

University of Central Florida

UCF Apocalypse Attack

Sachin Sivakumar, Andy Phan, Tyler Marks

```
1 Contest
                                                                 troubleshoot.txt
                                                                 Pre-submit:
2 Mathematics
                                                                Write a few simple test cases if sample is not enough.
                                                                 Are time limits close? If so, generate max cases.
                                                                 Is the memory usage fine?
3 Data structures
                                                                Could anything overflow?
                                                                 Make sure to submit the right file.
4 Geometry
                                                             6
                                                                 Wrong answer:
                                                                 Print your solution! Print debug output, as well.
                                                                Are you clearing all data structures between test cases?
5 Graphs
                                                                 Can your algorithm handle the whole range of input?
                                                                 Read the full problem statement again.
6 Numerical Methods
                                                                Do you handle all corner cases correctly?
                                                                 Have you understood the problem correctly?
                                                                 Any uninitialized variables?
7 Number theory
                                                            20
                                                                 Any overflows?
                                                                 Confusing N and M, i and j, etc.?
                                                                 Are you sure your algorithm works?
8 Combinatorial
                                                                 What special cases have you not thought of?
                                                                 Are you sure the STL functions you use work as you think?
9 Strings
                                                                Add some assertions, maybe resubmit.
                                                                 Create some testcases to run your algorithm on.
                                                                 Go through the algorithm for a simple case.
10 Various
                                                                Go through this list again.
                                                                 Explain your algorithm to a teammate.
                                                                 Ask the teammate to look at your code.
Contest (1)
                                                                 Go for a small walk, e.g. to the toilet.
                                                                 Is your output format correct? (including whitespace)
                                                                 Rewrite your solution from the start or let a teammate do it.
template.cpp
                                                         14 lines
#include <bits/stdc++.h>
                                                                 Runtime error:
using namespace std;
                                                                 Have you tested all corner cases locally?
                                                                 Any uninitialized variables?
#define rep(i, a, b) for(int i = a; i < (b); ++i)
                                                                 Are you reading or writing outside the range of any vector?
\#define all(x) begin(x), end(x)
                                                                 Any assertions that might fail?
#define sz(x) (int)(x).size()
                                                                 Any possible division by 0? (mod 0 for example)
typedef long long 11;
                                                                 Any possible infinite recursion?
typedef pair<int, int> pii;
                                                                 Invalidated pointers or iterators?
typedef vector<int> vi;
                                                                 Are you using too much memory?
                                                                 Debug with resubmits (e.g. remapped signals, see Various).
int main()
 cin.tie(0)->sync_with_stdio(0);
                                                                 Time limit exceeded:
 cin.exceptions(cin.failbit);
                                                                 Do you have any possible infinite loops?
                                                                 What is the complexity of your algorithm?
                                                                 Are you copying a lot of unnecessary data? (References)
                                                                 How big is the input and output? (consider scanf)
.bashrc
                                                                 Avoid vector, map. (use arrays/unordered_map)
alias c='g++ -Wall -Wconversion -Wfatal-errors -g -std=c++14 \
                                                                 What do your teammates think about your algorithm?
 -fsanitize=undefined, address'
xmodmap -e 'clear lock' -e 'keycode 66=less greater' #caps = <>
                                                                 Memory limit exceeded:
                                                                 What is the max amount of memory your algorithm should need?
.vimrc
                                                                 Are you clearing all data structures between test cases?
                                                          6 lines
set cin aw ai is ts=4 sw=4 tm=50 nu noeb bg=dark ru cul
sy on | im jk <esc> | im kj <esc> | no;:
" Select region and then type : Hash to hash your selection.
" Useful for verifying that there aren't mistypes.
ca Hash w !cpp -dD -P -fpreprocessed \| tr -d '[:space:]' \
\| md5sum \| cut -c-6
hash.sh
                                                          6 lines
\ensuremath{\text{\#}} Hashes a file, ignoring all whitespace and comments. Use for
# verifying that code was correctly typed.
# Usage:
# To make executable, run the command: chmod +x hash.sh
# To execute: ./hash.sh < file.cpp
cpp -dD -P -fpreprocessed | tr -d '[:space:]' | md5sum |cut -c-6
```

template .bashrc .vimrc hash troubleshoot

Mathematics (2)

Equations

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

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

$$ax + by = e$$

$$cx + dy = f$$

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

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

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

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

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

2.2 Recurrences

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

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

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

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

Length of median (divides triangle into two equal-area triangles):

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

Length of bisector (divides angles in two):

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

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

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

2.4.2 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:

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

2.6 Sums

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

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

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

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

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

2.7

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

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

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

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

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

2.8 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), $0 \le p \le 1$.

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

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

Poisson distribution

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

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

$$\mu = \lambda, \, \sigma^2 = \lambda$$

2.8.2 Continuous distributions Uniform distribution

If the probability density function is constant between a and 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 $\text{Exp}(\lambda)$, $\lambda > 0$.

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

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

Normal distribution

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

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

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

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

2.9 Markov chains

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

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

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

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

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

2.9.1 Quadrilaterals

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

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

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

Data structures (3)

OrderStatisticTree.h

Description: A set (not multiset!) with support for finding the n'th element, and finding the index of an element. To get a map, change null_type. Time: $\mathcal{O}(\log N)$ 782797, 16 lines

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

HashMap.h

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

```
#include <bits/extc++.h>
// To use most bits rather than just the lowest ones:
struct chash { // large odd number for C
 const uint64_t C = 11(4e18 * acos(0)) | 71;
 11 operator()(11 x) const { return __builtin_bswap64(x*C); }
__gnu_pbds::gp_hash_table<11,int,chash> h({},{},{},{},{1<<16});</pre>
```

PST.h

Description: Persistent segment tree with laziness

Time: $\mathcal{O}(\log N)$ per query, $\mathcal{O}((n+q)\log n)$ memory

3656e8, 39 lines

```
struct PST {
 PST *1 = 0, *r = 0;
 int lo, hi;
 11 \text{ val} = 0, 1zadd = 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);
   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;
   if (L <= lo && hi <= R) {
     n = new PST(*this);
     n->val += v;
     n->1zadd += v;
    } else {
     push();
     n = new PST(*this);
     n->1 = 1->add(L, R, v);
```

```
n->r = r->add(L, R, v);
   return n;
 void push() {
   if(lzadd == 0) return;
    l = l \rightarrow add(lo, hi, lzadd);
   r = r -> add(lo, hi, lzadd);
   lzadd = 0;
};
```

UnionFindRollback.h

Description: Disjoint-set data structure with undo. If undo is not needed, skip st, time() and rollback().

Usage: int t = uf.time(); ...; uf.rollback(t);

Time: $\mathcal{O}(\log(N))$ de4ad0, 21 lines

```
struct RollbackUF {
 vi e; vector<pii> st;
 RollbackUF(int n) : e(n, -1) {}
 int size(int x) { return -e[find(x)]; }
 int find(int x) { return e[x] < 0 ? x : find(e[x]); }
 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;
};
```

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;
 11 div(11 a, 11 b) { // floored division
   return a / b - ((a ^ b) < 0 && a % b); }
 bool isect(iterator x, iterator y) {
   if (y == end()) return x \rightarrow p = inf, 0;
   if (x->k == y->k) x->p = x->m > y->m ? inf : -inf;
   else x->p = div(y->m - x->m, x->k - y->k);
   return x->p >= y->p;
 void add(ll k, ll m) {
   auto z = insert(\{k, m, 0\}), y = z++, x = y;
   while (isect(y, z)) z = erase(z);
   if (x != begin() \&\& isect(--x, y)) isect(x, y = erase(y));
   while ((y = x) != begin() \&\& (--x)->p >= y->p)
      isect(x, erase(y));
```

```
ll query(ll x) {
    assert(!empty());
    auto 1 = *lower_bound(x);
    return 1.k * x + 1.m;
};
```

Treap.h

Description: A short self-balancing tree. It acts as a sequential container with log-time splits/joins, and is easy to augment with additional data.

```
struct node {
    int val, prior, sz = 1;
    node *left = nullptr, *right = nullptr;
    node(int val = 0): val(val), prior(rand()) {}
int getSz(node *cur) { return cur ? cur->sz : 0; }
void recalc(node *cur) { cur->sz = getSz(cur->left) + getSz(cur
    ->right) + 1; }
pair<node*, node*> split(node *cur, int v) {
    if(!cur) return {nullptr, nullptr};
    node *left, *right;
    if (getSz(cur->left) >= v) {
        right = cur;
        auto [L, R] = split(cur->left, v);
        left = L, right->left = R;
        recalc(right);
    else {
        left = cur;
        auto [L, R] = split(cur->right, v - getSz(cur->left) -
        left->right = L, right = R;
        recalc(left);
    return {left, right};
node* merge(node *t1, node *t2) {
    if(!t1 || !t2) return t1 ? t1 : t2;
    node *res;
    if(t1->prior > t2->prior) {
        res = t1;
        res->right = merge(t1->right, t2);
    else {
        res = t2;
        res->left = merge(t1, t2->left);
    recalc(res);
    return res:
```

FenwickTree.h

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

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

e62fac, 22 lines

```
struct FT {
 vector<ll> s;
 FT(int n) : s(n) {}
 void update(int pos, 11 dif) { // a[pos] += dif
   for (; pos < sz(s); pos |= pos + 1) s[pos] += dif;
```

```
UCF
  11 query(int pos) { // sum of values in [0, pos)
    11 \text{ res} = 0;
    for (; pos > 0; pos &= pos -1) res += s[pos-1];
    return res;
  int lower_bound(ll sum) \{// min \ pos \ st \ sum \ of \ [0, \ pos] >= sum
    // Returns n if no sum is >= sum, or -1 if empty sum is.
    if (sum <= 0) return -1;
    int pos = 0;
    for (int pw = 1 << 25; pw; pw >>= 1) {
      if (pos + pw \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 ele-
Time: \mathcal{O}(\log^2 N). (Use persistent segment trees for \mathcal{O}(\log N).)
```

ments a[i,j]. Requires that the elements to be updated are known in advance (call fakeUpdate() before init()).

```
"FenwickTree.h"
struct FT2 {
  vector<vi> ys; vector<FT> ft;
  FT2(int limx) : vs(limx) {}
  void fakeUpdate(int x, int v) {
    for (; x < sz(ys); x |= x + 1) ys[x].push_back(y);
    for (vi& v : ys) sort(all(v)), ft.emplace_back(sz(v));
  int ind(int x, int y) {
   return (int)(lower_bound(all(ys[x]), y) - ys[x].begin()); }
  void update(int x, int y, ll dif) {
    for (; x < sz(ys); x |= x + 1)
      ft[x].update(ind(x, y), dif);
  11 query(int x, int y) {
   11 \text{ sum} = 0;
    for (; x; x &= x - 1)
     sum += ft[x-1].query(ind(x-1, y));
    return sum;
};
```

RMQ.h

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

Usage: RMQ rmq(values); rmq.query(inclusive, exclusive);

Time: $\mathcal{O}(|V|\log|V|+Q)$ 510c32, 16 lines

```
template<class T>
struct RMO {
  vector<vector<T>> jmp;
  RMQ(const vector<T>& V) : jmp(1, V) {
    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> 0)
 int L = 0, R = 0, blk = 350; // \sim N/sqrt(Q)
 vi s(sz(Q)), res = s;
#define K(x) pii(x.first/blk, x.second ^ -(x.first/blk & 1))
 iota(all(s), 0);
  sort(all(s), [\&](int s, int t) \{ return K(O[s]) < K(O[t]); \});
  for (int qi : s) {
    pii a = O[ai];
    while (L > q.first) add(--L, 0);
    while (R < q.second) add(R++, 1);
    while (L < q.first) del(L++, 0);
    while (R > g.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;
 1:
 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;
```

02776c, 16 lines

af592b, 14 lines

Geometry (4)

Point.hpp

Description: Class to handle points in the plane. T can be e.g. double or long long. (Avoid int.) 47ec0a, 35 lines

template < class T > int $sgn(T x) \{ return (x > 0) - (x < 0); \}$ template < class T > struct Point { typedef Point P; T x, y; explicit Point(T x = 0, T y = 0): x(x), y(y) {} bool operator<(P p) const { return tie(x, y) < tie(p.x, p.y);</pre> bool operator == (P p) const { return tie(x, y) == tie(p.x, p.y); P operator+(P p) const { return P(x + p.x, y + p.y); } P operator-(P p) const { return P(x - p.x, y - p.y); } P operator*(T d) const { return P(x * d, y * d); } P operator/(T d) const { return P(x / d, y / d); } T dot(P p) const { return x * p.x + y * p.y; } T cross(P p) const { return x * p.y - y * p.x; } T cross(P a, P b) const { return (a - *this).cross(b - *this); T dist2() const { return x * x + y * y; } double dist() const { return sqrt((double)dist2()); } // angle to x-axis in interval [-pi, pi] double angle() const { return atan2(y, x); } P unit() const { return *this / dist(); } // makes dist()=1 P perp() const { return P(-y, x); } // rotates +90 degrees P normal() const { return perp().unit(); } // returns point rotated 'a' radians ccw around the origin P rotate(double a) const { return P(x * cos(a) - y * sin(a), x * sin(a) + y * cos(a));friend ostream& operator << (ostream& os, P p) { return os << "(" << p.x << "," << p.y << ")";

4.1 Lines and Segments

Description: Checks what side of the line a point is on. Returns -1 if the point is left. 1 if right, and 0 if on the line. Orientation is based off the unit

template<class P> int sideOf(P s, P e, P p) { auto cp = s.cross(e, p); return (cp > eps) - (cp < -eps);

Description: Returns whether or not a point is on a segment c2edf6, 3 lines

```
template < class P > bool on Seg(P s, P e, P p) {
    return abs(s.cross(e, p)) < eps && (s-p).dot(e-p) < eps;
```

lineIntersection.hpp

Description: Returns the intersection point of two lines using cramers rule. If the lines are parallel or are the same, (inf, inf) is returned.

```
template < class P > P lineInter(P s1, P e1, P s2, P e2) {
    auto det = -(e1-s1).cross(e2-s2);
   if(abs(det) < eps) return P(inf, inf);</pre>
   auto t = (e2-s2).cross(s2-s1) / det;
   return s1 + (e1-s1) * t;
```

doSegIntersection.hpp

Description: Checks if two segments intersect (inclusive of intersections at endpoints) 59a2a4, 4 lines

```
template<class P> bool doSegInter(P s1, P e1, P s2, P e2) {
   return sideOf(s1, e1, s2) != sideOf(s1, e1, e2) &&
          sideOf(s2, e2, s1) != sideOf(s2, e2, e1);
```

segIntersection.hpp

Description: Returns the intersection point of two segments.

Usage: Returns a vector of points. If no points, there is no intersection. If 1 point, the segments intersect at a distinct point. If 2 points, the segments intersect at a segment of points, where the 2 points are the end points. 71bb02, 9 lines template<class P> vector<P> segInter(P s1, P e1, P s2, P e2) {

```
if(doSegInter(s1, e1, s2, e2)) return {lineInter(s1, e1, s2
set<P> seg;
 if (onSeg(s1, e1, s2)) seg.insert(s2);
 if(onSeg(s1, e1, e2)) seg.insert(e2);
 if(onSeg(s2, e2, s1)) seg.insert(s1);
 if(onSeg(s2, e2, e1)) seg.insert(e1);
  return {all(seg)};
```

lineDistance.hpp

Description: Gets the distance between a point and a line. 8f3d9d, 3 lines

```
template < class P > 1d lineDist(P s, P e, P p) {
   return (e-s).cross(p-s) / (e-s).mag();
```

segDistance.hpp

Description: Gets the distance between a point and a segment d02cd5, 5 lines

```
template<class P> ld segDist(P s, P e, P p) {
   if (s == e) return (p - s).mag();
   auto d = (e-s).mag2(), t = min(d, max(0.0, (p-s).dot(e-s)))
   return ((p - s)*d - (e - s)*t).mag() / d;
```

4.2 Polygons

polygonArea.hpp

Description: Uses shoelace theorem to find the area of a polygon.

Usage: If area is negative, points are given cw, otherwise points are given ccw.

Memory: $\mathcal{O}(1)$ Time: $\mathcal{O}(n)$

8b5fb1, 6 lines template<class T> ld polyArea(vector<pnt<T>> poly) { int n = sz(poly); T area = 0; for (int i = 0; i < n; i++) area += poly[i].cross(poly[(i+1)%n]); return area / 2.0L;

inPolygon.hpp

Description: Uses the cutting-ray test to see if a point is inside a polygon. Usage: Returns 0 if outside, 1 if strictly inside, and 2 if on. Memory: O(1) Time: $\mathcal{O}\left(n\right)$ 1ff9f1, 11 lines

```
template<class P> int inPoly(vector<P> poly, P p) {
   bool good = false; int n = sz(poly);
```

```
auto crosses = [](P s, P e, P p) {
    return ((e.y \ge p.y) - (s.y \ge p.y)) * p.cross(s, e) >
for (int i = 0; i < n; i++) {
    if(onSeg(poly[i], poly[(i+1)%n], p)) return 2;
    good ^= crosses(poly[i], poly[(i+1)%n], p);
return good;
```

convexHull.hpp

Description: gets the smallest convex polygon containing all points using monotone chaining.

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

```
template<class P> vector<P> convexHull(vector<P> poly) {
    int n = sz(polv);
    vector<P> hull(n+1);
    sort(all(poly));
    int k = 0;
    for (int i = 0; i < n; i++) {
        while (k \ge 2 \&\& hull[k-2].cross(hull[k-1], poly[i]) <=
             0) k--;
        hull[k++] = poly[i];
    for (int i = n-1, t = k+1; i > 0; i--) {
        while (k \ge t \&\& hull[k-2].cross(hull[k-1], poly[i-1])
            <= 0) k--;
        hull[k++] = poly[i-1];
```

hullDiameter.hpp

return hull;

hull.resize(k-1);

Description: Gets the furthest set of points on a convex polygon using rotating calibers

Memory: $\mathcal{O}(1)$

```
Time: \mathcal{O}(n)
template<class P> pair<P, P> hullDiameter(vector<P> poly) {
    int n = sz(polv); if (n < 3) return 0;
```

```
int j = 1;
pair<ld, pair<P, P>> ans(0, pair<P, P>(poly[0], poly[0]));
rep(i, 0, j){
    for(;; j = (j+1)%n){
        ans = max(ans, {(poly[i]-poly[j]).mag2(), {poly[i],}
             poly[j]}});
        if((poly[(j+1)%n] - poly[j]).cross(poly[i+1] - poly
             [i]) >= 0)
            break:
return ans.second;
```

hullTangents.hpp

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, $n \ge 3$, and the point be on or outside the polygon.

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

```
"Point.h"
#define cmp(i, j) p.cross(h[i], h[j == n ? 0 : j]) * (R ?: -1)
template<bool R, class P> int getTangent(vector<P>& h, P p) {
 int n = sz(h), lo = 0, hi = n - 1, md;
```

```
7
```

```
if (cmp(0, 1) >= R && cmp(0, n - 1) >= !R) return 0;
while (md = (lo + hi + 1) / 2, lo < hi) {
   auto a = cmp(md, md + 1), b = cmp(md, lo);
   if (a >= R && cmp(md, md - 1) >= !R) return md;
   if (cmp(lo, lo + 1) < R)
        a < R&& b >= 0 ? lo = md : hi = md - 1;
   else a < R || b <= 0 ? lo = md : hi = md - 1;
}
return -1; // point strictly inside hull
}
template<class P> pii hullTangents(vector<P>& h, P p) {
   return {getTangent<0>(h, p), getTangent<1>(h, p)};
}
```

inHull.hpp

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

Time: $\mathcal{O}(\log N)$ 6d9710, 12 lines template < class P > bool inHull (const vector < P > & 1, P p, bool strict = true) { int a = 1, b = sz(1) - 1, r = !strict; if (sz(1) < 3) return r & & onSegment(1[0], 1.back(), p); if (sideof(1[0], 1[a], 1[b]) > 0) swap(a, b); if (sideof(1[0], 1[a], p) >= r || sideof(1[0], 1[b], p) <= -r) return false; while (abs(a - b) > 1) { int c = (a + b) / 2; (sideof(1[0], 1[c], p) > 0 ? b : a) = c; } return sgn(1[a].cross(1[b], p)) < r;

polygonClipping.hpp

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.

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

Time: $\mathcal{O}\left(n\right)$ fd7bb9, 12 lines

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

halfplaneIntersection.hpp

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

```
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 operator<(HalfPlane hp) {
      if(abs(d.x) < eps && abs(hp.d.x) < eps)
            return d.y > 0 && hp.d.y < 0;
   bool side = d.x < eps || (abs(d.x) <= eps && d.y > 0);
```

```
bool sideHp = hp.d.x < eps || (abs(hp.d.x) <= eps && hp
             .d.y > 0);
        if(side != sideHp) 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), l = 1, r = 0;
    vector<HalfPlane> dq(n+1);
    rep(i, 0, n) {
        while (1 < r \&\& !hps[i].contains(dq[r].inter(dq[r-1])))
        while (1 < r \&\& !hps[i].contains(dq[l].inter(dq[l+1])))
            1++;
        dq[++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[1].inter(dq[1+1]))) 1
    if(1 > r - 2) return {};
    vector<P> poly;
    rep(i, 1, r)
       poly.push_back(dg[i].inter(dg[i+1]));
    poly.push_back(dq[r].inter(dq[l]));
    return poly;
```

centerOfMass.hpp

Description: Returns the center of mass for a polygon.

Memory: $\mathcal{O}(1)$

```
Time: O(n)

ccce20, 8 lines

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

minkowskiSum.hpp

d08058, 43 lines

Description: returns the minkowski sum of several polygons 13cd02, 30 lines

```
if(size(dir) == 0)
    return {init};
sort(all(dir), [&](P a, P b)->bool {
    bool sideA = a.x > 0 || (a.x == 0 && a.y > 0);
    bool sideB = b.x > 0 || (b.x == 0 && b.y > 0);
    if(sideA != sideB)
        return sideA;
    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;
```

4.3 Circles

CircleLine.hpp

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>.

CircleIntersection.hpp

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

CirclePolygonIntersection.hpp

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

```
#define arg(p, q) atan2(p.cross(q), p.dot(q))
template<class P> 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 det = a * a - b;
        if (det <= 0) return arg(p, q) * 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 = p + d * t;
        return arg(p, u) * r2;
    };
    auto sum = 0.0;</pre>
```

```
rep(i, 0, sz(ps))
 sum += tri(ps[i] - c, ps[(i + 1) % sz(ps)] - c);
return sum;
```

CircleTangents.hpp

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. b0153d, 13 lines

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

MinimumEnclosingCircle.hpp

Description: Computes the minimum circle that encloses a set of points.

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

7f7b1a, 27 lines

```
typedef pnt<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;
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};
```

4.4 3D Geometry

Point3D.hpp

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); }</pre>
 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;
};
```

PolyhedronVolume.hpp

Description: Magic formula for the volume of a polyhedron. Faces should point outwards. 3058c3, 6 lines

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

4.5 Miscellaneous

ClosestPair.hpp

Description: Finds the closest pair of points.

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

```
"Point.h"
                                                      ac41a6, 17 lines
typedef Point<11> P;
pair<P, P> closest (vector<P> v) {
 assert(sz(v) > 1);
 sort(all(v), [](P a, P b) { return a.y < b.y; });</pre>
 pair<11, pair<P, P>> ret{LLONG MAX, {P(), P()}};
 int j = 0;
 for (P p : v) {
   P d{1 + (ll)sqrt(ret.first), 0};
   while (v[j].y \le p.y - d.x) S.erase(v[j++]);
   auto lo = S.lower_bound(p - d), hi = S.upper_bound(p + d);
   for (; lo != hi; ++lo)
     ret = min(ret, \{(*lo - p).dist2(), \{*lo, p\}\});
   S.insert(p);
 return ret.second:
```

FastDelaunay.hpp

Description: Fast Delaunay triangulation. Each circumcircle contains none of the input points. There must be no duplicate points. If all points are on a line, no triangles will be returned. Should work for doubles as well, though there may be precision issues in 'circ'. Returns triangles in order {t[0][0], $t[0][1], t[0][2], t[1][0], \dots\}$, all counter-clockwise.

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

```
"Point.h"
                                                      eefdf5, 88 lines
typedef Point<11> P;
typedef struct Quad* Q;
typedef __int128_t lll; // (can be ll if coords are < 2e4)
P arb(LLONG_MAX, LLONG_MAX); // not equal to any other point
struct Quad {
  Q rot, o; P p = arb; bool mark;
  P& F() { return r()->p; }
  Q& r() { return rot->rot; }
  Q prev() { return rot->o->rot; }
  Q next() { return r()->prev(); }
bool circ(P p, P a, P b, P c) { // is p in the circumcircle?
  111 p2 = p.dist2(), A = a.dist2()-p2,
      B = b.dist2()-p2, C = c.dist2()-p2;
  return p.cross(a,b) *C + p.cross(b,c) *A + p.cross(c,a) *B > 0;
Q makeEdge(P orig, P dest) {
  Q r = H ? H : new Quad{new Quad{new Quad{0}}}};
  H = r -> 0; r -> r() -> r() = r;
  rep(i,0,4) r = r - rot, r - rot = arb, r - rot = i & 1 ? r : r - rot);
  r->p = orig; r->F() = dest;
  return r;
void splice(Q a, Q b) {
  swap(a->o->rot->o, b->o->rot->o); swap(a->o, b->o);
Q connect(Q a, Q b) {
  Q = makeEdge(a->F(), b->p);
  splice(q, a->next());
  splice(q->r(), b);
  return q;
pair<0,0> rec(const vector<P>& s) {
  if (sz(s) \le 3)
    Q = makeEdge(s[0], s[1]), b = makeEdge(s[1], s.back());
    if (sz(s) == 2) return { a, a->r() };
    splice(a->r(), b);
    auto side = s[0].cross(s[1], s[2]);
    0 c = side ? connect(b, a) : 0;
    return {side < 0 ? c->r() : a, side < 0 ? c : b->r() };
#define H(e) e \rightarrow F(), e \rightarrow p
#define valid(e) (e->F().cross(H(base)) > 0)
  O A, B, ra, rb;
  int half = sz(s) / 2;
  tie(ra, A) = rec({all(s) - half});
  tie(B, rb) = rec({sz(s) - half + all(s)});
  while ((B->p.cross(H(A)) < 0 \&\& (A = A->next())) | |
         (A->p.cross(H(B)) > 0 && (B = B->r()->o)));
  Q base = connect(B->r(), A);
  if (A->p == ra->p) ra = base->r();
  if (B->p == rb->p) rb = base;
#define DEL(e, init, dir) Q e = init->dir; if (valid(e)) \
    while (circ(e->dir->F(), H(base), e->F())) {
      Q t = e->dir; \setminus
      splice(e, e->prev()); \
      splice(e->r(), e->r()->prev()); \
```

UCF

```
e->o = H; H = e; e = t; \
  for (;;) {
   DEL(LC, base->r(), o); DEL(RC, base, prev());
   if (!valid(LC) && !valid(RC)) break;
   if (!valid(LC) || (valid(RC) && circ(H(RC), H(LC))))
     base = connect(RC, base->r());
   else
     base = connect(base->r(), LC->r());
 return { ra, rb };
vector<P> triangulate(vector<P> pts) {
 sort(all(pts)); assert(unique(all(pts)) == pts.end());
  if (sz(pts) < 2) return {};
 Q e = rec(pts).first;
 vector<Q>q=\{e\};
 int qi = 0;
  while (e->o->F().cross(e->F(), e->p) < 0) e = e->o;
#define ADD { Q c = e; do { c->mark = 1; pts.push_back(c->p); \
 q.push_back(c->r()); c = c->next(); } while (c != e); }
  ADD; pts.clear();
  while (qi < sz(q)) if (!(e = q[qi++]) \rightarrow mark) ADD;
  return pts;
```

struct mcmf {

Graphs (5)

5.1 Network flow

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) e4f62e, 56 lines

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

MinCostMaxFlowDijkstra.h

Description: If SPFA TLEs, swap the find_path function in MCMF with the one below and in_q with seen. If negative edge weights can occur, initialize pi with the shortest path from the source to each node using Bellman-Ford. Negative weight cycles not supported.

efdefd, 24 line

```
bool findPath(int s, int t) {
 fill(all(dist), inf);
 fill(all(seen), 0);
 dist[s] = 0:
  __gnu_pbds::priority_queue<pair<ll, int>> pg;
 vector<decltype(pg)::point iterator> its(n);
 pq.push({0, s});
 while(!pq.empty()) {
   auto [d, cur] = pq.top(); pq.pop(); d *= -1;
   seen[cur] = 1;
   if(dist[cur] < d) continue;</pre>
    for(int idx: adj[cur]) {
     auto [nxt, cap, f, wt] = edges[idx];
     11 nxtD = d + wt + pi[cur] - pi[nxt];
     if(f >= cap || nxtD >= dist[nxt] || seen[nxt]) continue;
     dist[nxt] = nxtD;
     par[nxt] = {cur, idx};
     if (its[nxt] == pq.end()) its[nxt] = pq.push({-nxtD, nxt})
     else pq.modify(its[nxt], {-nxtD, nxt});
 rep(i, 0, n) pi[i] = min(pi[i] + dist[i], inf);
 return seen[t];
```

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.

```
d7f0f1, 42 lines
struct Dinic {
 struct Edge {
    int to, rev;
    11 c, oc;
    11 flow() { return max(oc - c, OLL); } // if you need flows
 };
 vi lvl, ptr, q;
 vector<vector<Edge>> adj;
 \label{eq:definition} \mbox{Dinic(int } \mbox{n) : } \mbox{lvl(n), ptr(n), q(n), adj(n) } \{ \}
 void addEdge(int a, int b, ll c, ll rcap = 0) {
    adj[a].push_back({b, sz(adj[b]), c, c});
    adj[b].push_back({a, sz(adj[a]) - 1, rcap, rcap});
 11 dfs(int v, int t, 11 f) {
    if (v == t || !f) return f;
    for (int& i = ptr[v]; i < sz(adj[v]); i++) {
      Edge& e = adj[v][i];
      if (lvl[e.to] == lvl[v] + 1)
        if (ll p = dfs(e.to, t, min(f, e.c))) {
          e.c -= p, adj[e.to][e.rev].c += p;
          return p;
    return 0;
 11 calc(int s, int t) {
    11 flow = 0; q[0] = s;
    rep(L,0,31) do { // 'int L=30' maybe faster for random data
     lvl = ptr = vi(sz(q));
      int qi = 0, qe = lvl[s] = 1;
      while (qi < qe && !lvl[t]) {
        int v = q[qi++];
```

```
for (Edge e : 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; }
};
```

GlobalMinCut.h

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

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

8b0e19, 21 lines

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

GomoryHu.h

Description: Given a list of edges representing an undirected flow graph, returns edges of the Gomory-Hu tree. The max flow between any pair of vertices is given by minimum edge weight along the Gomory-Hu tree path.

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

MatroidIntersection.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

- $\operatorname{check}(\operatorname{int} x)$: 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: $R^2 N(M2.add + M1.check + M2.check) + R^3 M1.add + R^2 M1.clear + RNM2.clear$

```
"../data-structures/UnionFind.h"
                                                     9812a7, 60 lines
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());
   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) ml.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;
};
```

5.2 Matching

hopcroftKarp.h

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

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

```
Time: \mathcal{O}\left(\sqrt{V}E\right)
bool dfs(int a, int L, vector<vi>& q, vi& btoa, vi& A, vi& B) {
 if (A[a] != L) return 0;
 A[a] = -1;
 for (int b : q[a]) if (B[b] == L + 1) {
   B[b] = 0:
   if (btoa[b] == -1 \mid | dfs(btoa[b], L + 1, q, btoa, A, B))
      return btoa[b] = a, 1;
 return 0;
int hopcroftKarp(vector<vi>& g, vi& btoa) {
 int res = 0:
 vi A(q.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(g)) if(A[a] == 0) cur.push_back(a);
    for (int lay = 1;; lay++) {
     bool islast = 0;
     next.clear();
      for (int a : cur) for (int b : q[a]) {
       if (btoa[b] == -1) {
         B[b] = lay;
          islast = 1;
        else if (btoa[b] != a && !B[b]) {
         B[b] = lay;
          next.push_back(btoa[b]);
      if (islast) break;
      if (next.empty()) return res;
      for (int a : next) A[a] = lay;
      cur.swap(next);
   rep(a, 0, sz(q))
      res += dfs(a, 0, q, btoa, A, B);
```

DFSMatching.h

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

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

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 : g[di])
   if (!vis[e] && find(e, g, btoa, vis)) {
      btoa[e] = di;
      return 1;
   }
   return 0;
}
int dfsMatching(vector<vi>& g, vi& btoa) {
   vi vis;
   rep(i,0,sz(g)) {
      vis.assign(sz(btoa), 0);
      for (int j : g[i])
      if (find(j, g, btoa, vis)) {
        btoa[j] = i;
        break;
    }
}
return sz(btoa) - (int)count(all(btoa), -1);
}
```

MinimumVertexCover.h

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

```
"DFSMatching.h"
                                                    da4196, 20 lines
vi cover(vector<vi>& g, int n, int m) {
 vi match(m, -1);
 int res = dfsMatching(q, match);
 vector<bool> lfound(n, true), seen(m);
 for (int it : match) if (it != -1) lfound[it] = false;
 vi q, cover:
 rep(i,0,n) if (lfound[i]) g.push back(i);
  while (!q.empty()) {
   int i = q.back(); q.pop_back();
   lfound[i] = 1;
    for (int e : q[i]) if (!seen[e] && match[e] != -1) {
      seen[e] = true;
      q.push_back(match[e]);
 rep(i,0,n) if (!lfound[i]) cover.push_back(i);
 rep(i,0,m) if (seen[i]) cover.push_back(n+i);
 assert(sz(cover) == res);
 return cover;
```

WeightedMatching.h

Description: Given a weighted bipartite graph, matches every node on the left with a node on the right such that no nodes are in two matchings and the sum of the edge weights is minimal. Takes cost[N][M], where cost[i][j] = cost for L[i] to be matched with R[j] and returns (min cost, match), where L[i] is matched with R[match[i]]. Negate costs for max cost. Requires $N \leq M$.

```
Time: \mathcal{O}\left(N^2M\right) 1e0fe9, 31 lines
```

```
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 "dammy" 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]) {
```

GeneralMatching SCC BiconnectedComponents 2sat

```
auto cur = a[i0 - 1][j - 1] - u[i0] - v[j];
    if (cur < dist[j]) dist[j] = cur, pre[j] = j0;
    if (dist[j] < delta) delta = dist[j], j1 = j;
}
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</pre>
```

GeneralMatching.h

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

```
"../numerical/MatrixInverse-mod.h"
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) {
     ll \ a = modpow(A[fi][fj], mod-2);
     rep(i,0,M) if (has[i] && A[i][fj]) {
       11 b = A[i][fj] * a % mod;
        rep(j, 0, M) A[i][j] = (A[i][j] - A[fi][j] * b) % mod;
     swap(fi,fj);
  return ret;
```

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

```
76b5c9, 24 lines
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]) {
   do {
     x = z.back(); z.pop_back();
     comp[x] = ncomps;
     cont.push back(x);
    } while (x != i);
    f(cont); cont.clear();
    ncomps++;
 return val[j] = low;
template < class G, class F> void scc(G& q, F f) {
 int n = sz(q);
 val.assign(n, 0); comp.assign(n, -1);
 Time = ncomps = 0;
 rep(i,0,n) if (comp[i] < 0) dfs(i, g, f);
```

BiconnectedComponents.h

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: O(E+V)
```

st.resize(si);

2965e5, 33 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, e, y, top = me; for (auto pa : ed[at]) if (pa.second != par) { tie(y, e) = pa;if (num[y]) { top = min(top, num[y]); if (num[y] < me)</pre> 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()));

```
} 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);
}
```

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 $(\sim x)$.

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

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

```
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;
    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. Input should be a vector of (dest, global edge index), where for undirected graphs, forward/backward edges have the same index. Returns a list of nodes in the Eulerian path/cycle with src at both start and end, or empty list if no cycle/path exists. To get edge indices back, add second to s and ret. Time: $\mathcal{O}(V+E)$

```
vi eulerWalk(vector<vector<pii>>& gr, int nedges, int src=0) {
  int n = sz(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(qr[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 DAG. Output tree is in adj, directed down from the root.

```
Time: \mathcal{O}(V+E)
```

dd16e<u>b, 52 lines</u>

```
struct DominatorTree {
  vector<vi> adj, ans; // input edges, edges of dominator tree
       (directed tree downwards from root)
  vector<vi> radj, child, sdomChild;
  vi label, rlabel, sdom, dom;
 vi par, bes;
  int co = 0;
  DominatorTree(int n) {
   adj = ans = radj = child = sdomChild =
     vector<vector<int>>(n);
   label = rlabel = sdom = dom = par = bes = vector<int>(n);
  void add_edge(int a, int b) { adj[a].pb(b); }
  int get(int x) {
   if (par[x] != x)
     int t = get(par[x]);
     par[x] = par[par[x]];
     if (sdom[t] < sdom[bes[x]]) bes[x] = t;</pre>
    return bes[x];
  void dfs(int x) { // create DFS tree
   label[x] = ++co;
```

```
rlabel[co] = x;
   sdom[co] = par[co] = bes[co] = co;
   for (auto y : adj[x]) {
     if (!label[y]) {
       dfs(y);
       child[label[x]].pb(label[y]);
     radj[label[y]].pb(label[x]);
 void init(int root) {
   dfs(root);
   for (int i = co; i >= 1; --i) {
     for (auto j : radj[i])
       sdom[i] = min(sdom[i], sdom[get(j)]);
     if (i > 1) sdomChild[sdom[i]].pb(i);
     for (auto j : sdomChild[i]) {
       int k = get(j);
       if (sdom[j] == sdom[k]) dom[j] = sdom[j];
       else dom[j] = k;
     for (auto j : child[i]) par[j] = i;
    for (int i = 2; i < co + 1; ++i) {
     if (dom[i] != sdom[i]) dom[i] = dom[dom[i]];
     ans[rlabel[dom[i]]].pb(rlabel[i]);
 }
};
```

5.4 Coloring

EdgeColoring.h

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

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

```
e210e2, 31 lines
vi edgeColoring(int N, vector<pii> eds) {
 vi cc(N + 1), ret(sz(eds)), fan(N), free(N), loc;
 for (pii e : eds) ++cc[e.first], ++cc[e.second];
 int u, v, ncols = *max_element(all(cc)) + 1;
 vector<vi> adj(N, vi(ncols, -1));
 for (pii e : eds) {
   tie(u, v) = e;
   fan[0] = v;
   loc.assign(ncols, 0);
   int at = u, end = u, d, c = free[u], ind = 0, i = 0;
   while (d = free[v], !loc[d] \&\& (v = adj[u][d]) != -1)
     loc[d] = ++ind, cc[ind] = d, fan[ind] = v;
   cc[loc[d]] = c;
   for (int cd = d; at != -1; cd ^= c ^ d, at = adj[at][cd])
     swap(adj[at][cd], adj[end = at][cd ^ c ^ d]);
   while (adj[fan[i]][d] != -1) {
     int left = fan[i], right = fan[++i], e = cc[i];
     adj[u][e] = left;
     adj[left][e] = u;
     adj[right][e] = -1;
     free[right] = e;
   adj[u][d] = fan[i];
   adj[fan[i]][d] = u;
   for (int y : {fan[0], u, end})
     for (int& z = free[y] = 0; adj[y][z] != -1; z++);
   for (tie(u, v) = eds[i]; adj[u][ret[i]] != v;) ++ret[i];
 return ret;
```

5.5 Heuristics

MaximalCliques.h

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

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

b0d5b1, 12 lines

```
typedef bitset<128> B;
template<class F>
void cliques(vector<B>& eds, F f, B P = \simB(), B X={}, B R={}) {
 if (!P.any()) { if (!X.any()) f(R); return; }
  auto g = (P | X)._Find_first();
  auto cands = P & ~eds[q];
  rep(i,0,sz(eds)) if (cands[i]) {
    cliques(eds, f, P & eds[i], X & eds[i], R);
    R[i] = P[i] = 0; X[i] = 1;
```

MaximumClique.h

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

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

f7c0bc, 49 lines

```
typedef vector<br/>bitset<200>> vb;
struct Maxclique {
 double limit=0.025, pk=0;
 struct Vertex { int i, d=0; };
 typedef vector<Vertex> vv;
  vb e;
  vv V;
  vector<vi> C:
  vi qmax, q, S, old;
  void init(vv& r) {
    for (auto \& v : r) v.d = 0;
    for (auto& v : r) for (auto j : r) v.d += e[v.i][j.i];
    sort(all(r), [](auto a, auto b) { return a.d > b.d; });
    int mxD = r[0].d;
    rep(i, 0, sz(r)) r[i].d = min(i, mxD) + 1;
 void expand(vv& R, int lev = 1) {
    S[lev] += S[lev - 1] - old[lev];
    old[lev] = S[lev - 1];
    while (sz(R)) {
      if (sz(q) + R.back().d <= sz(qmax)) return;</pre>
      g.push_back(R.back().i);
      vv T;
      for(auto v:R) if (e[R.back().i][v.i]) T.push_back({v.i});
        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();

6f34db, 46 lines

```
}
}
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 MinimumVertexCover.

5.6 Trees

BinaryLifting.h

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

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

bfce85, 25 lines

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

LCA.h

Description: Data structure for computing lowest common ancestors in a tree (with 0 as root). C should be an adjacency list of the tree, either directed or undirected.

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

```
"../data-structures/RMQ.h"

Struct LCA {
   int T = 0;
   vi time, path, ret;
   RMQ<int> rmq;

LCA(vector<vi>& C) : time(sz(C)), rmq((dfs(C,0,-1), ret)) {}
   void dfs(vector<vi>& C, int v, int par) {
        time[v] = T++;
        for (int y : C[v]) if (y != par) {
            path.push_back(v), ret.push_back(time[v]);
            dfs(C, y, v);
        }
   }
   int lca(int a, int b) {
        if (a == b) return a;
        tie(a, b) = minmax(time[a], time[b]);
   }
}
```

```
return path[rmq.query(a, b)];
}
//dist(a,b){return depth[a] + depth[b] - 2*depth[lca(a,b)];};
```

CentroidDecomp.h

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

c4d5af, 32 lines

```
template <class F> struct centroid {
   vector<vector<int>> adi;
   F f;
   vector<int> sub sz:
   centroid(const vector<vector<int>>& a_adj, F a_f)
        : adj(a_adj), f(a_f), sub_sz(ssize(adj), -1) {
        for (int i = 0; i < ssize(adj); i++)
            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];
   void dfs(int u) {
       calc_sz(u, -1);
       for (int p = -1, sz\_root = sub\_sz[u];;) {
            auto big_ch = find_if(begin(adj[u]), end(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(begin(adj[v]), end(adj[v]), u),
                rbegin(adj[v]));
            adj[v].pop_back();
            dfs(v);
};
```

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(1i)-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;
}
```

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.

Time: $\mathcal{O}\left((\log N)^2\right)$ ".../data-structures/LazySegmentTree.h"

```
template <bool VALS_EDGES> struct HLD {
 int N, tim = 0;
 vector<vi> adi;
 vi par, siz, depth, rt, pos;
 Node *tree;
 HLD(vector<vi> adj_)
   : N(sz(adj_)), adj(adj_), par(N, -1), siz(N, 1), depth(N),
      rt(N), pos(N), tree(new Node(0, N)) { dfsSz(0); dfsHld(0); }
 void dfsSz(int v) {
    if (par[v] != -1) adj[v].erase(find(all(adj[v]), par[v]));
    for (int& u : adj[v]) {
     par[u] = v, depth[u] = depth[v] + 1;
     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 (; rt[u] != rt[v]; v = par[rt[v]]) {
     if (depth[rt[u]] > depth[rt[v]]) swap(u, v);
     op(pos[rt[v]], pos[v] + 1);
   if (depth[u] > depth[v]) swap(u, v);
    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 $\mathcal{O}(\log N)$.

5909e2, 90 lines

```
struct Node { // Splay tree. Root's pp contains tree's parent.
 Node *p = 0, *pp = 0, *c[2];
 bool flip = 0;
 Node() { c[0] = c[1] = 0; fix(); }
  void fix() {
   if (c[0]) c[0] -> p = this;
   if (c[1]) c[1] -> p = this;
   // (+ update sum of subtree elements etc. if wanted)
  void 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];
     z -> c[h ^1] = b ? x : this;
   y - c[i ^ 1] = b ? this : x;
    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;
     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->splav();
   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()
   kev.w += delta;
   if (1) 1->delta += delta;
   if (r) r->delta += delta;
   delta = 0;
 Edge top() { prop(); return key; }
Node *merge(Node *a, Node *b) {
 if (!a || !b) return a ?: b;
 a->prop(), b->prop();
 if (a->key.w > b->key.w) swap(a, b);
 swap(a->1, (a->r = merge(b, a->r)));
 return a;
void pop(Node*\& a) { a->prop(); a = merge(a->1, a->r); }
pair<ll, vi> dmst(int n, int r, vector<Edge>& g) {
 RollbackUF uf(n);
 vector<Node*> heap(n);
  for (Edge e : 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>>> cvcs;
 rep(s,0,n) {
   int u = s, qi = 0, w;
    while (seen[u] < 0) {
      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 * cvc = 0;
```

```
int end = qi, time = uf.time();
    do cyc = merge(cyc, heap[w = path[--qi]]);
    while (uf.join(u, w));
    u = uf.find(u), heap[u] = cyc, seen[u] = -1;
        cycs.push_front({u, time, {&Q[qi], &Q[end]}});
}

rep(i,0,qi) in[uf.find(Q[i].b)] = Q[i];

for (auto& [u,t,comp] : cycs) { // restore sol (optional)
    uf.rollback(t);
    Edge inEdge = in[u];
    for (auto& e : comp) in[uf.find(e.b)] = e;
    in[uf.find(inEdge.b)] = inEdge;
}
rep(i,0,n) par[i] = in[i].a;
return {res, par};
}
```

15

Numerical Methods (6)

6.1 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}},-le9,le9) // solve $x^2-3x+2=0$ **Time:** $\mathcal{O}(n^2 \log(1/\epsilon))$

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; 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 $((f \le 0) ^ sign) 1 = m;$ else h = m;ret.push_back((1 + h) / 2); return ret;

PolvInterpolate.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:** $\mathcal{O}\left(n^2\right)$

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 < n.

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

```
"../number-theory/ModPow.h"
                                                     96548b, 20 lines
vector<1l> berlekampMassey(vector<1l> s) {
 int n = sz(s), L = 0, m = 0;
 vector<ll> C(n), B(n), T;
 C[0] = B[0] = 1;
 11 b = 1;
 rep(i,0,n) { ++m;
   11 d = s[i] % mod;
   rep(j,1,L+1) d = (d + C[j] * s[i - j]) % mod;
   if (!d) continue;
   T = C; 11 coef = d * modpow(b, mod-2) % mod;
    rep(j,m,n) C[j] = (C[j] - coef * B[j - m]) % mod;
   if (2 * L > i) continue;
   L = i + 1 - L; B = T; b = d; m = 0;
 C.resize(L + 1); C.erase(C.begin());
 for (11& x : C) x = (mod - x) % mod;
 return C;
```

LinearRecurrence.h

Description: Generates the k'th term of an n-order linear recurrence $S[i] = \sum_j S[i-j-1]tr[j]$, given $S[0\ldots \geq n-1]$ and $tr[0\ldots n-1]$. Faster than matrix multiplication. Useful together with Berlekamp–Massey.

Usage: linearRec($\{0, 1\}, \{1, 1\}, k$) // k'th Fibonacci number Time: $\mathcal{O}\left(n^2\log k\right)$ f4e444.26 lines

```
typedef vector<ll> Polv:
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);
 11 \text{ res} = 0;
 rep(i, 0, n) res = (res + pol[i + 1] * S[i]) % mod;
 return res:
```

6.2 Optimization

GoldenSectionSearch.h

HillClimbing.h

Description: Poor man's optimization for unimodal functions_{8eeeaf, 14 lines}

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

template<class F> pair<double, P> hillClimb(P start, F f) {
  pair<double, P> cur(f(start), start);
  for (double jmp = 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;
}
```

IntegrateAdaptiveTyler.h

Description: Gets area under a curve

e7heba 17 line

RungeKutta4.h

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

Sc1ac, 12 lines

```
template < class F, class T>
T solveSystem(F f, T x, ld 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^Tx subject to $Ax \leq b$, $x \geq 0$. Returns -inf if there is no solution, inf if there are arbitrarily good solutions, or the maximum value of c^Tx otherwise. The input vector is set to an optimal x (or in the unbounded case, an arbitrary solution fulfilling the constraints). Numerical stability is not guaranteed. For better performance, define variables such that x = 0 is viable.

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

bool simplex(int phase) {

for (;;) {

int x = m + phase - 1;

rep(j,0,n+1) if (N[j] != -phase) ltj(D[x]);

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

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

```
const T eps = 1e-8, inf = 1/.0;
#define MP make pair
struct LPSolver {
 int m, n;
 vi N, B;
 vvd D;
  LPSolver (const vvd& A, const vd& b, const vd& c) :
   m(sz(b)), n(sz(c)), N(n+1), B(m), D(m+2), vd(n+2)) {
     rep(i, 0, m) rep(j, 0, n) D[i][j] = A[i][j];
     rep(i, 0, m) \{ B[i] = n+i; D[i][n] = -1; D[i][n+1] = b[i]; \}
     rep(j, 0, n) \{ N[j] = j; D[m][j] = -c[j]; \}
     N[n] = -1; D[m+1][n] = 1;
  void pivot(int r, int s) {
   T *a = D[r].data(), inv = 1 / a[s];
   rep(i, 0, m+2) if (i != r \&\& abs(D[i][s]) > eps) {
     T *b = D[i].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]);
```

```
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;</pre>
    rep(i, 0, m) if (B[i] == -1) {
      int s = 0;
      rep(j,1,n+1) ltj(D[i]);
      pivot(i, s);
  bool ok = simplex(1); x = vd(n);
  rep(i, 0, m) if (B[i] < n) x[B[i]] = D[i][n+1];
  return ok ? D[m][n+1] : inf;
```

6.3 Matrices

Determinant.h

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

```
double det(vector<vector<double>>& a) {
   int n = sz(a); double res = 1;
   rep(i,0,n) {
    int b = i;
   rep(j,i+1,n) if (fabs(a[j][i]) > fabs(a[b][i])) b = j;
   if (i != b) swap(a[i], a[b]), res *= -1;
   res *= a[i][i];
   if (res == 0) return 0;
   rep(j,i+1,n) {
      double v = a[j][i] / a[i][i];
      if (v != 0) rep(k,i+1,n) a[j][k] -= v * a[i][k];
   }
}
return res;
}
```

IntDeterminant.h

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

```
Time: \mathcal{O}\left(N^3\right) 3313dc, 18 lines const 11 mod = 12345;

11 det(vector<vector<11>>& a) {

  int n = sz(a); 11 ans = 1;

  rep(i,0,n) {

    rep(j,i+1,n) {

    while (a[j][i] != 0) { // gcd step}

        11 t = a[i][i] / a[j][i];

        if (t) rep(k,i,n)

        a[i][k] = (a[i][k] - a[j][k] * t) % mod;

        swap(a[i], a[j]);

        ans *= -1;

    }
```

```
ans = ans * a[i][i] % mod;
if (!ans) return 0;
}
return (ans + mod) % mod;
}
```

SolveLinear.h

Description: Solves A*x=b. If there are multiple solutions, an arbitrary one is returned. Returns rank, or -1 if no solutions. Data in A and b is lost. **Time:** $\mathcal{O}\left(n^2m\right)$

```
typedef vector<double> vd;
const double eps = 1e-12;
int solveLinear(vector<vd>& A, vd& b, vd& x) {
 int n = sz(A), m = sz(x), rank = 0, br, bc;
 if (n) assert(sz(A[0]) == m);
 vi col(m); iota(all(col), 0);
 rep(i,0,n) {
    double v, bv = 0;
    rep(r,i,n) rep(c,i,m)
     if ((v = fabs(A[r][c])) > bv)
       br = r, bc = c, bv = v;
    if (bv <= eps) {
      rep(j,i,n) if (fabs(b[j]) > eps) return -1;
      break;
    swap(A[i], A[br]);
    swap(b[i], b[br]);
    swap(col[i], col[bc]);
    rep(j,0,n) swap(A[j][i], A[j][bc]);
    bv = 1/A[i][i];
    rep(j,i+1,n) {
      double fac = A[j][i] * bv;
      b[j] = fac * b[i];
      rep(k, i+1, m) A[j][k] -= fac*A[i][k];
    rank++;
  x.assign(m, 0);
  for (int i = rank; i--;) {
   b[i] /= A[i][i];
    x[col[i]] = b[i];
    rep(j, 0, i) b[j] -= A[j][i] * b[i];
 return rank; // (multiple solutions if rank < m)
```

SolveLinear2.h

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

SolveLinearBinary.h

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

```
typedef bitset<1000> bs;
```

6af945, 10 lines

```
int solveLinear(vector<bs>& A, vi& b, bs& x, int m) {
 int n = sz(A), rank = 0, br;
  assert(m \le sz(x));
  vi col(m); iota(all(col), 0);
  rep(i,0,n) {
   for (br=i; br<n; ++br) if (A[br].any()) break;</pre>
   if (br == n) {
     rep(j,i,n) if(b[j]) return -1;
     break;
    int bc = (int)A[br]._Find_next(i-1);
    swap(A[i], A[br]);
    swap(b[i], b[br]);
    swap(col[i], col[bc]);
    rep(j, 0, n) if (A[j][i] != A[j][bc]) {
     A[j].flip(i); A[j].flip(bc);
    rep(j,i+1,n) if (A[j][i]) {
     b[j] ^= b[i];
     A[j] ^= A[i];
    rank++;
  x = bs();
  for (int i = rank; i--;) {
   if (!b[i]) continue;
   x[col[i]] = 1;
   rep(j,0,i) b[j] ^= A[j][i];
  return rank; // (multiple solutions if rank < m)
```

MatrixInverse.h

Description: Invert matrix A. Returns rank; result is stored in A unless singular (rank < n). Can easily be extended to prime moduli; for prime powers, repeatedly set $A^{-1} = A^{-1}(2I - AA^{-1})$ (mod p^k) where A^{-1} starts as the inverse of A mod p, and k is doubled in each step. Time: $\mathcal{O}(n^3)$

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

```
rep(k,0,n) tmp[j][k] -= v*tmp[i][k];
}
rep(i,0,n) rep(j,0,n) A[col[i]][col[j]] = tmp[i][j];
return n;
```

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"
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;
    A[i].swap(A[r]); tmp[i].swap(tmp[r]);
    rep(j,0,n) swap(A[j][i], A[j][c]), swap(tmp[j][i], tmp[j][c
    swap(col[i], col[c]);
    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) {
   11 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
        : 0);
 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
```

```
\begin{split} \{a_i\} &= \operatorname{tridiagonal}(\{1,-1,-1,\ldots,-1,1\},\{0,c_1,c_2,\ldots,c_n\},\\ \{b_1,b_2,\ldots,b_n,0\},\{a_0,d_1,d_2,\ldots,d_n,a_{n+1}\}). \end{split}
```

Fails if the solution is not unique. If $|d_i| > |p_i| + |q_{i-1}|$ for all i, or $|d_i| > |p_{i-1}| + |q_i|$, or the matrix is positive definite, the algorithm is numerically stable and neither tr nor the check for diag[i] == 0 is needed. Time: $\mathcal{O}(N)$

```
typedef double T;
vector<T> tridiagonal(vector<T> diag, const vector<T>& super,
    const vector<T>& sub, vector<T> b) {
  int n = sz(b); vi tr(n);
  rep(i,0,n-1) {
    if (abs(diag[i]) < 1e-9 * abs(super[i])) { // diag[i] == 0}
      b[i+1] -= b[i] * diag[i+1] / super[i];
      if (i+2 < n) b[i+2] -= b[i] * sub[i+1] / super[i];</pre>
      diag[i+1] = sub[i]; tr[++i] = 1;
      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;
}
x[i] -= eps;
}
return J;
```

NewtonsMethod.h

Description: Solves a system on non-linear equations

```
jacobianMatrix.h
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;
   }
}
```

6.4 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: $\operatorname{conv}(a, b) = c$, where $c[x] = \sum_x a[i]b[x-i]$. 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 $(\sum_x a_i^2 + \sum_y b_i^2) \log_2 N < 9 \cdot 10^{14}$ (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 *= 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 *= 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);

FastFourierTransformMod.h

return res;

"FastFourierTransform.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)

```
typedef vector<1l> v1;
template<int M> v1 convMod(const v1 &a, const v1 &b) {
   if (a.empty() || b.empty()) return {};
   v1 res(ss(a) + sz(b) - 1);
   int B=32-_builtin_clz(sz(res)), n=1<<B, cut=int(sqrt(M));
   vector<C> L(n), R(n), outs(n), out1(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);
    out1[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(out1), fft(outs);
   rep(i,0,sz(res)) {
```

```
11 av = ll(real(outl[i])+.5), cv = ll(imag(outs[i])+.5);
    11 bv = ll(imag(outl[i])+.5) + ll(real(outs[i])+.5);
    res[i] = ((av % M * cut + bv) % M * cut + cv) % M;
}
return res;
}
```

NumberTheoreticTransform.h

Description: ntt(a) computes $\hat{f}(k) = \sum_x a[x]g^{xk}$ for all k, where $g = \operatorname{root}^{(mod-1)/N}$. N must be a power of 2. Useful for convolution modulo specific nice primes of the form 2^ab+1 , where the convolution result has size at most 2^a . For arbitrary modulo, see FFTMod. $\operatorname{conv}(a, b) = c$, 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)
                                                     ced03d, 33 lines
"../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 - \underline{builtin_clz(n)};
  static vl rt(2, 1);
  for (static int k = 2, s = 2; k < n; k *= 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;
  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
  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)] = (11)L[i] * R[i] % mod * inv %
  return {out.begin(), out.begin() + s};
```

FastSubsetTransform.h

b82773, 22 lines

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)
```

```
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;
}
```

Number theory (7)

7.1 Modular arithmetic

ModInverse.h

Description: Pre-computation of modular inverses. Assumes LIM \leq mod and that mod is a prime.

```
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;
```

ModLog.h

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. **Time:** $\mathcal{O}(\sqrt{m})$

```
11 modLog(11 a, 11 b, 11 m) {
    11 n = (11) sqrt(m) + 1, e = 1, f = 1, j = 1;
    unordered_map<11, 11> 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

Description: Sums of mod'ed arithmetic progressions. modsum(to, c, k, m) = $\sum_{i=0}^{\text{to}-1} (ki+c)\%m$. divsum is similar but for floored division.

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

5c5bc5, 16 line

```
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 bbbd8f, 11 lines

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

ModSart.h

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

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

```
"ModPow.h"
                                                        19a793, 24 lines
ll sgrt(ll a, ll p) {
 a \% = p; if (a < 0) a += p;
 if (a == 0) return 0;
 assert (modpow(a, (p-1)/2, p) == 1); // else no solution
 if (p % 4 == 3) return modpow(a, (p+1)/4, p);
 // a^{(n+3)/8} \text{ or } 2^{(n+3)/8} * 2^{(n-1)/4} \text{ works if } p \% 8 == 5
 11 s = p - 1, n = 2;
 int r = 0, m;
 while (s % 2 == 0)
    ++r, s /= 2;
  while (modpow(n, (p - 1) / 2, p) != p - 1) ++n;
 11 x = modpow(a, (s + 1) / 2, p);
 ll b = modpow(a, s, p), g = modpow(n, s, p);
 for (;; r = m) {
    11 t = b;
    for (m = 0; m < r && t != 1; ++m)
     t = t * t % p;
    if (m == 0) return x;
   11 \text{ gs} = \text{modpow}(g, 1LL << (r - m - 1), p);
   q = qs * qs % p;
   x = x * qs % p;
   b = b * q % p;
```

7.2 Primality

FastEratosthenes.h

Description: Prime sieve for generating all primes smaller than LIM.

Time: LIM=1e9 ≈ 1.5 s 6b2912, 20 lines

```
const int LIM = 1e6;
bitset<LIM> isPrime:
vi eratosthenes() {
 const int S = (int)round(sqrt(LIM)), R = LIM / 2;
 vi pr = \{2\}, sieve(S+1); pr.reserve(int(LIM/log(LIM) \star1.1));
  vector<pii> cp;
  for (int i = 3; i <= S; i += 2) if (!sieve[i]) {
    cp.push_back({i, i * i / 2});
    for (int j = i * i; j <= S; j += 2 * i) sieve[j] = 1;
  for (int L = 1; L <= R; L += S) {
    array<bool, S> block{};
    for (auto &[p, idx] : cp)
     for (int i=idx; i < S+L; idx = (i+=p)) block[i-L] = 1;
    rep(i, 0, min(S, R - L))
      if (!block[i]) pr.push_back((L + i) * 2 + 1);
 for (int i : pr) isPrime[i] = 1;
 return pr;
```

LinearSieve.h

Description: Finds smallest prime factor of each integer **Time:** $\mathcal{O}\left(N\right)$

```
const int LIM = 1000000;
vi lp(LIM+1), primes;
rep(i, 2, LIM + 1) {
   if (lp[i] == 0) primes.push_back(lp[i] = i);
   for (int j = 0; j < sz(primes) && i * primes[j] <= LIM &&
        primes[j] <= lp[i]; ++j)
   lp[i * primes[j]] = primes[j];
```

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;
}
```

Factor.h

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

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

```
"ModMulLL.h", "MillerRabin.h"
                                                      a33cf6, 18 lines
ull pollard(ull n) {
 auto f = [n] (ull x) \{ return modmul(x, x, n) + 1; \};
 ull x = 0, y = 0, t = 30, prd = 2, i = 1, q;
 while (t++ % 40 || __gcd(prd, n) == 1) {
   if (x == y) x = ++i, y = f(x);
    if ((q = modmul(prd, max(x,y) - min(x,y), n))) prd = q;
    x = f(x), y = f(f(y));
 return __gcd(prd, n);
vector<ull> factor(ull n) {
 if (n == 1) return {};
 if (isPrime(n)) return {n};
 ull x = pollard(n);
  auto 1 = factor(x), r = factor(n / x);
 l.insert(l.end(), all(r));
 return 1:
```

GetFactors.h

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

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

void getFactors(auto &pF, auto &primes, auto &factors, int i =
 0, int n = 1) {
 if(i == sz(pF)) {
 factors.push_back(n);
 return;
 }

 for(int j = 0, pow = 1; i <= pf[j]; j++, pow *= primes[j])
 getFactors(pF, primes, factors, i+1, n * pow);
}</pre>

7.3 Divisibility

euclid.h

32eeca, 8 lines

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

```
11 euclid(ll a, ll b, ll &x, ll &y) {
  if (!b) return x = 1, y = 0, a;
```

CRT phiFunction ContinuedFractions FracBinarySearch // If b > a/2, we have a semi-convergent that gives us a

```
ll d = euclid(b, a % b, y, 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 $0 \le x \le \operatorname{lcm}(m, n)$. Assumes $mn \le 2^{62}$

Time: $\log(n)$

```
04d93a, 7 lines
11 crt(l1 a, l1 m, l1 b, l1 n) {
  if (n > m) swap(a, b), swap(m, n);
  ll x, y, g = euclid(m, n, x, y);
  assert((a - b) % g == 0); // else no solution
  x = (b - a) % n * x % n / q * m + a;
  return x < 0 ? x + m*n/g : x;
```

7.3.1 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 $\leq n$ that are coprime with n. $\phi(1) = 1$, p prime $\Rightarrow \phi(p^k) =$ $(p-1)p^{k-1}$, m, n coprime $\Rightarrow \phi(mn) = \phi(m)\phi(n)$. If $n = p_1^{k_1} p_2^{k_2} ... p_r^{k_r}$ then $\phi(n) = (p_1 - 1)p_1^{k_1 - 1}...(p_r - 1)p_r^{k_r - 1}$. $\phi(n) = n \cdot \prod_{p|n} (1 - 1/p)$. $\sum_{d \mid n} \phi(d) = n, \ \sum_{1 \le k \le n, \gcd(k,n) = 1} k = n\phi(n)/2, n > 1$ Euler's thm: a, ncoprime $\Rightarrow a^{\phi(n)} \equiv 1 \pmod{n}$. Fermat's little thm: p prime $\Rightarrow a^{p-1} \equiv 1$ $\pmod{p} \ \forall a.$

const int LIM = 5000000; int phi[LIM]; void calculatePhi() { rep(i, 0, LIM) phi[i] = i&1 ? i : i/2;for (int i = 3; i < LIM; i += 2) if(phi[i] == i)</pre> for (int j = i; j < LIM; j += i) phi[j] -= phi[j] / i;

7.4 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, $p_{k+1}q_k - q_{k+1}p_k = (-1)^k$. $(p_k/q_k \text{ alternates between }$ > 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: $\mathcal{O}(\log N)$

if (a > b) {

```
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; dy = x;
  for (;;) {
    ll lim = min(P ? (N-LP) / P : inf, Q ? (N-LQ) / Q : inf),
       a = (ll) floor(y), b = min(a, lim),
       NP = b*P + LP, NQ = b*Q + LQ;
```

```
// 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, NO};
LP = P; P = NP;
LQ = Q; Q = NQ;
```

FracBinarySearch.h

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

Usage: fracBS([](Frac f) { return f.p>=3*f.q; }, 10); // {1,3} Time: $\mathcal{O}(\log(N))$ 27ab3e, 25 lines

```
struct Frac { ll p, q; };
template<class F>
Frac fracBS(F f, 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) {
      adv += step;
     Frac mid{lo.p * adv + hi.p, lo.q * adv + hi.q};
     if (abs(mid.p) > N || mid.q > N || dir == !f(mid)) {
       adv -= step; si = 2;
   hi.p += lo.p * adv;
   hi.q += lo.q * adv;
    dir = !dir;
    swap(lo, hi);
   A = B; B = !!adv;
 return dir ? hi : lo;
```

7.5 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.

7.6 Primes

dd6c5e, 21 lines

p = 962592769 is such that $2^{21} \mid p - 1$, which may be useful. For hashing use 970592641 (31-bit number), 31443539979727 (45-bit), 3006703054056749 (52-bit). There are 78498 primes less than 1000000.

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

7.7 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.

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:

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

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

IntPerm multinomial

Combinatorial (8)

8.1 Permutations

8.1.1 Factorial

	123							
n!	1 2 6	24 1	20 72	0 5040	4032	20 362	2880 36	528800
	11							
n!	4.0e7	4.8e	8 6.26	e9 8.7ε	e10 1.5	3e12 2	2.1e13	3.6e14
								171
n!	2e18	2e25	3e32	8e47	3e64 9	9e157	6e262	>DBL_MAX

IntPerm.h

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

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

int permToInt(vi& v) {
 int use = 0, i = 0, r = 0;
 for(int x:v) r = r * ++i + __builtin_popcount(use & -(1<<x)),
 use |= 1 << x;
 return r;</pre>

8.1.2 Cycles

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

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

8.1.3 Derangements

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

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

8.1.4 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

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

.2 Partitions and subsets

8.2.1 Partition function

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

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

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

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

8.2.3 Binomials

multinomial.h

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

11 multinomial(vi& v) {
 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;
}

8.3 General purpose numbers

8.3.1 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:

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

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

8.3.2 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 $c(n,2) = 0, 0, 1, 3, 11, 50, 274, 1764, 13068, 109584, \dots$

8.3.3 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. $\pi(j) > j$.

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

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

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

8.3.4 Stirling numbers of the second kind

Partitions of n distinct elements into exactly k groups.

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

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

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

8.3.5 Bell numbers

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

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

8.3.6 Labeled unrooted trees

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

8.3.7 Catalan numbers

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

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

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

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

Strings (9)

KMP.h

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

Time: $\mathcal{O}(n)$ d4375c, 16 lines vi pi(const string& s) { vi p(sz(s));rep(i,1,sz(s))int g = p[i-1];while (g && s[i] != s[g]) g = p[g-1];p[i] = g + (s[i] == s[g]);return p: vi match (const string& s, const string& pat) { vi p = $pi(pat + ' \setminus 0' + s)$, res; rep(i, sz(p) - sz(s), sz(p))if (p[i] == sz(pat)) res.push_back(i - 2 * sz(pat)); return res;

Zfunc.h

Description: z[x] computes the length of the longest common prefix of s[i:]and s, except z[0] = 0. (abacaba -> 0010301)

Time: $\mathcal{O}(n)$ 3ae526, 12 lines vi Z(string S) { vi z(sz(S)); int 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

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

Time: $\mathcal{O}(N)$ e7ad79, 13 lines array<vi, 2> manacher(const string& s) { int n = sz(s); $array < vi, 2 > p = {vi(n+1), vi(n)};$ 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;

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

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

2052a7, 24 lines

struct Eertree {

```
vector<int> slink = \{0, 0\}, len = \{-1, 0\};
 vector<vector<int>> down:
 int cur = 0, t = 0;
 Eertree(string &str) : down(2, vector<int>(26, -1)) {
    for (int i = 0; i < str.size(); i++) {</pre>
     char c = str[i]; int ci = c - 'a';
     while (t \le 0 | | str[t-1] != c)
     t = i - len[cur = slink[cur]];
     if (down[cur][ci] == -1) {
     down[cur][ci] = slink.size();
     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 = slink.size() - 1;
     } else cur = down[cur][ci];
     t = i - len[cur] + 1;
};
```

MinRotation.h

Description: Finds the lexicographically smallest rotation of a string. Usage: rotate(v.begin(), v.begin()+minRotation(v), v.end()); Time: $\mathcal{O}(N)$ d07a42, 8 lines

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

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

```
38db9f, 23 lines
struct SuffixArray {
 vi sa, lcp;
 SuffixArrav(string& s, int lim=256) { // or basic_string<int>
   int n = sz(s) + 1, k = 0, a, b;
   vi x(all(s)+1), y(n), ws(max(n, lim)), rank(n);
   sa = lcp = y, iota(all(sa), 0);
    for (int j = 0, p = 0; p < n; j = max(1, j * 2), lim = p) {
     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++;
   rep(i,1,n) rank[sa[i]] = i;
    for (int i = 0, j; i < n - 1; lcp[rank[i++]] = k)
     for (k \&\& k--, j = sa[rank[i] - 1];
         s[i + k] == s[j + k]; k++);
};
```

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 $K = \text{Alphabet Size}_{1914a9, 22 \text{ lines}}$

```
struct st { int len, pos, term; st *link; map<char, st*> next;
st *suffixAutomaton(string &str) {
 st *last = new st(), *root = last;
  for(auto c : str) {
    st *p = last, *cur = last = new st{last->len + 1, last->len}
         };
    while(p && !p->next.count(c))
      p->next[c] = cur, p = p->link;
    if (!p) cur->link = root;
      st *a = p->next[c];
      if (p\rightarrow len + 1 == q\rightarrow len) cur\rightarrow link = q;
        st *clone = new st{p->len+1, q->pos, 0, q->link, q->}
        for (; p && p->next[c] == q; p = p->link)
          p->next[c] = clone;
        g->link = cur->link = clone;
  while(last) last->term = 1, last = last->link;
  return root;
```

SuffixTree.h

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

Time: $\mathcal{O}(26N)$

aae0b8, 50 lines

```
struct SuffixTree {
 enum { N = 200010, ALPHA = 26 }; //N \sim 2*maxlen+10
 int toi(char c) { return c - 'a'; }
  string a; //v = cur \ node, q = cur \ position
  int t[N][ALPHA], 1[N], r[N], p[N], s[N], v=0, q=0, m=2;
  void ukkadd(int i, int c) { suff:
    if (r[v] \le 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]; }</pre>
      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 (l[node] <= i1 && i1 < r[node]) return 1;</pre>
    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 = ( 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 ~ 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 {};
  H h = 0, pw = 1;
  rep(i,0,length)
  h = h * C + str[i], pw = pw * C;
  vector<H> ret = {h};
  rep(i,length,sz(str)) {
    ret.push_back(h = h * C + str[i] - pw * str[i-length]);
  return ret;
```

```
H hashString(string& s){H h{}; for(char c:s) h=h*C+c; return h;}
```

AhoCorasick-Tyler.h

Description: Aho-Corasick automaton, used for multiple pattern matching. Initialize with AhoCorasick ac(patterns); the automaton start node will be at index 0. find(word) returns for each position the index of the longest word that ends there, or -1 if none. findAll(-, word) finds all words (up to $N\sqrt{N}$ many if no duplicate patterns) that start at each position (shortest first). Duplicate patterns are allowed; empty patterns are not. To find the longest words that start at each position, reverse all input. For large alphabets, split each symbol into chunks, with sentinel bits for symbol boundaries.

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

```
const int ABSIZE = 26;
struct node {
    int nxt[ABSIZE]:
   vi ids = {};
    int prv = -1, link = -1;
    int linkMemo[ABSIZE];
   node (int prv = -1, char c = '$'): prv(prv), c(c) {
        fill(all(nxt), -1);
        fill(all(linkMemo), -1);
vector<node> trie(1);
void addWord(string &s, int id) {
   int cur = 0;
    for(char c: s) {
       int idx = c - 'a';
       if(trie[cur].nxt[idx] == -1) {
            trie[cur].nxt[idx] = sz(trie);
            trie.emplace_back(cur, c);
        cur = trie[cur].nxt[idx];
   trie[cur].ids.push_back(id);
int getLink(int cur);
int calc(int cur, char c) {
   int idx = c - 'a';
    auto &ret = trie[cur].linkMemo[idx];
   if (ret != -1) return ret;
   if(trie[cur].nxt[idx] != -1)
       return ret = trie[cur].nxt[idx];
    return ret = cur == 0 ? 0 : calc(getLink(cur), c);
int getLink(int cur) {
   auto &ret = trie[cur].link;
   if (ret != -1) return ret;
   if(cur == 0 || trie[cur].prv == 0) return ret = 0;
    return ret = calc(getLink(trie[cur].prv), trie[cur].c);
```

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;

IntervalCover.h

if (R != r2) is.emplace (R, r2);

Description: Compute indices of smallest set of intervals covering another interval. Intervals should be [inclusive, exclusive). To support [inclusive, inclusive], change (A) to add | | R.empty(). Returns empty set on failure (or if G is empty).

Time: $\mathcal{O}(N \log N)$ 9e9d8d, 19 lines

```
template<class T>
vi cover(pair<T, T> G, vector<pair<T, T>> I) {
  vi S(sz(I)), R;
  iota(all(S), 0);
  sort(all(S), [&](int a, int b) { return I[a] < I[b]; });</pre>
  T cur = G.first;
  int at = 0;
  while (cur < G.second) { // (A)
   pair<T, int> mx = make_pair(cur, -1);
   while (at < sz(I) && I[S[at]].first <= cur) {
     mx = max(mx, make_pair(I[S[at]].second, S[at]));
   if (mx.second == -1) return {};
   cur = mx.first;
   R.push_back (mx.second);
  return R;
```

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)\{\ldots\});
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) {
   g(i, to, p);
   i = to; p = q;
   int mid = (from + to) >> 1;
   rec(from, mid, f, g, 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, g, i, p, q);
 q(i, to, q);
```

10.2 Misc. algorithms

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

```
2932a0, 17 lines
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:
```

FastKnapsack.h

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

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

```
b20ccc, 16 lines
int knapsack(vi w, int t) {
 int a = 0, b = 0, x;
 while (b < sz(w) & & a + w[b] <= t) a += w[b++];
 if (b == sz(w)) return a;
 int m = *max element(all(w));
 vi u, v(2*m, -1);
 v[a+m-t] = b;
 rep(i,b,sz(w))
   rep(x,0,m) v[x+w[i]] = max(v[x+w[i]], u[x]);
   for (x = 2*m; --x > m;) rep(j, max(0,u[x]), v[x])
     v[x-w[j]] = max(v[x-w[j]], j);
 for (a = t; v[a+m-t] < 0; a--);
 return a:
```

10.3 Dynamic programming

KnuthDP.h

Description: When doing DP on intervals: $a[i][j] = \min_{i < k < j} (a[i][k] + a[i][k])$ 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 $f(b,c) \leq f(a,d)$ 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: $\mathcal{O}(N^2)$

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. Time: $\mathcal{O}\left(\left(N + (hi - lo)\right) \log N\right)$

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

10.4 Debugging tricks

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

10.5 Optimization tricks

builtin ia32 ldmxcsr(40896); disables denormals (which make floats 20x slower near their minimum value).

10.5.1 Bit backs

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

10.5.2Pragmas

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

FastMod.h

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

```
typedef unsigned long long ull;
struct FastMod {
  ull b, m;
  FastMod(ull b) : b(b), m(-1ULL / b) {}
  ull reduce (ull a) { // a \% b + (0 \text{ or } b)
    return a - (ull) ((__uint128_t(m) * a) >> 64) * b;
};
```

FastInput.h

Description: Read an integer from stdin. Usage requires your program to pipe in input from file.

Usage: ./a.out < input.txt

Time: About 5x as fast as cin/scanf.

```
7b3c70, 17 lines
inline char gc() { // like getchar()
  static char buf[1 << 16];
  static size t bc, be;
  if (bc >= be) {
   buf[0] = 0, bc = 0;
   be = fread(buf, 1, sizeof(buf), stdin);
  return buf[bc++]; // returns 0 on EOF
int readInt() {
 int a, c;
  while ((a = gc()) < 40);
  if (a == '-') return -readInt();
  while ((c = qc()) >= 48) a = a * 10 + c - 480;
  return a - 48;
```

BumpAllocator.h

Description: When you need to dynamically allocate many objects and don't care about freeing them. "new X" otherwise has an overhead of something like 0.05us + 16 bytes per allocation. 745db2, 8 lines

```
// Either globally or in a single class:
static char buf[450 << 20];
void* operator new(size t s) {
 static size_t i = sizeof buf;
  assert(s < i);
  return (void*) &buf[i -= s];
void operator delete(void*) {}
```

```
template < class T > struct ptr {
  unsigned ind;
```

```
SmallPtr.h
Description: A 32-bit pointer that points into BumpAllocator memory.
"BumpAllocator.h"
                                                            2dd6c9, 10 lines
```

```
ptr(T*p = 0) : ind(p ? unsigned((char*)p - buf) : 0) {
  assert(ind < sizeof buf);
T& operator*() const { return *(T*)(buf + ind); }
T* operator->() const { return &**this; }
T& operator[](int a) const { return (&**this)[a]; }
explicit operator bool() const { return ind; }
```

BumpAllocatorSTL.h

Description: BumpAllocator for STL containers.

```
Usage: vector<vector<int, small<int>>> ed(N);
                                                     bb66d4, 14 lines
```

```
char buf[450 << 20] alignas(16);</pre>
size t buf ind = sizeof buf;
template<class T> struct small {
 typedef T value_type;
 small() {}
 template < class U > small(const U&) {}
 T* allocate(size_t n) {
   buf ind -= n * sizeof(T);
   buf_ind &= 0 - alignof(T);
   return (T*) (buf + buf_ind);
 void deallocate(T*, size t) {}
```

SIMD.h

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

```
551b82, 43 lines
#pragma GCC target ("avx2") // or sse4.1
#include "immintrin.h"
typedef __m256i mi;
#define L(x) _mm256_loadu_si256((mi*)&(x))
// High-level/specific methods:
// load(u)?\_si256, store(u)?\_si256, setzero\_si256, \_mm\_malloc
// blendv_(epi8|ps|pd) (z?y:x), movemask_epi8 (hibits of bytes)
// i32gather_epi32(addr, x, 4): map addr[] over 32-b parts of x
// sad_epu8: sum of absolute differences of u8, outputs 4xi64
// maddubs_epi16: dot product of unsigned i7's, outputs 16xi15
// madd_epi16: dot product of signed i16's, outputs 8xi32
// extractf128_si256(, i) (256->128), cvtsi128_si32 (128->lo32)
// permute2f128_si256(x,x,1) swaps 128-bit lanes
// shuffle_epi32(x, 3*64+2*16+1*4+0) == x for each lane
// shuffle_epi8(x, y) takes a vector instead of an imm
// Methods that work with most data types (append e.g. _epi32):
// set1, blend (i8?x:y), add, adds (sat.), mullo, sub, and/or,
// and not, abs, min, max, sign(1,x), cmp(gt|eq), unpack(lo|hi)
int sumi32 (mi m) { union {int v[8]; mi m;} u; u.m = m;
 int ret = 0; rep(i,0,8) ret += u.v[i]; return ret; }
mi zero() { return _mm256_setzero_si256(); }
mi one() { return _mm256_set1_epi32(-1); }
bool all_zero(mi m) { return _mm256_testz_si256(m, m); }
bool all_one(mi m) { return _mm256_testc_si256(m, one()); }
11 example_filteredDotProduct(int n, short* a, short* b) {
```

```
int i = 0; 11 r = 0;
mi zero = _mm256_setzero_si256(), acc = zero;
while (i + 16 \le n) {
  mi \ va = L(a[i]), \ vb = L(b[i]); \ i += 16;
  va = _mm256_and_si256(_mm256_cmpgt_epi16(vb, va), va);
  mi vp = _mm256_madd_epi16(va, vb);
  acc = _mm256_add_epi64(_mm256_unpacklo_epi32(vp, zero),
    _mm256_add_epi64(acc, _mm256_unpackhi_epi32(vp, zero)));
union \{11 \ v[4]; \ mi \ m;\} \ u; \ u.m = acc; \ rep(i,0,4) \ r += u.v[i];
for (;i < n; ++i) if (a[i] < b[i]) r += a[i] *b[i]; // <- equiv
return r;
```