sideP = p.y < 0 || (p.y == 0 && p.x < 0); \
   bool sideQ = q.y < 0 \mid \mid (q.y == 0 \&\& q.x < 0); 
   if(sideP != sideQ) return sideP; \
   return p.cross(q) > 0; }
 rep(i, 0, n)
    sort(all(adj[i]), cmp(i));
 rep(i, 0, n) for(int j: adj[i])
   ed.emplace_back(i, j);
 sort (all (ed));
 auto get_idx = [&](int i, int j) -> int {
   return lower_bound(all(ed), pii(i, j))-begin(ed);
 vector<vector<P>> faces;
 vi used(sz(ed));
 rep(i, 0, n) for(int j: adj[i]) {
```

```
if(used[get_idx(i, j)])
   continue;
used[get_idx(i, j)] = true;
vector<P> face = {pts[i]};
int prv = i, cur = j;
while(cur != i) {
   face.push_back(pts[cur]);
   auto it = lower_bound(all(adj[cur]), prv, cmp(cur));
   if(it == begin(adj[cur]))
      it = end(adj[cur]);
   prv = cur, cur = *prev(it);
   used[get_idx(prv, cur)] = true;
}
faces.push_back(face);
}
#undef cmp
return faces;
```

8.6 3D

PolyhedronVolume.h

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

```
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. ${}_{8058ae,\ 32\ lines}$

```
template<class T> struct Point3D {
 typedef Point3D P:
 typedef const P& R;
 T x, y, z;
 explicit Point3D(T x=0, T y=0, T z=0) : x(x), y(y), z(z) {}
 bool operator<(R p) const {
   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 $dx = \sin(t2) \cdot \cos(f2) - \sin(t1) \cdot \cos(f1)$;

```
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)

9.1 Applications of String Functions Searching for a Sub-string in a String (KMP)

- Concatenate the sub-string to the beginning of the string plus another character that does not occur in either string.
- Run the prefix function on the string and count the number of occurrences of the length of the original sub-string.

Counting the Number of Prefixes in a String

- · Run the prefix function on the string.
- For each string of length i, do: ans[pi[i-1]] += ans[i] since each occurrence of a longer prefix implies the occurrence of the shorter prefix.
- Add one for each original (non-counted) prefix.

```
vll ans(n+1, 1); // Set to one for the actual prefix for (int i = 0; i; i; i++) ans[pi[i]]++; for (int i = n-1; i; i; i; i; i ans[pi[i-1]] += ans[i];
```

The Number of Different Substrings in a String

- · Start with an empty string.
- Add the next character of the original string to our string and reverse the string.
- Run the prefix function to find the max length of the substrings we have already found.
- The number of new substrings = accumulate(|S| + 1 maxPi).

Compressing a String

- Given a string s of length n, we want to find the shortest "compressed" representation of the string.
- If after running the prefix function, n is divisible by (n pi[n-1]), then the string is compressible, with minimum size (n - pi[n-1]).
- Otherwise, it is not compressible.

Creating Automation for String Matching

- After running the prefix function on one or many strings, an automation should be able to preprocess the strings in O(n) and then do matching in O(m) per query.
- This is especially useful for **many repeated searches** with the same pattern (KMP runs in O(n+m) per query).
- Build a table storing aut[state][charNumber], where state is the index of the string we are searching for and char is the current character.
- Depending on the character we see, move forwards or backwards to a new state (e.g., seeing an a moves us from state 1 → 2, but seeing a b moves us from state 2 → 0).

MatchingAutomation.h

```
Description: Creates substring searching automation
```

```
void computeAutomaton(string s, vector<vector<int>>& aut)
{
    s += '#';
    int n = s.size();
    vector<int> pi = prefix_function(s);
    aut.assign(n, vector<int>(26));

for (int i = 0; i < n; i++)
    for (int c = 0; c < 26; c++)</pre>
```

KMP.h

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

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.

```
vi pi(const string& s) {
vi pi(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) Used for Finding Periods CSES **Time:** $\mathcal{O}(n)$

```
vi Z(const string& S) {
  vi z(sz(S));
  int 1 = -1, r = -1;
  rep(i,1,sz(S)) {
    z[i] = i >= r ? 0 : min(r - i, z[i - 1]);
    while (i + z[i] < sz(S) && S[i + z[i]] == S[z[i]])
    z[i]++;
  if (i + z[i] > r)
    1 = i, r = i + z[i];
  }
  return z;
}
```

Manacher.h

c3fe8a, 12 lines

Description: For each position in a string, computes p[0][i] = half length of longest even palindrome around pos i, <math>p[1][i] = longest odd (half rounded down).

```
Time: \mathcal{O}(N) e7ad79, 13 lines
```

```
array<vi, 2> manacher(const string& s) {
  int n = sz(s);
  array<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][1+t]);
    int L = i-p[z][i], R = i+p[z][i]-!z;
  while (L>=1 && R+1<n && s[L-1] == s[R+1])
    p[z][i]++, L--, R++;
    if (R>r) l=L, r=R;
  }
  return p;
}
```

Eertree.h

Description: Generates an eartree on str. cur is accurate at the end of the main loop before the final assignment to t.

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

```
struct Eertree {
  vi slink = \{0, 0\}, len = \{-1, 0\};
  vvi down:
  int cur = 0, t = 0;
  Eertree(string &str) : down(2, vi(26, -1)) {
    for (int i = 0; i < sz(str); i++) {
     char c = str[i]; int ci = c - 'a';
     while (t \le 0 \mid | str[t-1] != c)
       t = i - len[cur = slink[cur]];
      if (down[cur][ci] == -1) {
        down[cur][ci] = sz(slink);
        down.emplace back(26, -1);
        len.push_back(len[cur] + 2);
        if (len.back() > 1) {
          do t = i - len[cur = slink[cur]];
          while (t \le 0 \mid | str[t-1] != c);
          slink.push_back(down[cur][ci]);
       } else slink.push back(1);
        cur = sz(slink) - 1;
     } else cur = down[cur][ci];
      t = i - len[cur] + 1;
};
```

MinRotation.h

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

Description: Finds the lexicographically smallest rotation of a string. **Usage:** rotate(v.begin(), v.begin()+minRotation(v), v.end());

int minRotation(string s) {
 int a=0, N=sz(s); s += s;
 rep(b,0,N) rep(k,0,N) {
 if (a+k == b || s[a+k] < s[b+k]) {b += max(0, k-1); break;}
 if (s[a+k] > s[b+k]) { a = b; break; }
 }
 return a;

SuffixArrav.h

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

```
635552, 22 lines
struct SuffixArray {
  vi sa, lcp;
  SuffixArray(string s, int lim=256) { // or vector<int>
    s.push\_back(0); int n = sz(s), k = 0, a, b;
   vi x(all(s)), y(n), ws(max(n, lim));
    sa = lcp = y, iota(all(sa), 0);
    for (int j = 0, p = 0; p < n; j = max(1, j * 2), lim = p) {
     p = j, iota(all(y), n - j);
     rep(i,0,n) if (sa[i] >= j) y[p++] = sa[i] - j;
      fill(all(ws), 0);
     rep(i, 0, n) ws[x[i]] ++;
      rep(i, 1, lim) ws[i] += ws[i - 1];
      for (int i = n; i--;) sa[--ws[x[y[i]]]] = y[i];
     swap(x, y), p = 1, x[sa[0]] = 0;
     rep(i,1,n) = sa[i-1], b = sa[i], x[b] =
        (y[a] == y[b] \&\& y[a + j] == y[b + j]) ? p - 1 : p++;
    for (int i = 0, j; i < n - 1; lcp[x[i++]] = k)
```

```
for (k \&\& k--, j = sa[x[i] - 1];

s[i + k] == s[j + k]; k++);

}
```

SuffixAutomaton.h

Description: Creates a partial DFA (DAG) that accepts all suffixes, with suffix links. One-to-one map between a path from the root and a substring. len is the longest-length substring ending here. pos is the first index in the string matching here. term is whether this node is a terminal (aka a suffix) **Time:** construction takes $\mathcal{O}(N \log K)$, where $K = \text{Alphabet Sigs}_{483 \text{afe}}$, $_{27 \text{ lines}}$

```
struct st {int len, pos, term, link=-1; map<char, int> next;};
struct SuffixAutomaton {
 vector<st> a;
 SuffixAutomaton(string &str) {
   a.resize(1);
   int last = 0;
    for(auto c : str) {
     int p = last, cur = last = sz(a);
     a.push_back({a[p].len + 1, a[p].len});
     while (p \ge 0 \&\& !a[p].next.count(c))
       a[p].next[c] = cur, p = a[p].link;
      if (p == -1) a [cur]. link = 0;
     else {
       int g = a[p].next[c];
       if (a[p].len + 1 == a[q].len) a[cur].link = q;
         a.push\_back({a[p].len+1, a[q].pos, 0, a[q].link,}
           a[q].next});
         for(; p \ge 0 \&\& a[p].next[c] == q; p = a[p].link)
           a[p].next[c] = sz(a)-1;
         a[q].link = a[cur].link = sz(a)-1;
     }
   while(last >= 0) a[last].term = 1, last = a[last].link;
};
```

SuffixTree.h

d07a42, 8 lines

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

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

```
struct SuffixTree {
 enum { N = 200010, ALPHA = 26 }; //N \sim 2*maxlen+10
 int toi(char c) { return c - 'a'; }
 string a; //v = cur \ node, q = cur \ position
 int t[N][ALPHA],1[N],r[N],p[N],s[N],v=0,q=0,m=2;
 void ukkadd(int i, int c) { suff:
   if (r[v]<=q) {
     if (t[v][c]==-1) { t[v][c]=m; l[m]=i;
       p[m++]=v; v=s[v]; q=r[v]; goto suff; }
     v=t[v][c]; q=l[v];
   if (q==-1 || c==toi(a[q])) q++; else {
     l[m+1]=i; p[m+1]=m; l[m]=l[v]; r[m]=q;
     p[m]=p[v]; t[m][c]=m+1; t[m][toi(a[q])]=v;
     l[v]=q; p[v]=m; t[p[m]][toi(a[l[m]])]=m;
     v=s[p[m]]; q=l[m];
     while (q < r[m]) \{ v = t[v][toi(a[q])]; q + = r[v] - l[v]; \}
     if (q==r[m]) s[m]=v; else s[m]=m+2;
     q=r[v]-(q-r[m]); m+=2; goto suff;
```

```
SuffixTree(string a) : a(a) {
    fill(r,r+N,sz(a));
    memset(s, 0, sizeof s);
    memset(t, -1, sizeof t);
    fill(t[1],t[1]+ALPHA,0);
    s[0] = 1; 1[0] = 1[1] = -1; r[0] = r[1] = p[0] = p[1] = 0;
    rep(i,0,sz(a)) ukkadd(i, toi(a[i]));
  // example: find longest common substring (uses ALPHA = 28)
  pii best;
  int lcs(int node, int i1, int i2, int olen) {
    if (1[node] <= i1 && i1 < r[node]) return 1;
    if (1[node] <= i2 && i2 < r[node]) return 2;</pre>
    int mask = 0, len = node ? olen + (r[node] - 1[node]) : 0;
    rep(c, 0, ALPHA) if (t[node][c] != -1)
     mask |= lcs(t[node][c], i1, i2, len);
    if (mask == 3)
      best = max(best, {len, r[node] - len});
    return mask;
  static pii LCS(string s, string t) {
    SuffixTree st(s + (char) ('z' + 1) + t + (char) ('z' + 2));
    st.lcs(0, sz(s), sz(s) + 1 + sz(t), 0);
    return st.best;
};
```

Hashing.h

Description: Self-explanatory methods for string hashing.

```
// Arithmetic mod 2^64-1. 2x slower than mod 2^64 and more
// code, but works on evil test data (e.g. Thue-Morse, where
// ABBA... and BAAB... of length 2^10 hash the same mod 2^64).
// "typedef ull H;" instead if you think test data is random,
// or work mod 10^9+7 if the Birthday paradox is not a problem.
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 	ext{ operator} * (H 	ext{ o}) { auto } m = (\underline{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;}
```

HashInterval.h

Description: Various self-explanatory methods for string hashing.

```
"Hashing.h"
                                                      122649, 12 lines
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];
};
```

AhoCorasick-Kactl.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. $\operatorname{find}(x)$ 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.empty());
   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 = v;
      else {
        N[ed].back = y;
        (N[ed].end == -1 ? N[ed].end : backp[N[ed].start])
         = N[v].end;
        N[ed].nmatches += N[y].nmatches;
        q.push(ed);
vi find(string word) {
  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;
```

LyndonFactorization.h

Description: Computes the Lyndon Factorization of a string. A Lyndon word is a nonempty string that is strictly smaller in lexicographic order than any of its proper suffixes. Returns the starting indices of the Lyndon words in the string. Time: $\mathcal{O}\left(n\right)$

```
09e827, 12 lines
vi duval(string &s) {
 vi ans:
 for(int start = 0; start < sz(s);) {</pre>
   int i = start+1, j = start;
   for(; i < sz(s) && s[i] >= s[j]; i++)
     if(s[i] > s[j]) j = start;
     else j++;
   for (int sz = i-j; start + sz <= i; start += sz)
     ans.push_back(start);
 return ans;
```

SuffixAutomaton.h

Description: Creates a partial DFA (DAG) that accepts all suffixes, with suffix links. One-to-one map between a path from the root and a substring. len is the longest-length substring ending here. pos is the first index in the string matching here. term is whether this node is a terminal (aka a suffix) Time: construction takes $\mathcal{O}(N \log K)$, where $K = \text{Alphabet Size}_{383 \text{afe}, 27 \text{ lines}}$

```
struct st {int len, pos, term, link=-1; map<char, int> next;};
struct SuffixAutomaton {
 vector<st> a;
 SuffixAutomaton(string &str) {
   a.resize(1);
```

```
int last = 0;
    for(auto c : str) {
      int p = last, cur = last = sz(a);
      a.push_back({a[p].len + 1, a[p].len});
      while(p >= 0 && !a[p].next.count(c))
        a[p].next[c] = cur, p = a[p].link;
      if (p == -1) a [cur]. link = 0;
      else {
        int q = a[p].next[c];
        if (a[p].len + 1 == a[q].len) a[cur].link = q;
          a.push\_back({a[p].len+1, a[q].pos, 0, a[q].link,}
            a[q].next});
          for(; p \ge 0 \&\& a[p].next[c] == q; p = a[p].link)
            a[p].next[c] = sz(a)-1;
          a[q].link = a[cur].link = sz(a)-1;
    while(last >= 0) a[last].term = 1, last = a[last].link;
};
```

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) {
   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]; });
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.

```
Usage: constantIntervals(0, sz(v), [&](int x){return v[x];}, [&](int lo, int hi, T val){...});
```

```
Time: O\left(k\log\frac{n}{k}\right)
                                                       753a4c, 19 lines
template<class F, class G, class T>
void rec(int from, int to, F& f, G& g, int& i, T& p, T q) {
  if (p == q) return;
  if (from == to) {
    q(i, to, p);
    i = to; p = q;
    int mid = (from + to) >> 1;
    rec(from, mid, f, q, i, p, f(mid));
    rec(mid+1, to, f, q, i, p, q);
template < class F, class G>
void constantIntervals(int from, int to, F f, G g) {
 if (to <= from) return;
 int i = from; auto p = f(i), q = f(to-1);
 rec(from, to-1, f, q, 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)) 9155b4, 11 lines
```

```
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>
```

LIS.h

Description: Compute indices for the longest increasing subsequence. **Time:** $\mathcal{O}(N \log N)$

```
template<class I> vi lis(const vector<I>& S) {
   if (S.empty()) return {};
   vi prev(sz(S));
   typedef pair<I, int> p;
   vector res;
   rep(i,0,sz(S)) {
      // change 0 -> i for longest non-decreasing subsequence
      auto it = lower_bound(all(res), p{S[i], 0});
      if (it == res.end()) res.emplace_back(), it = res.end()-1;
      *it = {S[i], i};
      prev[i] = it == res.begin() ? 0 : (it-1)->second;
   }
   int L = sz(res), cur = res.back().second;
   vi ans(L);
   while (L--) ans[L] = cur, cur = prev[cur];
   return ans;
```

10.3 Dynamic programming

KnuthDP.h

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

```
 \begin{array}{ll} \textbf{Time: } \mathcal{O}\left((N+(hi-lo))\log N\right) & & & \\ \textbf{struct DP } \{\ //\ Modify\ at\ will: \\ & \textbf{int lo(int ind)}\ \{\ \text{return 0; }\} \\ & \textbf{int hi(int ind)}\ \{\ \text{return ind; }\} \\ & 11\ f(\textbf{int ind, int k})\ \{\ \text{return dp[ind][k]; }\} \end{array}
```

```
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<1l, 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); }
};
```