



**FPT UNIVERSITY**

FPT University

Dolphin

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2	Mathematics
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## Contest (1)

template.cpp

42 lines

```
#include <bits/stdc++.h>
using namespace std;

#define rep(i, a, b) for(int i = a; i < (b); ++i)
#define all(x) begin(x), end(x)
#define sz(x) (int)(x).size()
#define PI acos(-1)

using ll = long long;
using db = double;

typedef complex<double> base;
typedef vector<base> vb;
typedef pair<int, int> pii;
typedef vector<int> vi;

const int ALPHABET_SIZE = 26;
const int BASE = 31;
const int MAXN = 100000001;
const int INF = 1e9;
const int NBIT = 18;
const int N = 1<<18;
const int MOD = (int)1e9+7;

void solve() {

}

int main() {
    #ifndef ONLINE_JUDGE
    freopen("INPUT.INP", "r", stdin);
    freopen("OUTPUT.OUT", "w", stdout);
}
```

```
1     #endif
1     ios_base::sync_with_stdio(false);
1     cin.tie(0); cout.tie(0);
2     int test = 1;
2     cin >> test;
2     while (test--) {
3         solve();
3     }
4 }
```

## Mathematics (2)

### 2.1 Equations

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

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

$$\begin{aligned} 12 \quad ax + by = e &\Rightarrow x = \frac{ed - bf}{ad - bc} \\ 17 \quad cx + dy = f &\Rightarrow y = \frac{af - ec}{ad - bc} \end{aligned}$$

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  $a_n = c_1 a_{n-1} + \dots + c_k a_{n-k}$ , and  $r_1, \dots, r_k$  are distinct roots of  $x^k - c_1 x^{k-1} - \dots - c_k$ , there are  $d_1, \dots, d_k$  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

$$\begin{aligned} \sin(v+w) &= \sin v \cos w + \cos v \sin w \\ \cos(v+w) &= \cos v \cos w - \sin v \sin w \end{aligned}$$

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

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

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

$$\begin{aligned} a \cos x + b \sin x &= r \cos(x-\phi) \\ a \sin x + b \cos x &= r \sin(x+\phi) \end{aligned}$$

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

## 2.4 Geometry

### 2.4.1 Triangles

Side lengths:  $a, b, c$

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

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

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

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

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

Length of bisector (divides angles in two):

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

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

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

### 2.4.2 Quadrilaterals

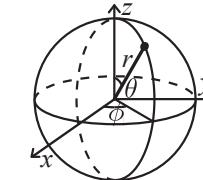
$$\text{law of tangents: } \frac{a+b}{a-b} = \frac{c+d}{c-d}$$

With side lengths  $a, b, c, d$ , diagonals  $e, f$ , diagonals angle  $\theta$ , area  $A$  and magic flux  $F = b^2 + d^2 - \frac{a^2 + c^2}{4}$ :

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

### 2.4.3 Spherical coordinates

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



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

## 2.5 Derivatives/Integrals

$$\begin{aligned} \frac{d}{dx} \arcsin x &= \frac{1}{\sqrt{1-x^2}} & \frac{d}{dx} \arccos x &= -\frac{1}{\sqrt{1-x^2}} \\ \frac{d}{dx} \tan x &= 1 + \tan^2 x & \frac{d}{dx} \arctan x &= \frac{1}{1+x^2} \\ \int \tan ax \, dx &= -\frac{\ln |\cos ax|}{a} & \int x \sin ax \, dx &= \frac{\sin ax - ax \cos ax}{a^2} \\ \int e^{-x^2} \, dx &= \frac{\sqrt{\pi}}{2} \operatorname{erf}(x) & \int xe^{ax} \, dx &= \frac{e^{ax}}{a^2} (ax - 1) \end{aligned}$$

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 Series

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

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

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

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

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

## template

### 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  $\sigma$  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  $Y$ ,

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

#### 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  $\text{Bin}(n, p)$ ,  $n = 1, 2, \dots$ ,  $0 \leq p \leq 1$ .

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

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

$\text{Bin}(n, p)$  is approximately  $\text{Po}(np)$  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  $\text{Fs}(p)$ ,  $0 \leq p \leq 1$ .

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

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

##### Poisson distribution

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

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

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

### 2.8.2 Continuous distributions

##### Uniform distribution

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

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

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

##### Exponential distribution

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

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

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

##### Normal distribution

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

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

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

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

## 2.9 Markov chains

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

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

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

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

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

## Data structures (3)

### OrderStatisticTree.h

**Description:** A set (not multiset!) with support for finding the  $n$ 'th element, and finding the index of an element. To get a map, change `null_type`.

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

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
}
```

### SegmentTree.h

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

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

0f4bdb, 29 lines

```
struct Tree {
    typedef int T;
    static constexpr T unit = INT_MIN;
    T f(T a, T b) { return max(a, b); } // max
    // static constexpr T unit = INT_MAX;
    // T f(T a, T b) { return min(a, b); } // min
    static constexpr T unit = 0;
    // T f(T a, T b) { return a + b; } // sum
    static constexpr T unit = 0;
    // T f(T a, T b) { return __gcd(a, b); } // GCD
    static constexpr T unit = 1;
    // T f(T a, T b) { return (a*b)/__gcd(a, b); } // GCD
    vector<T> s; int n;
    Tree(int n = 0, T def = unit) : s(2*n, def), n(n) {}
    void update(int pos, T val) {
        // condition
        // for (s[pos += n] = (val > 0 ? 1 : 0); pos /= 2;) {
        for (s[pos += n] = val; pos /= 2;) {
            s[pos] = f(s[pos * 2], s[pos * 2 + 1]);
        }
    }
    T query(int b, int e) { // query [b, e)
        T ra = unit, rb = unit;
        for (b += n, e += n; b < e; b /= 2, e /= 2) {
            if (b % 2) ra = f(ra, s[b++]);
            if (e % 2) rb = f(s[-e], rb);
        }
        return f(ra, rb);
    }
};
```

### LazySegmentTree.h

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

**Usage:** `Node* tr = new Node(v, 0, sz(v))`

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

..../various/BumpAllocator.h" 34ecf5, 48 lines

```
const int inf = 1e9; // Change to 0 for sum or INT_MAX for min
struct Node {
    Node* l = 0, *r = 0;
    int lo, hi, mset = inf, madd = 0, val = -inf; // Change val to match the problem
    Node(int lo, int hi) : lo(lo), hi(hi) {} // Default value for a large interval
    Node(vi& v, int lo, int hi) : lo(lo), hi(hi) {
        if (lo + 1 < hi) {
            int mid = lo + (hi - lo) / 2;
            l = new Node(v, lo, mid); r = new Node(v, mid, hi);
            val = max(l->val, r->val); // Combine function (change to + for sum or min queries)
        } else val = v[lo];
    }
    int query(int L, int R) {
        if (R <= lo || hi <= L) return -inf; // Default return value (change to 0 for sum or inf for min)
        if (L <= lo && hi <= R) return val;
        push();
        return max(l->query(L, R), r->query(L, R)); // Combine function
    }
    void set(int L, int R, int x) {
        if (R <= lo || hi <= L) return;
        if (L <= lo && hi <= R) mset = val = x, madd = 0;
        else {
            push(), l->set(L, R, x), r->set(L, R, x);
            val = max(l->val, r->val); // Combine function
        }
    }
    void add(int L, int R, int x) {
        if (R <= lo || hi <= L) return;
        if (L <= lo && hi <= R) {
            if (mset != inf) mset += x;
            else madd += x;
            val += x; // Modify logic for add if necessary
        } else {
            push(), l->add(L, R, x), r->add(L, R, x);
            val = max(l->val, r->val); // Combine function
        }
    }
    void push() {
        if (!l) {
            int mid = lo + (hi - lo) / 2;
            l = new Node(lo, mid); r = new Node(mid, hi);
        }
        if (mset != inf)
            l->set(lo, hi, mset), r->set(lo, hi, mset), mset = inf;
        else if (madd)
            l->add(lo, hi, madd), r->add(lo, hi, madd), madd = 0;
    }
};
```

### UnionFind.h

**Description:** Disjoint-set data structure.

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

7aa27c, 14 lines

```
struct UF {
    vi e;
    UF(int n) : e(n, -1) {}
```

```
bool sameSet(int a, int b) { return find(a) == find(b); }
int size(int x) { return -e[find(x)]; }
int find(int x) { return e[x] < 0 ? x : e[x] = find(e[x]); }
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);
    e[a] += e[b]; e[b] = a;
    return true;
}
};
```

### 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;
        }
};
```

### SubMatrix.h

**Description:** Calculate submatrix sums quickly, given upper-left and lower-right corners (half-open).

**Usage:** `SubMatrix<int> m(matrix);`  
`m.sum(0, 0, 2, 2); // top left 4 elements`

**Time:**  $\mathcal{O}(N^2 + Q)$

c59ada, 13 lines

```
template<class T>
struct SubMatrix {
    vector<vector<T>> p;
    SubMatrix(vector<vector<T>> v) {
        int R = sz(v), C = sz(v[0]);
        p.assign(R+1, vector<T>(C+1));
        rep(r, 0, R) rep(c, 0, C)
            p[r+1][c+1] = v[r][c] + p[r][c+1] + p[r+1][c] - p[r][c];
    }
    T sum(int u, int l, int d, int r) {
        return p[d][r] - p[d][l] - p[u][r] + p[u][l];
    }
};
```

### Matrix.h

**Description:** Basic operations on square matrices.

**Usage:** `Matrix<int, 3> A;`  
`A.d = {{1,2,3}, {4,5,6}, {7,8,9}};`  
`vector<int> vec = {1,2,3};`  
`vec = (A^N) * vec;`

c43c7d, 26 lines

```
template<class T, int N> struct Matrix {
```

```

typedef Matrix M;
array<array<T, N>, N> d{};
M operator*(const M& m) const {
    M a;
    rep(i, 0, N) rep(j, 0, N)
        rep(k, 0, N) a.d[i][j] += d[i][k]*m.d[k][j];
    return a;
}
vector<T> operator*(const vector<T>& vec) const {
    vector<T> ret(N);
    rep(i, 0, N) rep(j, 0, N) ret[i] += d[i][j] * vec[j];
    return ret;
}
M operator^(ll p) const {
    assert(p >= 0);
    M a, b(*this);
    rep(i, 0, N) a.d[i][i] = 1;
    while (p) {
        if (p&1) a = a*b;
        b = b*b;
        p >>= 1;
    }
    return a;
}
};

LineContainer.h

```

**Description:** Container where you can add lines of the form  $kx+m$ , and query maximum values at points  $x$ . Useful for dynamic programming (“convex hull trick”).  
**Time:**  $\mathcal{O}(\log N)$

sec1c7, 30 lines

```

struct Line {
    mutable ll k, m, p;
    bool operator<(const Line& o) const { return k < o.k; }
    bool operator<(ll x) const { return p < x; }
};

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

```

FenwickTree.h

**Description:** Computes partial sums  $a[0] + a[1] + \dots + a[pos - 1]$ , and updates single elements  $a[i]$ , taking the difference between the old and new value.  
**Time:** Both operations are  $\mathcal{O}(\log N)$ .

e62fac, 22 lines

```

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

FenwickTree2d.h

```

**Description:** Computes sums  $a[i,j]$  for all  $i < I, j < J$ , and increases single elements  $a[i,j]$ . Requires that the elements to be updated are known in advance (call fakeUpdate() before init()).  
**Time:**  $\mathcal{O}(\log^2 N)$ . (Use persistent segment trees for  $\mathcal{O}(\log N)$ .)

"FenwickTree2d.h"

157f07, 22 lines

```

struct FT2 {
    vector<vi> ys; vector<FT> ft;
    FT2(int limx) : ys(limx) {}
    void fakeUpdate(int x, int y) {
        for (; x < sz(ys); x |= x + 1) ys[x].push_back(y);
    }
    void init() {
        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);
    }
    ll query(int x, int y) {
        ll sum = 0;
        for (; x; x &= x - 1)
            sum += ft[x-1].query(ind(x-1, y));
        return sum;
    }
};

```

## Numerical (4)

### 4.1 Polynomials and recurrences

Polynomial.h

c9b7b0, 17 lines

```

struct Poly {
    vector<double> a;
    double operator()(double x) const {
        double val = 0;
        for (int i = sz(a); i--;) (val *= x) += a[i];
        return val;
    }
    void diff() {
        rep(i, 1, sz(a)) a[i-1] = i*a[i];
        a.pop_back();
    }
};

```

```

    }
    void divroot(double x0) {
        double b = a.back(), c; a.back() = 0;
        for (int i=sz(a)-1; i--;) c = a[i], a[i] = a[i+1]*x0+b, b=c;
        a.pop_back();
    }
};

PolyRoots.h

```

**Description:** Finds the real roots to a polynomial.  
**Usage:** polyRoots({{2,-3,1}}, -1e9, 1e9) // solve  $x^2-3x+2 = 0$   
**Time:**  $\mathcal{O}(n^2 \log(1/\epsilon))$

"Polynomial.h" b00bfe, 23 lines

```

vector<double> polyRoots(Poly p, double xmin, double xmax) {
    if (sz(p.a) == 2) { return {-p.a[0]/p.a[1]}; }
    vector<double> ret;
    Poly der = p;
    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(l) > 0;
        if (sign ^ (p(h) > 0)) {
            rep(it, 0, 60) { // while (h - l > 1e-8)
                double m = (l + h) / 2, f = p(m);
                if ((f <= 0) ^ sign) l = m;
                else h = m;
            }
            ret.push_back((l + h) / 2);
        }
    }
    return ret;
}

```

PolyInterpolate.h

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

08bf48, 13 lines

```

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

```

BerlekampMassey.h

**Description:** Recovers any  $n$ -order linear recurrence relation from the first  $2n$  terms of the recurrence. Useful for guessing linear recurrences after brute-forcing the first terms. Should work on any field, but numerical stability for floats is not guaranteed. Output will have size  $\leq n$ .  
**Usage:** berlekampMassey({0, 1, 1, 3, 5, 11}) // {1, 2}

"../number-theory/ModPow.h" 96548b, 20 lines

```

vector<ll> berlekampMassey(vector<ll> s) {
    int n = sz(s), L = 0, m = 0;
    vector<ll> C(n), B(n), T;

```

```
C[0] = B[0] = 1;
ll b = 1;
rep(i,0,n) { ++m;
  ll d = s[i] % mod;
  rep(j,1,L+1) d = (d + C[j] * s[i - j]) % mod;
  if (!d) continue;
  T = C; ll 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 (ll& 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  $S[0 \dots \geq n - 1]$  and  $tr[0 \dots n - 1]$ . Faster than matrix multiplication. Useful together with Berlekamp–Massey.

**Usage:** linearRec({0, 1}, {1, 1}, k) //  $k$ 'th Fibonacci number  
**Time:**  $\mathcal{O}(n^2 \log k)$

f4e444, 26 lines

```
typedef vector<ll> Poly;
ll linearRec(Poly S, Poly tr, ll k) {
  int n = sz(tr);
}
```

```
auto combine = [&](Poly a, Poly b) {
  Poly res(n * 2 + 1);
  rep(i,0,n+1) rep(j,0,n+1)
    res[i + j] = (res[i + j] + a[i] * b[j]) % mod;
  for (int i = 2 * n; i > n; --i) rep(j,0,n)
    res[i - 1 - j] = (res[i - 1 - j] + res[i] * tr[j]) % mod;
  res.resize(n + 1);
  return res;
};
```

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

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

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

**4.2 Matrices****Determinant.h**

**Description:** Calculates determinant of a matrix. Destroys the matrix.

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

bd5cec, 15 lines

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

**IntDeterminant.h**

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

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

3313de, 18 lines

```
const ll mod = 12345;
ll det(vector<vector<ll>>& a) {
  int n = sz(a); ll ans = 1;
  rep(i,0,n) {
    rep(j,i+1,n) {
      while (a[j][i] != 0) { // gcd step
        ll t = a[i][i] / a[j][i];
        if (t) rep(k,i,n)
          a[i][k] = (a[i][k] - a[j][k] * t) % mod;
        swap(a[i], a[j]);
        ans *= -1;
      }
      ans = ans * 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:**  $\mathcal{O}(n^2 m)$

44c9ab, 38 lines

```
typedef vector<double> vd;
const double eps = 1e-12;

int solveLinear(vector<vd>& A, vd& b, vd& x) {
  int n = sz(A), m = sz(x), rank = 0, br, bc;
  if (n) assert(sz(A[0]) == m);
  vi col(m); iota(all(col), 0);
  rep(i,0,n) {
    double v, bv = 0;
    rep(r,i,n) rep(c,i,m)
      if ((v = fabs(A[r][c])) > bv)
        br = r, bc = c, bv = v;
    if (bv <= eps) {
      rep(j,i,n) if (fabs(b[j]) > eps) return -1;
      break;
    }
    swap(A[i], A[br]);
    swap(b[i], b[br]);
    swap(col[i], col[bc]);
    rep(j,0,n) swap(A[j][i], A[j][bc]);
    bv = 1/A[i][i];
    rep(j,i+1,n) {
      double fac = A[j][i] * bv;
      b[j] -= fac * b[i];
      rep(k,i+1,m) A[j][k] -= fac * A[i][k];
    }
    rank++;
  }
  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];
  }
  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 SolveLinear, make the following changes:

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

**SolveLinearBinary.h**

**Description:** Solves  $Ax = b$  over  $\mathbb{F}_2$ . If there are multiple solutions, one is returned arbitrarily. Returns rank, or -1 if no solutions. Destroys  $A$  and  $b$ .

**Time:**  $\mathcal{O}(n^2 m)$

fa2d7a, 34 lines

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

```
int solveLinear(vector<bs>& A, vi& b, bs& x, int m) {
  int n = sz(A), rank = 0, br;
  assert(m <= sz(x));
  vi col(m); iota(all(col), 0);
  rep(i,0,n) {
    for (br=i; br<n; ++br) if (A[br].any()) break;
    if (br == n) {
      rep(j,i,n) if (b[j]) return -1;
      break;
    }
    int bc = (int)A[br]._Find_next(i-1);
    swap(A[i], A[br]);
    swap(b[i], b[br]);
    swap(col[i], col[bc]);
    rep(j,0,n) if (A[j][i] != A[j][bc]) {
      A[j].flip(i); A[j].flip(bc);
    }
    rep(j,i+1,n) if (A[j][i]) {
      b[j] ^= b[i];
      A[j] ^= 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 ( $\text{rank} < n$ ). Can easily be extended to prime moduli; for prime powers, repeatedly set  $A^{-1} = A^{-1}(2I - AA^{-1}) \pmod{p^k}$  where  $A^{-1}$  starts as the inverse of  $A$  mod  $p$ , and  $k$  is doubled in each step.

**Time:**  $\mathcal{O}(n^3)$

ebfff6, 35 lines

```
int matInv(vector<vector<double>>& A) {
  int n = sz(A); vi col(n);
  vector<vector<double>> tmp(n, vector<double>(n));
  rep(i,0,n) tmp[i][i] = 1, col[i] = i;
  rep(i,0,n) {
    int r = i, c = i;
```

```

rep(j,i,n) rep(k,i,n)
  if (fabs(A[j][k]) > fabs(A[r][c]))
    r = j, c = k;
  if (fabs(A[r][c]) < 1e-12) return i;
A[i].swap(A[r]); tmp[i].swap(tmp[r]);
rep(j,0,n)
  swap(A[j][i], A[j][c]), swap(tmp[j][i], tmp[j][c]);
swap(col[i], col[c]);
double v = A[i][i];
rep(j,i+1,n) {
  double f = A[j][i] / v;
  A[j][i] = 0;
  rep(k,i+1,n) A[j][k] -= f*A[i][k];
  rep(k,0,n) tmp[j][k] -= f*tmp[i][k];
}
rep(j,i+1,n) A[i][j] /= v;
rep(j,0,n) tmp[i][j] /= v;
A[i][i] = 1;
}

for (int i = n-1; i > 0; --i) rep(j,0,i) {
  double v = A[j][i];
  rep(k,0,n) tmp[j][k] -= v*tmp[i][k];
}

rep(i,0,n) rep(j,0,n) A[col[i]][col[j]] = tmp[i][j];
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:**  $\mathcal{O}(N^3)$

[.../number-theory/ModPow.h](#) 0b7b13, 37 lines

```

int matInv(vector<vector<ll>>& 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,1,n) rep(k,i,n) if (A[j][k]) {
      r = j; c = k; goto found;
    }
    return i;
  }
  found:
  A[i].swap(A[r]); tmp[i].swap(tmp[r]);
  rep(j,0,n)
    swap(A[j][i], A[j][c]), swap(tmp[j][i], tmp[j][c]);
  swap(col[i], col[c]);
  ll v = modpow(A[i][i], mod - 2);
  rep(j,i+1,n) {
    ll f = A[j][i] * v % mod;
    A[j][i] = 0;
    rep(k,i+1,n) A[j][k] = (A[j][k] - f*A[i][k]) % mod;
    rep(k,0,n) tmp[j][k] = (tmp[j][k] - f*tmp[i][k]) % mod;
  }
  rep(j,i+1,n) A[i][j] = A[i][j] * v % mod;
  rep(j,0,n) tmp[i][j] = tmp[i][j] * v % mod;
  A[i][i] = 1;

  for (int i = n-1; i > 0; --i) rep(j,0,i) {
    ll v = A[j][i];
    rep(k,0,n) tmp[j][k] = (tmp[j][k] - v*tmp[i][k]) % mod;
  }
}

```

```

rep(i,0,n) rep(j,0,n)
  A[col[i]][col[j]] = tmp[i][j] % mod + (tmp[i][j] < 0)*mod;
return n;
}

```

## Tridiagonal.h

**Description:**  $x = \text{tridiagonal}(d, p, q, b)$  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

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

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

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

Fails if the solution is not unique.

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

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

[8f9fa8, 26 lines](#)

```

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

```

## 4.3 Fourier transforms

### FastFourierTransform.h

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

**Time:**  $\mathcal{O}(N \log N)$  with  $N = |A| + |B|$  ( $\sim 1s$  for  $N = 2^{22}$ )

```

typedef complex<double> C;
typedef vector<double> vd;
void fft(vector<C>& a) {

```

```

int n = sz(a), L = 31 - __builtin_clz(n);
static vector<complex<long double>> R(2, 1);
static vector<C> rt(2, 1); // (^ 10% faster if double)
for (static int k = 2; k < n; k *= 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) {
    C z = rt[j+k] * a[i+j+k]; // (25% faster if hand-rolled)
    a[i + j + k] = a[i + j] - z;
    a[i + j] += z;
  }
}
vd conv(const vd& a, const vd& b) {
  if (a.empty() || b.empty()) return {};
  vd res(sz(a) + sz(b) - 1);
  int L = 32 - __builtin_clz(sz(res)), n = 1 << L;
  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]);
  fft(out);
  rep(i,0,sz(res)) res[i] = imag(out[i]) / (4 * n);
  return res;
}

```

### FastFourierTransformMod.h

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

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

["FastFourierTransform.h"](#) b82773, 22 lines

```

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

```

### NumberTheoreticTransform.h

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

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

```
.../number-theory/ModPow.h" ced03d, 35 lines
const ll mod = (119 << 23) + 1, root = 62; // = 998244353
// For p < 2^30 there is also e.g. 5 << 25, 7 << 26, 479 << 21
// and 483 << 21 (same root). The last two are > 10^9.
typedef vector<ll> vl;
void ntt(vl &a) {
    int n = sz(a), L = 31 - __builtin_clz(n);
    static vl rt(2, 1);
    for (static int k = 2, s = 2; k < n; k *= 2, s++) {
        rt.resize(n);
        ll z[] = {1, modpow(root, mod >> s)};
        rep(i, k, 2*k) rt[i] = rt[i / 2] * z[i & 1] % mod;
    }
    vi rev(n);
    rep(i, 0, n) rev[i] = (rev[i / 2] | (i & 1) << L) / 2;
    rep(i, 0, n) if (i < rev[i]) swap(a[i], a[rev[i]]);
    for (int k = 1; k < n; k *= 2)
        for (int i = 0; i < n; i += 2 * k) rep(j, 0, k) {
            ll z = rt[j + k] * a[i + j + k] % mod, &ai = a[i + j];
            a[i + j + k] = ai - z + (z > ai ? mod : 0);
            ai += (ai + z >= mod ? z - mod : z);
        }
    }
    vl conv(const vl &a, const vl &b) {
        if (a.empty() || b.empty()) return {};
        int s = sz(a) + sz(b) - 1, B = 32 - __builtin_clz(s),
            n = 1 << B;
        int inv = modpow(n, mod - 2);
        vl L(a), R(b), out(n);
        L.resize(n), R.resize(n);
        ntt(L), ntt(R);
        rep(i, 0, n)
            out[-i & (n - 1)] = (ll)L[i] * R[i] % mod * inv % mod;
        ntt(out);
        return {out.begin(), out.begin() + s};
    }
}
```

## Number theory (5)

### 5.1 Modular arithmetic

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

```
"euclid.h" 35bfea, 18 lines
const ll mod = 17; // change to something else
struct Mod {
    ll x;
    Mod(ll xx) : x(xx) {}
    Mod operator+(Mod b) { return Mod((x + b.x) % mod); }
    Mod operator-(Mod b) { return Mod((x - b.x + mod) % mod); }
    Mod operator*(Mod b) { return Mod((x * b.x) % mod); }
    Mod operator/(Mod b) { return *this * invert(b); }
    Mod invert(Mod a) {
        ll x, y, g = euclid(a.x, mod, x, y);
        assert(g == 1); return Mod((x + mod) % mod);
    }
    Mod operator^(ll e) {
        if (!e) return Mod(1);
        ...
```

```
        Mod r = *this ^ (e / 2); r = r * r;
        return e & 1 ? *this * r : r;
    };
}
```

### ModInverse.h

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

6f684f, 3 lines

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

### ModPow.h

b83e45, 8 lines

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

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

### ModLog.h

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

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

c040b8, 11 lines

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

### ModSum.h

**Description:** Sums of mod'ed arithmetic progressions.

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

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

5c5bc5, 16 lines

```
typedef unsigned long long ull;
ull sumsq(ull to) { return to / 2 * ((to-1) | 1); }

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

ull 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  $a \cdot b \pmod{c}$  (or  $a^b \pmod{c}$ ) for  $0 \leq a, b \leq 7.2 \cdot 10^{18}$ .

**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;
}
```

### ModSqrt.h

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

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

"ModPow.h" 19a793, 24 lines

```
ll sqrt(ll a, ll p) {
```

```
    a %= p; if (a < 0) a += p;
    if (a == 0) return 0;
    assert(modpow(a, (p-1)/2, p) == 1); // else no solution
    if (p % 4 == 3) return modpow(a, (p+1)/4, p);
    // a^(n+3)/8 or 2^(n+3)/8 * 2^(n-1)/4 works if p % 8 == 5
    ll s = p - 1, n = 2;
```

```
    int r = 0, m;
    while (s % 2 == 0)
        ++r, s /= 2;
    while (modpow(n, (p - 1) / 2, p) != p - 1) ++n;
    ll x = modpow(a, (s + 1) / 2, p);
    ll b = modpow(a, s, p), g = modpow(n, s, p);
    for (; r = m) {
        ll t = b;
        for (m = 0; m < r && t != 1; ++m)
            t = t * t % p;
        if (m == 0) return x;
        ll gs = modpow(g, 1LL << (r - m - 1), p);
        g = gs * gs % p;
        x = x * gs % p;
        b = b * g % p;
    }
}
```

## 5.2 Primality

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

**Time:**  $\text{LIM} = 1e9 \approx 1.5s$

6b2912, 20 lines

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

## 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  $a^b \pmod{c}$ .

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

"ModMullL.h", "MillerRabin.h" d8d98d, 18 lines

```
ull pollard(ull n) {
    ull x = 0, y = 0, t = 30, prd = 2, i = 1, q;
    auto f = [&](ull x) { return modmul(x, x, n) + i; };
    while (t++ % 40 || __gcd(prd, n) == 1) {
        if (x == y) x = ++i, y = f(x);
        if ((q = modmul(prd, max(x,y) - min(x,y), n))) prd = q;
        x = f(x), y = f(f(y));
    }
    return __gcd(prd, n);
}

vector<ull> factor(ull n) {
    if (n == 1) return {};
    if (isPrime(n)) return {n};
    ull x = pollard(n);
    auto l = factor(x), r = factor(n / x);
    l.insert(l.end(), all(r));
    return l;
}
```

## 5.3 Divisibility

## euclid.h

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

33ba8f, 5 lines

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

## CRT.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  $0 \leq x < \text{lcm}(m, n)$ . Assumes  $mn < 2^{62}$ .

**Time:**  $\log(n)$

"euclid.h" 04d93a, 7 lines

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

}

## 5.3.1 Bézout's identity

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

$$ax + by = d$$

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

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

## phiFunction.h

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

$$\sum_{d|n} \phi(d) = n, \sum_{1 \leq k \leq n, \gcd(k,n)=1} k = n\phi(n)/2, n > 1$$

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

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

cf7d6d, 8 lines

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

## 5.4 Fractions

## ContinuedFractions.h

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

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

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

dd6c5e, 21 lines

```
typedef double d; // for N ~ 1e7; long double for N ~ 1e9
pair<11, 11> approximate(d x, 11 N) {
    11 LP = 0, LQ = 1, P = 1, Q = 0, inf = LLONG_MAX; d y = x;
    for (;;) {
        11 lim = min(P ? (N-LP) / P : inf, Q ? (N-LQ) / Q : inf),
            a = (11)floor(y), b = min(a, lim),
            NP = b*P + LP, NQ = b*Q + LQ;
        if (a > b) {
            // If b > a/2, we have a semi-convergent that gives us a
            // better approximation; if b = a/2, we *may* have one.
            // Return {P, Q} here for a more canonical approximation.
            return (abs(x - (d)NP / (d)NQ) < abs(x - (d)P / (d)Q)) ?
                make_pair(NP, NQ) : make_pair(P, Q);
        }
        if (abs(y = 1/(y - (d)a)) > 3*N) {
            return {NP, NQ};
        }
        LP = P; P = NP;
        LQ = Q; Q = NQ;
    }
}
```

## FracBinarySearch.h

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

**Usage:** `fracBS([](Frac f) { return f.p>=3*f.q; }, 10); // {1,3}`

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

struct Frac { 11 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;
}
```

## 5.5 Pythagorean Triples

The Pythagorean triples are uniquely generated by

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

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

## 5.6 Primes

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

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

## 5.7 Estimates

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

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

## 5.8 Möbius Function

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

Möbius Inversion:

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

# FPT

Other useful formulas/forms:

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

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

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

## Combinatorial (6)

### 6.1 Permutations

#### 6.1.1 Factorial

$n$	1	2	3	4	5	6	7	8	9	10
$n!$	1	2	6	24	120	720	5040	40320	362880	3628800
$n$		11	12	13	14	15	16	17		
$n!$	4.0e7	4.8e8	6.2e9	8.7e10	1.3e12	2.1e13	3.6e14			
$n$	20	25	30	40	50	100	150		171	
$n!$	2e18	2e25	3e32	8e47	3e64	9e157	6e262	>DBL_MAX		

#### IntPerm.h

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

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

044568, 6 lines

```
int permToInt(vi& v) {
    int use = 0, i = 0, r = 0;
    for(int x:v) r = r * ++i + __builtin_popcount(use & -(1<<x)),
        use |= 1 << x; // (note: minus, not ~!)
    return r;
}
```

#### 6.1.2 Cycles

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

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

#### 6.1.3 Derangements

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

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

#### 6.1.4 Burnside's lemma

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

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

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

## IntPerm multinomial

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

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

### 6.2 Partitions and subsets

#### 6.2.1 Partition function

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

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

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

$n$	0	1	2	3	4	5	6	7	8	9	20	50	100
$p(n)$	1	1	2	3	5	7	11	15	22	30	627	~2e5	~2e8

#### 6.2.2 Lucas' Theorem

Let  $n, m$  be non-negative integers and  $p$  a prime. Write  $n = n_k p^k + \dots + n_1 p + n_0$  and  $m = m_k p^k + \dots + m_1 p + m_0$ . Then  $\binom{n}{m} \equiv \prod_{i=0}^k \binom{n_i}{m_i} \pmod{p}$ .

#### 6.2.3 Binomials

multinomial.h

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

```
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;
}
```

### 6.3 General purpose numbers

#### 6.3.1 Bernoulli numbers

EGF of Bernoulli numbers is  $B(t) = \frac{t}{e^t - 1}$  (FFT-able).

$$B[0, \dots] = [1, -\frac{1}{2}, \frac{1}{6}, 0, -\frac{1}{30}, 0, \frac{1}{42}, \dots]$$

Sums of powers:

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

Euler-Maclaurin formula for infinite sums:

$$\begin{aligned} \sum_{i=m}^{\infty} f(i) &= \int_m^{\infty} f(x) dx - \sum_{k=1}^{\infty} \frac{B_k}{k!} f^{(k-1)}(m) \\ &\approx \int_m^{\infty} f(x) dx + \frac{f(m)}{2} - \frac{f'(m)}{12} + \frac{f''(m)}{720} + O(f^{(5)}(m)) \end{aligned}$$

#### 6.3.2 Stirling numbers of the first kind

Number of permutations on  $n$  items with  $k$  cycles.

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

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

$$c(8, k) = 8, 0, 5040, 13068, 13132, 6769, 1960, 322, 28, 1$$

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

#### 6.3.3 Eulerian numbers

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

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

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

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

#### 6.3.4 Stirling numbers of the second kind

Partitions of  $n$  distinct elements into exactly  $k$  groups.

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

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

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

#### 6.3.5 Bell numbers

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

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

#### 6.3.6 Labeled unrooted trees

# on  $n$  vertices:  $n^{n-2}$

# on  $k$  existing trees of size  $n_i$ :  $n_1 n_2 \dots n_k n^{k-2}$

# with degrees  $d_i$ :  $(n-2)! / ((d_1-1)! \dots (d_{n-1}-1)!)$

#### 6.3.7 Catalan numbers

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

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

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

- sub-diagonal monotone paths in an  $n \times n$  grid.
- strings with  $n$  pairs of parenthesis, correctly nested.
- binary trees with  $n+1$  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.
- permutations of  $[n]$  with no 3-term increasing subseq.

## Games Theory (7)

### Grundy.h

**Description:** Calculates Grundy numbers for impartial games.

**Usage:** GrundyCalculator gc(n, m);  
gc.add\_move(a, b); // Add move from position a to b  
gc.compute\_grundy(); // Calculate all Grundy numbers

**Time:**  $\mathcal{O}(N + M)$

4e0230, 25 lines

### struct GrundyCalculator {

```
int N, M;
vector<vector<int>> moves;
vector<int> grundy;

GrundyCalculator(int n, int m) : N(n), M(m),
    moves(n+1), grundy(n+1) {}

void add_move(int a, int b) { moves[a].push_back(b); }

int mex(const vector<int> &s) {
    vector<bool> present(s.size() + 1);
    for(int x : s) if(x < sz(present)) present[x] = true;
    rep(i, 0, sz(present)) if(!present[i]) return i;
    return sz(present);
}

void compute_grundy() {
    rep(i, 0, N+1) {
        vector<int> next;
        for(int b : moves[i]) next.push_back(grundy[b]);
        grundy[i] = mex(next);
    }
}
```

### GrundyF.h

**Description:** Functional implementation of Grundy numbers calculation

**Usage:** grundy(n, S) returns grundy number for position n with subtraction set S

**Time:**  $\mathcal{O}(N * |S|)$  where |S| is size of subtraction set

7c066e, 8 lines

```
int grundy(int n, vector<int>& S) {
    if (n == 0) return 0;
    unordered_set<int> s;
    for (int x : S)
        if (n >= x) s.insert(grundy(n - x, S));
    for (int i = 0;; i++)
        if (!s.count(i)) return i;
}
```

### TurningTurtles.h

**Description:** Grundy numbers for Turning Turtles game A turtle can be flipped to change its direction

**Usage:** grundy(n) returns grundy number for n turtles in a row

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

4f9cb1, 4 lines

```
int grundy(int n) {
    if (n == 0) return 0;
    return n % 2 == 0 ? n / 2 : grundy(n / 2);
}
```

### GreenHackenbush.h

**Description:** Grundy numbers for Green Hackenbush on trees

**Usage:** grundy(root) returns grundy number for tree rooted at root

**Time:**  $\mathcal{O}(N)$  where N is number of nodes

51bc65, 10 lines

```
struct TreeNode {
    vector<TreeNode*> children;
```

```
};

int grundy(TreeNode* node) {
    int x = 0;
    for (auto child : node->children)
        x ^= (1 + grundy(child));
    return x;
}
```

### ChompGame.h

**Description:** Grundy numbers for Chomp game on a rectangle

**Usage:** grundy(a, b) returns grundy number for axb chocolate bar

**Time:**  $\mathcal{O}(A * B)$  with memoization

714eeb, 11 lines

```
map<pair<int, int>, int> memo;
int grundy(int a, int b) {
    if (a == 0 || b == 0) return 0;
    if (memo.count({a, b})) return memo[{a, b}];
    unordered_set<int> s;
    rep(i, 1, a+1) rep(j, 1, b+1)
        s.insert(grundy(i - 1, j - 1));
    rep(i, 0, INT_MAX) if (!s.count(i))
        return memo[{a, b}] = i;
    return 0;
}
```

### KaylesGame.h

**Description:** Grundy numbers for Kayles game on a binary string

**Usage:** grundy(s) returns grundy number for position represented by string s

**Time:**  $\mathcal{O}(N * 2^N)$  where N is string length

588282, 18 lines

```
map<string, int> memo;
int grundy(string s) {
    if (memo.count(s)) return memo[s];
    unordered_set<int> sgs;
    rep(i, 0, sz(s)) {
        if (s[i] == '1') {
            s[i] = '0';
            sgs.insert(grundy(s));
            if (i + 1 < sz(s) && s[i + 1] == '1') {
                s[i + 1] = '0';
                sgs.insert(grundy(s));
                s[i + 1] = '1';
            }
            s[i] = '1';
        }
        for (int i = 0;; i++) if (!sgs.count(i)) return memo[s] = i;
    }
}
```

### TakeAwayGamewithPrimeMoves.h

**Description:** Grundy numbers for Take Away game with prime moves Can only take prime number of stones

**Usage:** grundy(n) returns grundy number for n stones

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

5101fd, 15 lines

```
bool is_prime(int n) {
    if (n < 2) return false;
    for (int i = 2; i * i <= n; i++)
        if (n % i == 0) return false;
    return true;
}

int grundy(int n) {
    if (n == 0) return 0;
    unordered_set<int> s;
    for (int i = 2; i <= n; i++)
```

```
    if (is_prime(i)) s.insert(grundy(n - i));
    rep(i, 0, INT_MAX) if (!s.count(i)) return i;
    return 0;
}
```

### SubtractionGames.h

**Description:** Grundy numbers for subtraction games

**Usage:** grundy(n, S) returns grundy number for position n with subtraction set S

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

7c066e, 8 lines

```
int grundy(int n, vector<int>& S) {
    if (n == 0) return 0;
    unordered_set<int> s;
    for (int x : S)
        if (n >= x) s.insert(grundy(n - x, S));
    for (int i = 0;; i++)
        if (!s.count(i)) return i;
}
```

### Cram.h

**Description:** Grundy numbers for Cram game on wxh board

**Usage:** grundy(w, h) returns grundy number for board of width w and height h

**Time:**  $\mathcal{O}(1)$

757141, 3 lines

```
int grundy(int w, int h) {
    return (w * h) % 2;
}
```

## Dynamic Programming (8)

### DivideAndConquerDP.h

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

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

d38d2b, 18 lines

```
struct DP { // Modify at will:
    int lo(int ind) { return 0; }
    int hi(int ind) { return ind; }
    ll 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) / 2;
        pair<ll, 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); }
};
```

### KnuthDP.h

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

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

## LIS.h

**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<p> 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  $\leq t$  such that S is the sum of some subset of the weights.

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

b20ccc, 16 lines

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

## mobiusDP.h

**Description:** DP with Möbius Transform over subsets.

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

9254e2, 7 lines

```
const int N = 20;
ll f[1<<N];

void mobius_transform() {
    rep(i, N) rep(mask, 1<<N) if(mask&(1<<i))
        f[mask] -= f[mask^(1<<i)];
}
```

## NTTDP.h

**Description:** DP with convolution optimized using Number Theoretic Transform (NTT).

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

ff8776, 35 lines

```
const int mod=998244353, root=3;
void ntt(vector<int>& a, bool inv) {
    int na=a.size(), j=0;
    for(int i=1;i<n;++i) {
        int bit=n>>1;
        for(;j&bit;bit>>1) j^=bit;
        j^=bit; if(i<j) swap(a[i],a[j]);
    }
}
```

```
}
for(int len=2;len<=n;len<<=1) {
    int wlen=powmod(root, (mod-1)/len);
    if(inv) wlen=powmod(wlen, mod-2);
    for(int i=0;i<n;i+=len) {
        int w=1;
        for(int j=0;j<len/2;j++) {
            int u=a[i+j], v=(int)(1LL*a[i+j+len/2]*w%mod);
            a[i+j]=u+v<mod?u+v:u+v-mod;
            a[i+j+len/2]=u-v>=0?u-v:u-v+mod;
            w=(int)(1LL*w*wlen%mod);
        }
    }
    if(inv) {
        int nrev=powmod(n, mod-2);
        for(int x:a) x=(int)(1LL*x*nrev%mod);
    }
}

void dp_ntt(vector<int>& a, vector<int>& b) {
    int n=1;
    while(n<a.size()+b.size()) n<<=1;
    a.resize(n); b.resize(n);
    ntt(a, false); ntt(b, false);
    rep(i, n) a[i]=(int)(1LL*a[i]*b[i]%mod);
    ntt(a, true);
}
```

## palindromeDP.h

**Description:** DP for counting palindromic substrings.

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

59f2e9, 10 lines

```
const int N = 5000;
char s[N];
bool p[N][N];

void pal_dp() {
    int n=strlen(s);
    repd(i, n) rep(j, i, n) {
        p[i][j]=(s[i]==s[j])&&(j-i<2||p[i+1][j-1]);
    }
}
```

## SOSDP.h

**Description:** Sum over Subsets DP (SOS DP).

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

49795d, 7 lines

```
const int N = 20;
ll f[1<<N];

void sos_dp() {
    rep(i, N) rep(mask, 1<<N) if(mask&(1<<i))
        f[mask]+=f[mask^(1<<i)];
}
```

## bitmaskDP.h

**Description:** TSP using DP with bitmasking.

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

4527c7, 20 lines

```
const int N = 20, INF = 1e9;
int n, d[N][N], dp[1<<N][N];

void tsp() {
    rep(m, 1<<n) rep(u, n) dp[m][u]=INF;
    dp[1][0]=0;
    rep(m, 1<<n) rep(u, n) if(m&(1<<u))
        rep(v, n) if(! (m&(1<<v)))
            dp[m][v]=min(dp[m](1<<v)][v], dp[m][u]+d[u][v]);
}

ll dp[1<<N];
void bitmask_dp() {
    int n; dp[0]=base_case;
    rep(mask, 1<<n) {
        // process dp[mask]
        rep(i, n) if(!(mask&(1<<i)))
            dp[mask|(1<<i)]=update(dp[mask|(1<<i)], dp[mask], i);
    }
}

```

## flowerDP.h

**Description:** Top-down DP for flower arrangement.

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

0ff6f4, 17 lines

```
const int N = 1e3, K = 1e3, INF = 1e9;
```

```
int A[K+1][N+1];
```

```
int memo[K+1][N+1];
```

```
bool vis[K+1][N+1];
```

```
int dp(int i, int j) {
```

```
    if (i == 0) return 0;
```

```
    if (j < i) return -INF;
```

```
    if (vis[i][j]) return memo[i][j];
```

```
    vis[i][j] = true;
```

```
    return memo[i][j] = max(dp(i, j - 1), dp(i - 1, j - 1) + A[i][j]);
```

```
int maxAesthetic(int n, int k) {
```

```
    memset(vis, 0, sizeof(vis));
```

```
    return dp(k, n);
```

## SubsetSumDP.h

**Description:** Checks if a subset sum equals S.

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

2937e7, 10 lines

```
const int N = 1000, S = 1e5;
```

```
int a[N];
```

```
bitset<S+1> dp;
```

```
bool subsetSum(int n, int sum) {
```

```
    dp.reset(); dp[0] = 1;
```

```
    for (int i = 0; i < n; ++i)
```

```
        dp |= dp << a[i];
```

```
    return dp[sum];
```

```
}
```

## TargetSumDP.h

**Description:** Determines if target sum S can be achieved using + and -.

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

51ecff, 15 lines

```
const int N = 1000, T = 1e4;
```

```
int a[N];
```

```
bitset<2*T+1> dp;
```

```
bool targetSum(int n, int S) {
```

```
    const int offset = T;
    dp.reset(); dp[offset] = 1;
```

```
    for (int i = 0; i < n; ++i)
```

```
        bitset<2*T+1> next;
```

```
        next |= dp << a[i];
```

```
        next |= dp >> a[i];
```

```
        dp = next;
```

```
}
```

```
return dp[S + offset];
```

}

## PartitionDP.h

**Description:** Minimizes difference between two subsets.**Time:**  $\mathcal{O}(N * T)$ 

cb63d6, 14 lines

```
const int N = 1000, T = 1e5;
int a[N];
bitset<T+1> dp;
```

```
int minDifference(int n) {
    int total = accumulate(a, a + n, 0);
    dp.reset(); dp[0] = 1;
    for (int i = 0; i < n; ++i)
        dp |= dp << a[i];
    int half = total / 2;
    for (int s = half; s >= 0; --s)
        if (dp[s]) return total - 2 * s;
    return total;
}
```

## stringDP.h

**Description:** Computes edit distance between A and B.**Time:**  $\mathcal{O}(M * N)$ 

ed07be, 13 lines

```
const int N = 1e3;
char A[N], B[N];
```

```
int editDistance(int m, int n) {
    vector<vector<int>> dp(m+1, vector<int>(n+1));
    rep(i, m+1) dp[i][0] = i;
    rep(j, n+1) dp[0][j] = j;
    rep(i, 1, m+1) rep(j, 1, n+1) {
        if (A[i-1] == B[j-1]) dp[i][j] = dp[i-1][j-1];
        else dp[i][j] = 1 + min({dp[i-1][j], dp[i][j-1], dp[i-1][j-1]});
    }
    return dp[m][n];
}
```

## LCSDP.h

**Description:** Computes the length of LCS of A and B.**Time:**  $\mathcal{O}(M * N)$ 

027631, 11 lines

```
const int N = 1e3;
char A[N], B[N];
```

```
int LCS(int m, int n) {
    vector<vector<int>> dp(m+1, vector<int>(n+1));
    rep(i, 1, m+1) rep(j, 1, n+1) {
        if (A[i-1] == B[j-1]) dp[i][j] = dp[i-1][j-1] + 1;
        else dp[i][j] = max(dp[i-1][j], dp[i][j-1]);
    }
    return dp[m][n];
}
```

## CoinDP.h

**Description:** Minimum coins to make amount S.**Time:**  $\mathcal{O}(N * S)$ 

4efa3b, 11 lines

```
const int N = 100, S = 1e5, INF = 1e9;
int coins[N];

int coinChange(int n, int amount) {
    vector<int> dp(amount+1, INF);
    dp[0] = 0;
    for (int i = 0; i < n; ++i)
        for (int s = coins[i]; s <= amount; ++s)
            dp[s] = min(dp[s], dp[s - coins[i]] + 1);
```

```
return dp[amount] == INF ? -1 : dp[amount];
}
```

## newratingDP.h

**Description:** Computes the maximum possible rating after skipping an optimal interval.**Time:**  $\mathcal{O}(N)$  per test case

e6915d, 20 lines

```
void newRating() {
    int t; cin >> t;
    while (t--) {
        int n; cin >> n;
        vector<int> a(n), s(n);
        for (int i = 0; i < n; ++i) cin >> a[i];

        int x = 0, x_total = 0;
        for (int i = 0; i < n; ++i) {
            if (a[i] > x) { s[i] = 1; x += 1; }
            else if (a[i] == x) { s[i] = 0; }
            else { s[i] = -1; x -= 1; }
        }
        x_total = x;

        int min_sum = INT_MAX, curr_sum = 0;
        for (int i = 0; i < n; ++i) {
            curr_sum += s[i];
            if (curr_sum > 0) curr_sum = s[i];
            min_sum = min(min_sum, curr_sum);
        }
        if (min_sum == INT_MAX) min_sum = *min_element(s.begin(), s.end());
        int max_rating = x_total - min_sum;
        cout << max_rating << '\n';
    }
}
```

## Graph (9)

## 9.1 Fundamentals

## BellmanFord.h

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

830a8f, 23 lines

```
const ll inf = LLONG_MAX;
struct Ed { int a, b, w, s() { return a < b ? a : -a; } };
struct Node { ll dist = inf; int prev = -1; };

void bellmanFord(vector<Node>& nodes, vector<Ed>& eds, int s) {
    nodes[s].dist = 0;
    sort(all(eds), [] (Ed a, Ed b) { return a.s() < b.s(); });

    int lim = sz(nodes) / 2 + 2; // /3+100 with shuffled vertices
    rep(i, 0, lim) for (Ed ed : eds) {
        Node cur = nodes[ed.a], &dest = nodes[ed.b];
        if (abs(cur.dist) == inf) continue;
        ll d = cur.dist + ed.w;
        if (d < dest.dist) {
            dest.prev = ed.a;
            dest.dist = (i < lim-1 ? d : -inf);
        }
    }
    rep(i, 0, lim) for (Ed e : eds) {
        if (nodes[e.a].dist == -inf)
            nodes[e.b].dist = -inf;
    }
}
```

}

## FloydWarshall.h

**Description:** Calculates all-pairs shortest path in a directed graph that might have negative edge weights. Input is an distance matrix  $m$ , where  $m[i][j] = \text{inf}$  if  $i$  and  $j$  are not adjacent. As output,  $m[i][j]$  is set to the shortest distance between  $i$  and  $j$ ,  $\text{inf}$  if no path, or  $-\text{inf}$  if the path goes through a negative-weight cycle.**Time:**  $\mathcal{O}(N^3)$ 

531245, 12 lines

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

## TopoSort.h

**Description:** Topological sorting. Given is an oriented graph. Output is an ordering of vertices, such that there are edges only from left to right. If there are cycles, the returned list will have size smaller than  $n - \text{nodes reachable from cycles}$  will not be returned.**Time:**  $\mathcal{O}(|V| + |E|)$ 

d678d8, 8 lines

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

## 9.2 Network flow

## PushRelabel.h

**Description:** Push-relabel using the highest label selection rule and the gap heuristic. Quite fast in practice. To obtain the actual flow, look at positive values only.**Time:**  $\mathcal{O}(V^2\sqrt{E})$ 

0ae1d4, 48 lines

```
struct PushRelabel {
    struct Edge {
        int dest, back;
        ll f, c;
    };
    vector<vector<Edge>> g;
    vector<ll> ec;
    vector<Edge*> cur;
    vector<vi> hs; vi H;
    PushRelabel(int n) : g(n), ec(n), cur(n), hs(2*n), H(n) {}

    void addEdge(int s, int t, ll cap, ll rcap=0) {
        if (s == t) return;
        g[s].push_back({t, sz(g[t]), 0, cap});
        g[t].push_back({s, sz(g[s])-1, 0, rcap});
    }

    void addFlow(Edge& e, ll f) {
        Edge &back = g[e.dest][e.back];
        if (!ec[e.dest] && f) hs[H[e.dest]].push_back(e.dest);
        e.f += f; e.c -= f; ec[e.dest] += f;
        back.f -= f; back.c += f; ec[back.dest] -= f;
    }
}
```

```

    }
ll calc(int s, int t) {
    int v = sz(g); H[s] = v; ec[t] = 1;
    vi co(2*v); co[0] = v-1;
    rep(i,0,v) cur[i] = g[i].data();
    for (Edge& e : g[s]) addFlow(e, e.c);

    for (int hi = 0;;) {
        while (hs[hi].empty()) if (!hi--) return -ec[s];
        int u = hs[hi].back(); hs[hi].pop_back();
        while (ec[u] > 0) // discharge u
            if (cur[u] == g[u].data() + sz(g[u])) {
                H[u] = 1e9;
                for (Edge& e : g[u]) if (e.c && H[u] > H[e.dest]+1)
                    H[u] = H[e.dest]+1, cur[u] = &e;
                if (++co[H[u]], !--co[hi] && hi < v)
                    rep(i,0,v) if (hi < H[i] && H[i] < v)
                        --co[H[i]], H[i] = v + 1;
                hi = H[u];
            } else if (cur[u]->c && H[u] == H[cur[u]->dest]+1)
                addFlow(*cur[u], min(ec[u], cur[u]->c));
            else ++cur[u];
    }
    bool leftOfMinCut(int a) { return H[a] >= sz(g); }
};

MinCostMaxFlow.h

```

**Description:** Min-cost max-flow. If costs can be negative, call setpi before maxflow, but note that negative cost cycles are not supported. To obtain the actual flow, look at positive values only.

**Time:**  $O(FE \log(V))$  where F is max flow.  $O(VE)$  for setpi. 58385b, 79 lines

```
#include <bits/extc++.h>
```

```

const ll INF = numeric_limits<ll>::max() / 4;

struct MCMF {
    struct edge {
        int from, to, rev;
        ll cap, cost, flow;
    };
    int N;
    vector<vector<edge>> ed;
    vi seen;
    vector<ll> dist, pi;
    vector<edge*> par;

MCMF(int N) : N(N), ed(N), seen(N), dist(N), pi(N), par(N) {}

void addEdge(int from, int to, ll cap, ll cost) {
    if (from == to) return;
    ed[from].push_back(edge{ from,to,sz(ed[to]),cap,cost,0 });
    ed[to].push_back(edge{ to,from,sz(ed[from])-1,0,-cost,0 });
}

void path(int s) {
    fill(all(seen), 0);
    fill(all(dist), INF);
    dist[s] = 0; ll di;

__gnu_pbds::priority_queue<pair<ll, int>> q;
vector<decltype(q)::point_iterator> its(N);
q.push({ 0, s });

while (!q.empty()) {
    s = q.top().second; q.pop();
    seen[s] = 1; di = dist[s] + pi[s];
    for (edge& e : ed[s]) if (!seen[e.to]) {

```

## MinCostMaxFlow EdmondsKarp Dinic

```

        ll val = di - pi[e.to] + e.cost;
        if (e.cap - e.flow > 0 && val < dist[e.to]) {
            dist[e.to] = val;
            par[e.to] = &e;
            if (its[e.to] == q.end())
                its[e.to] = q.push({ -dist[e.to], e.to });
            else
                q.modify(its[e.to], { -dist[e.to], e.to });
        }
    }
    rep(i,0,N) pi[i] = min(pi[i] + dist[i], INF);
}

pair<ll, ll> maxflow(int s, int t) {
    ll totflow = 0, totcost = 0;
    while (path(s), seen[t]) {
        ll fl = INF;
        for (edge* x = par[t]; x; x = par[x->from])
            fl = min(fl, x->cap - x->flow);

        totflow += fl;
        for (edge* x = par[t]; x; x = par[x->from]) {
            x->flow += fl;
            ed[x->to][x->rev].flow -= fl;
        }
    }
    rep(i,0,N) for (edge& e : ed[i]) totcost += e.cost * e.flow;
    return {totflow, totcost/2};
}

```

// If some costs can be negative, call this before maxflow:

```
void setpi(int s) { // (otherwise, leave this out)
    fill(all(pi), INF); pi[s] = 0;
    int it = N, ch = 1, ll v;
    while (ch-- && it--)
        rep(i,0,N) if (pi[i] != INF)
            for (edge& e : ed[i]) if (e.cap)
                if ((v = pi[i] + e.cost) < pi[e.to])
                    pi[e.to] = v, ch = 1;
    assert(it >= 0); // negative cost cycle
}
```

## EdmondsKarp.h

**Description:** Flow algorithm with guaranteed complexity  $O(VE^2)$ . To get edge flow values, compare capacities before and after, and take the positive values only. 482fe0, 36 lines

```
template<class T> T edmondsKarp(vector<unordered_map<int, T>>& graph, int source, int sink) {
    assert(source != sink);
    T flow = 0;
    vi par(sz(graph)), q = par;

    for (;;) {
        fill(all(par), -1);
        par[source] = 0;
        int ptr = 1;
        q[0] = source;

        rep(i,0,ptr) {
            int x = q[i];
            for (auto e : graph[x]) {
                if (par[e.first] == -1 && e.second > 0) {
                    par[e.first] = x;
                    q[ptr++] = e.first;
                    if (e.first == sink) goto out;

```

```

                }
            }
        }
        rep(i,0,N) pi[i] = min(pi[i] + dist[i], INF);
    }

    pair<ll, ll> maxflow(int s, int t) {
        ll totflow = 0, totcost = 0;
        while (path(s), seen[t]) {
            ll fl = INF;
            for (edge* x = par[t]; x; x = par[x->from])
                fl = min(fl, x->cap - x->flow);

            totflow += fl;
            for (edge* x = par[t]; x; x = par[x->from]) {
                x->flow += fl;
                ed[x->to][x->rev].flow -= fl;
            }
        }
        return {totflow, totcost/2};
    }
}
```

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

## MinCut.h

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

## GlobalMinCut.h

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

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

```
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) -> 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;
}
```

## 9.3 Matching

## hopcroftKarp.h

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

**Usage:**  $vi btoa(m, -1); \text{hopcroftKarp}(g, btoa);$

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

```
f612e4, 42 lines
```

```
bool dfs(int a, int L, vector<vi>& g, vi& btoa, vi& A, vi& B) {
    if (A[a] != L) return 0;
    A[a] = -1;
    for (int b : g[a]) if (B[b] == L + 1) {
        B[b] = 0;
        if (btoa[b] == -1 || dfs(btoa[b], L + 1, g, btoa, A, B))
            return btoa[b] = a, 1;
    }
    return 0;
}
```

```
int hopcroftKarp(vector<vi>& g, vi& btoa) {
    int res = 0;
    vi A(g.size()), B(btoa.size()), cur, next;
    for (;;) {
        fill(all(A), 0);
        fill(all(B), 0);
        cur.clear();
        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 : g[a]) {
                if (btoa[b] == -1) {
                    B[b] = lay;

```

```
                    islast = 1;
                }
                else if (btoa[b] != a && !B[b]) {
                    B[b] = lay;
                    next.push_back(btoa[b]);
                }
            }
            if (islast) break;
            if (next.empty()) return res;
            for (int a : next) A[a] = lay;
            cur.swap(next);
        }
        rep(a, 0, sz(g))
            res += dfs(a, 0, g, btoa, A, B);
    }
}
```

## DFSMatching.h

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

**Usage:**  $vi btoa(m, -1); \text{dfsMatching}(g, btoa);$

**Time:**  $\mathcal{O}(VE)$

```
522b98, 22 lines
```

```
bool find(int j, vector<vi>& g, vi& btoa, vi& vis) {
    if (btoa[j] == -1) return 1;
    vis[j] = 1; int di = btoa[j];
    for (int e : 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(g, match);
    vector<bool> lfound(n, true), seen(m);
    for (int it : match) if (it != -1) lfound[it] = false;
    vi q, cover;
    rep(i, 0, n) if (!lfound[i]) q.push_back(i);
    while (!q.empty()) {
        int i = q.back(); q.pop_back();
        lfound[i] = 1;
        for (int e : g[i]) if (!seen[e] && match[e] != -1) {
            seen[e] = true;
            q.push_back(match[e]);
        }
    }
    rep(i, 0, n) if (!lfound[i]) cover.push_back(i);
}
```

```
rep(i, 0, m) if (seen[i]) cover.push_back(n+i);
assert(sz(cover) == res);
return cover;
}
```

## WeightedMatching.h

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

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

```
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 "dummy" worker 0
        vi dist(m, INT_MAX), pre(m, -1);
        vector<bool> done(m + 1);
        do { // dijkstra
            done[j0] = true;
            int i0 = p[j0], j1, delta = INT_MAX;
            rep(j, 1, m) if (!done[j]) {
                auto cur = a[i0 - 1][j - 1] - u[i0] - v[j];
                if (cur < dist[j]) dist[j] = cur, pre[j] = j0;
                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]); // update alternating path
    int j1 = pre[j0];
    p[j0] = p[j1], j0 = j1;
}
rep(j, 1, m) if (p[j]) ans[p[j] - 1] = j - 1;
return {-v[0], ans}; // min cost
}
```

## GeneralMatching.h

**Description:** Matching for general graphs. Fails with probability  $N/\text{mod}$ .

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

```
../numerical/MatrixInverse-mod.h
cb1912, 40 lines
vector<pip> generalMatching(int N, vector<pip>& ed) {
    vector<vector<ll>> mat(N, vector<ll>(N));
    for (pip pa : ed) {
        int a = pa.first, b = pa.second, r = rand() % mod;
        mat[a][b] = r, mat[b][a] = (mod - r) % mod;
    }
    int r = matInv(A = mat), M = 2*N - r, fi, fj;
    assert(r % 2 == 0);
    if (M != N) do {
        mat.resize(M, vector<ll>(M));
        rep(i, 0, N) {
            mat[i].resize(M);
            rep(j, N, M) {
                int r = rand() % mod;
                mat[i][j] = r, mat[j][i] = (mod - r) % mod;
            }
        }
    } while (matInv(A = mat) != M);
}
```

```

vi has(M, 1); vector<pii> ret;
rep(i,0,M/2) {
    rep(j,i+1,M) if (has[i])
        fi = i; fj = j; goto done;
} assert(0); done:
if (fj < N) ret.emplace_back(fi, fj);
has[fi] = has[fj] = 0;
rep(sw,0,2) {
    ll a = modpow(A[fi][fj], mod-2);
    rep(i,0,M) if (has[i] && A[i][fj]) {
        ll b = A[i][fj] * a % mod;
        rep(j,0,M) A[i][j] = (A[i][j] - A[fi][j] * b) % mod;
    }
    swap(fi,fj);
}
return ret;
}

```

## 9.4 DFS algorithms

### SCC.h

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

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

**Time:**  $\mathcal{O}(E + V)$  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 : g[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 != j);
        f(cont); cont.clear();
        ncomps++;
    }
    return val[j] = low;
}
template<class G, class F> void scc(G& g, F f) {
    int n = sz(g);
    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:**  $\mathcal{O}(E + V)$  c6b7c7, 32 lines

```

vi num, st;
vector<vector<pii>> ed;
int Time;
template<class F>
int dfs(int at, int par, F& f) {
    int me = num[at] = ++Time, top = me;
    for (auto [y, e] : ed[at]) if (e != par) {
        if (num[y]) {
            top = min(top, num[y]);
            if (num[y] < me)
                st.push_back(e);
        } else {
            int si = sz(st);
            int up = dfs(y, e, f);
            top = min(top, up);
            if (up == me)
                st.push_back(e);
            f(vi(st.begin() + si, st.end()));
            st.reserve(si);
        }
        else if (up < me) st.push_back(e);
        else /* e is a bridge */
    }
    return top;
}

template<class F>
void bicomps(F f) {
    num.assign(sz(ed), 0);
    rep(i,0,sz(ed)) if (!num[i]) dfs(i, -1, f);
}

```

### 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:**  $\mathcal{O}(V + E)$  780b64, 15 lines

```

vi eulerWalk(vector<vector<pii>>& gr, int nedges, int src=0) {
    int n = sz(gr);
    vi D(n), its(n), eu(nedges), ret, s = {src};
    D[src]++;
    while (!s.empty()) {
        int x = s.back(), y, e, &it = its[x], end = sz(gr[x]);
        if (it == end) ret.push_back(x); s.pop_back(); continue;
        tie(y, e) = gr[x][it++];
        if (!eu[e]) {
            D[x]--, D[y]++;
            eu[e] = 1; s.push_back(y);
        }
        for (int x : D) if (x < 0 || sz(ret) != nedges+1) return {};
    }
    return {ret.rbegin(), ret.rend()};
}

```

## 9.5 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:**  $\mathcal{O}(3^{n/3})$ , much faster for sparse graphs b0d5b1, 12 lines

```

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

```

```

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

```

### MaximumClique.h

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

**Time:** Runs in about 1s for n=155 and worst case random graphs (p=.90). Runs faster for sparse graphs. f7c0bc, 49 lines

```

typedef vector<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;
            q.push_back(R.back().i);
            vv T;
            for (auto v:R) if (e[R.back().i][v.i]) T.push_back({v.i});
            if (sz(T)) {
                if (S[lev]++ / ++pk < limit) init(T);
                int j = 0, mxk = 1, mnk = max(sz(qmax) - sz(q) + 1, 1);
                C[1].clear(), C[2].clear();
                for (auto v : T) {
                    int k = 1;
                    auto f = [&](int i) { return e[v.i][i]; };
                    while (any_of(all(C[k]), f)) k++;
                    if (k > mxk) mxk = k, C[mxk + 1].clear();
                    if (k < mnk) T[j++].i = v.i;
                    C[k].push_back(v.i);
                }
                if (j > 0) T[j - 1].d = 0;
                rep(k,mnk,mxk + 1) for (int i : C[k])
                    T[j].i = i, T[j++].d = k;
                expand(T, lev + 1);
            } else if (sz(q) > sz(qmax)) qmax = q;
            q.pop_back(), R.pop_back();
        }
    }
    vi maxClique() { init(V), expand(V); return qmax; }
    Maxclique(vb conn) : e(conn), C(sz(e)+1), S(sz(C)), old(S) {
        rep(i,0,sz(e)) V.push_back({i});
    }
}

```

**MaximumIndependentSet.h**

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

## 9.6 Trees

**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:**  $O(N \log N + Q)$

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

0f62fb, 21 lines

```
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)];}
};

HLD.h
```

**Description:** Heavy-Light Decomposition for path updates and queries on trees.

**Usage:** HeavyLightDecomposition hld(n); // n is number of nodes

hld.add\_edge(u,v); // add edge between u and v

hld.DFS(1,1); // build DFS tree with root 1

hld.HLD(1); // build HLD with root 1

hld.update\_up\_HLD(u,v,val); // update path from u to v with value val

hld.query\_up\_HLD(u,v); // query sum on path from u to v

**Time:**  $O(\log^2 N)$  per operation

acd451, 177 lines

```
/*
 1-indexed tree, root = 1
 You are given a rooted tree consisting of n nodes. The
 nodes are numbered 1,2,...,n, and node 1 is the
 root. Each node has a value.
 Your task is to process following types of queries:
```

Update each vertice in path u to v by 1  
calculate sum of each edge

\*/

```
struct ITLazy{
    long long sum, lazy;
};
```

```
struct HeavyLightDecomposition{
    int N, maxx_height;
    int num_chain, pos_in_base;
    vector<int> depth;
    vector<int> numChild;
    vector<int> chainHead, chainInd;
    vector<int> Base, posInBase;
```

**MaximumIndependentSet LCA HLD**

```
vector<vector<int>> Anc;
vector<ITLazy> IT;
vector<vector<int>> Adj;

HeavyLightDecomposition(int N) : N(N) {
    const int LOG2 = int(log2(N));
    depth.resize(N+1, 0);
    numChild.resize(N+1, 0);
    chainHead.resize(N+1, 0); chainInd.resize(N+1, 0);
    Base.resize(N+1, 0); posInBase.resize(N+1, 0);
    IT.resize(4*N+1, {0, 0});
    Adj.resize(N+1);
    Anc.resize(N+1, vector<int>(LOG2+1, 0));

    maxx_height = -1;
    depth[1] = 1;
    num_chain = 1;
    pos_in_base = 0;
    Anc[1][0] = 1;
}

void add_edge(int u, int v){
    Adj[u].push_back(v);
    Adj[v].push_back(u);
}

void DFS(int u, int par){
    numChild[u] = 1;
    for(int v: Adj[u]){
        if(v == par) continue;
        depth[v] = depth[u] + 1;
        Anc[v][0] = u;
        maxx_height = max(maxx_height, depth[v]);
        for(int j=1, j <= int(log2(maxx_height)), j++) {
            Anc[v][j] = Anc[Anc[v][j-1]][j-1];
        }
        DFS(v, u);
        numChild[u] = numChild[u] + numChild[v];
    }
}

void HLD(int u){
    if(!chainHead[num_chain]){
        chainHead[num_chain] = u;
    }

    chainInd[u] = num_chain;
    posInBase[u] = ++pos_in_base;
    Base[pos_in_base] = u;
    int special_vertices = -1;
    for(int v: Adj[u]){
        if(v == Anc[u][0]) continue;
        if(special_vertices == -1 || numChild[v] > numChild[special_vertices]){
            special_vertices = v;
        }
    }

    if(special_vertices != -1) HLD(special_vertices);

    for(int v: Adj[u]){
        if(v == Anc[u][0] || v == special_vertices)
            continue;
        num_chain++;
        HLD(v);
    }
}

int Jump(int u, int v){
```

```
assert(depth[u] <= depth[v]);
int delta = depth[v] - depth[u];
for(int i=log2(maxx_height); i >= 0; i--){
    if(delta >> i & 1){
        v = Anc[v][i];
    }
}
return v;
}

int LCA(int u, int v){
    if(u == v) return u;
    if(depth[u] > depth[v]) swap(u, v);
    v = Jump(u, v);
    if(u == v) return u;
    for(int i=log2(depth[u]); i >= 0; i--){
        if(Anc[u][i] != Anc[v][i]){
            u = Anc[u][i];
            v = Anc[v][i];
        }
    }
    return Anc[u][0];
}

void lazyUpdate(int id, int L, int R, int mid){
    IT[id << 1].lazy += IT[id].lazy;
    IT[id << 1 | 1].lazy += IT[id].lazy;

    IT[id << 1].sum += IT[id].lazy * 1ll * (mid - L + 1);
    IT[id << 1 | 1].sum += IT[id].lazy * 1ll * (R - mid);

    IT[id].lazy = 0;
}

void update(int id, int L, int R, int u, int v, long long val){
    if(v < L || R < u) return;
    if(u <= L && R <= v){
        IT[id].lazy += val;
        IT[id].sum += 1ll * (R - L + 1) * val;
        return;
    }
    int mid = (L + R) >> 1;
    lazyUpdate(id, L, R, mid);
    update(id << 1, L, mid, u, v, val);
    update(id << 1 | 1, mid + 1, R, u, v, val);
    IT[id].sum = IT[id << 1].sum + IT[id << 1 | 1].sum;
}

void update_up_HLD(int beg, int en, long long val){
    while(1 + 1 == 2){
        if(chainInd[beg] == chainInd[en]){
            update(1, 1, N, posInBase[en], posInBase[beg], val);
            break;
        }
        else{
            update(1, 1, N, posInBase[chainHead[chainInd[beg]]], posInBase[beg], val);
            beg = Anc[chainHead[chainInd[beg]]][0];
        }
    }
}

long long get(int id, int L, int R, int u, int v){
    if(v < L || R < u) return 0;
    if(u <= L && R <= v) return IT[id].sum;
    int mid = (L + R) >> 1;
```

```

lazyUpdate(id, L, R, mid);
long long t1 = get(id << 1, L, mid, u, v);
long long t2 = get(id << 1 | 1, mid + 1, R, u, v);
return t1 + t2;
}

long long query_up_HLD(int beg, int en){
assert(depth[beg] >= depth[en]);
long long res = 0;
while(l + 1 == 2){
    if(chainInd[beg] == chainInd[en]){
        res = res + get(1, 1, N, posInBase[en],
                        posInBase[beg]);
        return res;
    }
    else{
        res = res + get(1, 1, N, posInBase[chainHead[
            chainInd[beg]]], posInBase[beg]);
        beg = Anc[chainHead[chainInd[beg]]][0];
    }
}
return res;
}

```

## DirectedMST.h

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

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

.../data-structures/UnionFindRollback.h 39e620, 60 lines

```

struct Edge { int a, b; ll w; };
struct Node {
    Edge key;
    Node *l, *r;
    ll delta;
    void prop() {
        key.w += delta;
        if (l) l->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->l, (a->r = merge(b, a->r)));
    return a;
}

void pop(Node*& a) { a->prop(); a = merge(a->l, 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});
    ll res = 0;
    vi seen(n, -1), path(n), par(n);
    seen[r] = r;
    vector<Edge> Q(n), in(n, {-1, -1}), comp;
    deque<tuple<int, int, vector<Edge>>> cycs;
    rep(s, 0, n) {
        int u = s, qi = 0, w;
        while (seen[u] < 0) {
            if (!heap[u]) return {-1, {}};
            Edge e = heap[u]->top();
            heap[u]->delta -= e.w, pop(heap[u]);
            Q[qi] = e, path[qi++]

```

```

                = u, seen[u] = s;
        }
        res += e.w, u = uf.find(e.a);
        if (seen[u] == s) {
            Node* cyc = 0;
            int end = qi, time = uf.time();
            do cyc = merge(cyc, heap[w = path[--qi]]);
            while (uf.join(u, w));
            u = uf.find(u), heap[u] = cyc, seen[u] = -1;
            cycs.push_front({u, time, {&Q[qi], &Q[en]}});
        }
        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};
}

```

```

        res += e.w, u = uf.find(e.a);
        if (seen[u] == s) {
            Node* cyc = 0;
            int end = qi, time = uf.time();
            do cyc = merge(cyc, heap[w = path[--qi]]);
            while (uf.join(u, w));
            u = uf.find(u), heap[u] = cyc, seen[u] = -1;
            cycs.push_front({u, time, {&Q[qi], &Q[en]}});
        }
        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};
}

```

## 9.7 Math

## 9.7.1 Number of Spanning Trees

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

## 9.7.2 Erdős–Gallai theorem

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

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

## Geometry (10)

## 10.1 Geometric primitives

## Point.h

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

```

47ec0a, 28 lines
template <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); }
    bool operator==(P p) const { return tie(x,y)==tie(p.x,p.y); }
    P operator+(P p) const { return P(x+p.x, y+p.y); }
    P operator-(P p) const { return P(x-p.x, y-p.y); }
    P operator*(T d) const { return P(x*d, y*d); }
    P operator/(T d) const { return P(x/d, y/d); }
    T dot(P p) const { return x*p.x + y*p.y; }
    T cross(P p) const { return x*p.y - y*p.x; }
    T cross(P a, P b) const { return (a-*this).cross(b-*this); }

```

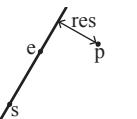
```

    T dist2() const { return x*x + y*y; }
    double dist() const { return sqrt((double)dist2()); }
    // angle to x-axis in interval [-pi, pi]
    double angle() const { return atan2(y, x); }
    P unit() const { return *this/dist(); } // makes dist()==1
    P perp() const { return P(-y, x); } // rotates +90 degrees
    P normal() const { return perp().unit(); }
    // returns point rotated 'a' radians ccw around the origin
    P rotate(double a) const {
        return P(x*cos(a)-y*sin(a), x*sin(a)+y*cos(a)); }
    friend ostream& operator<<(ostream& os, P p) {
        return os << "(" << p.x << ", " << p.y << ")";
    }
};
```

## lineDistance.h

## Description:

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



"Point.h"

```

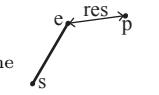
template<class P>
double lineDist(const P& a, const P& b, const P& p) {
    return (double)(b-a).cross(p-a)/(b-a).dist();
}
```

## SegmentDistance.h

## Description:

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

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



"Point.h"

```

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

## SegmentIntersection.h

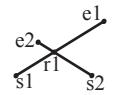
## Description:

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

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

```

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



```

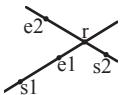
if (onSegment(a, b, c)) s.insert(c);
if (onSegment(a, b, d)) s.insert(d);
return {all(s)};
}

```

## lineIntersection.h

**Description:**

If a unique intersection point of the lines going through  $s_1, e_1$  and  $s_2, e_2$  exists  $\{1, \text{point}\}$  is returned. If no intersection point exists  $\{0, (0, 0)\}$  is returned and if infinitely many exists  $\{-1, (0, 0)\}$  is returned. The wrong position will be returned if P is an integer point and the desired point doesn't have integer coordinates. Products of three coordinates are used in intermediate steps so watch out for overflow if using int or ll.



**Usage:** auto res = lineInter(s1,e1,s2,e2);  
**if** (res.first == 1)  
cout << "intersection point at "<< res.second << endl;  
"Point.h" a01f81, 8 lines

```

template<class P>
pair<int, P> lineInter(P s1, P e1, P s2, P e2) {
    auto d = (e1 - s1).cross(e2 - s2);
    if (d == 0) // if parallel
        return {-(s1.cross(e1, s2) == 0), P(0, 0)};
    auto p = s2.cross(e1, e2), q = s2.cross(e2, s1);
    return {1, (s1 * p + e1 * q) / d};
}

```

## sideOf.h

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

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

"Point.h" 3af81c, 9 lines

```

template<class P>
int sideOf(const P& s, P e, P p) { return sgn(s.cross(e, p)); }

template<class P>
int sideOf(const P& s, const P& e, const P& p, double eps) {
    auto a = (e-s).cross(p-s);
    double l = (e-s).dist()*eps;
    return (a > 1) - (a < -1);
}

```

## OnSegment.h

**Description:** Returns true iff  $p$  lies on the line segment from  $s$  to  $e$ . Use  $(\text{segDist}(s, e, p) \leq \text{epsilon})$  instead when using Point<double>.

"Point.h" c597e8, 3 lines

```

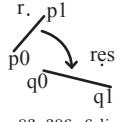
template<class P> bool onSegment(P s, P e, P p) {
    return p.cross(s, e) == 0 && (s - p).dot(e - p) <= 0;
}

```

## linearTransformation.h

**Description:**

Apply the linear transformation (translation, rotation and scaling) which takes line  $p_0-p_1$  to line  $q_0-q_1$  to point  $r$ .



```

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

```

## LineProjectionReflection.h

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

"Point.h" b5562d, 5 lines

```

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

```

## Angle.h

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

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

of0602, 35 lines

```

struct Angle {
    int x, y;
    int t;
    Angle(int x, int y, int t=0) : x(x), y(y), t(t) {}
    Angle operator-(Angle b) const { return {x-b.x, y-b.y, t}; }
    int half() const {
        assert(x || y);
        return y < 0 || (y == 0 && x < 0);
    }
    Angle t90() const { return {-y, x, t + (half() && x >= 0)? 0 : 90}; }
    Angle t180() const { return {-x, -y, t + half()}; }
    Angle t360() const { return {x, y, t + 1}; }
};

bool operator<(Angle a, Angle b) {
    // add a.dist2() and b.dist2() to also compare distances
    return make_tuple(a.t, a.half(), a.y * (ll)b.x) <
        make_tuple(b.t, b.half(), a.x * (ll)b.y);
}

// Given two points, this calculates the smallest angle between
// them, i.e., the angle that covers the defined line segment.
pair<Angle, Angle> segmentAngles(Angle a, Angle b) {
    if (b < a) swap(a, b);
    return (b < a.t180()?
        make_pair(a, b) : make_pair(b, a.t360()));
}

Angle operator+(Angle a, Angle b) { // point a + vector b
    Angle r(a.x + b.x, a.y + b.y, a.t);
    if (a.t180() < r.r) r.t--;
    return r.t180() < a ? a.t360() : r;
}

Angle angleDiff(Angle a, Angle b) { // angle b - angle a
    int tu = b.t - a.t; a.t = b.t;
    return {a.x*b.x + a.y*b.y, a.x*b.y - a.y*b.x, tu - (b < a)? 0 : 360};
}

```

### 10.2 Circles

#### CircleIntersection.h

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

"Point.h" 84d6d3, 11 lines

```

typedef Point<double> P;
bool circleInter(P a, P b, double r1, double r2, pair<P, P>* out) {
    if (a == b) { assert(r1 != r2); return false; }
    P vec = b - a;
    double d2 = vec.dist2(), sum = r1+r2, dif = r1-r2,
    p = (d2 + r1*r1 - r2*r2)/(d2*2), h2 = r1*r1 - p*p*d2;
}

```

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

P mid = a + vec\*p, per = vec.perp() \* sqrt(fmax(0, h2) / d2);
\*out = {mid + per, mid - per};
**return** **true**;

}

## CircleTangents.h

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

"Point.h" b0153d, 13 lines

```

template<class P>
vector<pair<P, P>> tangents(P c1, double r1, P c2, double r2) {
    P d = c2 - c1;
    double dr = r1 - r2, d2 = d.dist2(), h2 = d2 - dr * dr;
    if (d2 == 0 || h2 < 0) return {};
    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;
}

```

## CircleLine.h

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

"Point.h" e0cfba, 9 lines

```

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

```

## CirclePolygonIntersection.h

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

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

"../../../../content/geometry/Point.h" a1ee63, 19 lines

```

typedef Point<double> P;
#define arg(p, q) atan2(p.cross(q), p.dot(q))
double circlePoly(P c, double r, vector<P> ps) {
    auto tri = [&](P p, P q) {
        auto r2 = r * r / 2;
        P d = q - p;
        auto a = d.dot(p)/d.dist2(), b = (p.dist2()-r*r)/d.dist2();
        auto det = a * a - b;
        if (det <= 0) return arg(p, q) * r2;
        auto s = max(0, -a-sqrt(det)), t = min(1, -a+sqrt(det));
        if (t < 0 || l1 <= s) return arg(p, q) * r2;
        P u = p + d * s, v = p + d * t;
        return arg(p,u) * r2 + u.cross(v)/2 + arg(v,q) * r2;
    };
    auto sum = 0.0;
    rep(i, 0, sz(ps))
        sum += tri(ps[i] - c, ps[(i + 1) % sz(ps)] - c);
    return sum;
}

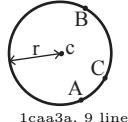
```

}

## circumcircle.h

**Description:**

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



1caa3a, 9 lines

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

## MinimumEnclosingCircle.h

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

```
"circumcircle.h" 09dd0a, 17 lines
pair<P, double> mec(vector<P> ps) {
    shuffle(all(ps), mt19937(time(0)));
    P o = 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};
}
```

## 10.3 Polygons

## InsidePolygon.h

**Description:** Returns true if p lies within the polygon. If strict is true, it returns false for points on the boundary. The algorithm uses products in intermediate steps so watch out for overflow.**Usage:** `vector<P> v = {P{4,4}, P{1,2}, P{2,1}};``bool in = inPolygon(v, P{3, 3}, false);`**Time:**  $\mathcal{O}(n)$ 

```
"Point.h", "OnSegment.h", "SegmentDistance.h" 2bf504, 11 lines
template<class P>
bool inPolygon(vector<P> &p, P a, bool strict = true) {
    int cnt = 0, n = sz(p);
    rep(i, 0, n) {
        P q = p[(i + 1) % n];
        if (onSegment(p[i], q, a)) return !strict;
        //or: if (segDist(p[i], q, a) <= eps) return !strict;
        cnt += ((a.y<p[i].y) - (a.y<q.y)) * a.cross(p[i], q) > 0;
    }
    return cnt;
}
```

## PolygonArea.h

**Description:** Returns twice the signed area of a polygon. Clockwise enumeration gives negative area. Watch out for overflow if using int as T!`"Point.h" f12300, 6 lines`

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

## PolygonCenter.h

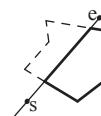
**Description:** Returns the center of mass for a polygon.**Time:**  $\mathcal{O}(n)$ 

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

## PolygonCut.h

**Description:**

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

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

f2b7d4, 13 lines

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

## PolygonUnion.h

**Description:** Calculates the area of the union of  $n$  polygons (not necessarily convex). The points within each polygon must be given in CCW order. (Epsilon checks may optionally be added to sideOf/sgn, but shouldn't be needed.)**Time:**  $\mathcal{O}(N^2)$ , where  $N$  is the total number of points`"Point.h", "sideOf.h"` 3931c6, 33 lines

```
typedef Point<double> P;
double rat(P a, P b) { return sgn(b.x) ? a.x/b.x : a.y/b.y; }
double polyUnion(vector<vector<P>> &poly) {
    double ret = 0;
    rep(i, 0, sz(poly)) rep(v, 0, sz(poly[i])) {
        P A = poly[i][v], B = poly[i][(v + 1) % sz(poly[i])];
        vector<pair<double, int>> segs = {{0, 0}, {1, 0}};
        rep(j, 0, sz(poly)) if (i != j) {
            rep(u, 0, sz(poly[j])) {
                P C = poly[j][u], D = poly[j][(u + 1) % sz(poly[j])];
                int sc = sideOf(A, B, C), sd = sideOf(A, B, D);
                if (sc != sd) {
                    double sa = C.cross(D, A), sb = C.cross(D, B);
                    if (min(sc, sd) < 0)
                        segs.emplace_back(sa / (sa - sb), sgn(sc - sd));
                } else if (!sc && !sd && j < i && sgn((B-A).dot(D-C)) > 0) {
                    segs.emplace_back(rat(C - A, B - A), 1);
                    segs.emplace_back(rat(D - A, B - A), -1);
                }
            }
        }
    }
    return ret;
}
```

```
} 71446b, 14 lines
}
sort(all(segs));
for (auto &s : segs) s.first = min(max(s.first, 0.0), 1.0);
int cnt = segs[0].second;
rep(j, 1, sz(segs)) {
    if (!cnt) sum += segs[j].first - segs[j - 1].first;
    cnt += segs[j].second;
}
ret += A.cross(B) * sum;
}
return ret / 2;
}
```

## ConvexHull.h

**Description:**

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

**Time:**  $\mathcal{O}(n \log n)$ `"Point.h"` 310954, 13 lines

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

## HullDiameter.h

**Description:** Returns the two points with max distance on a convex hull (ccw, no duplicate/collinear points).**Time:**  $\mathcal{O}(n)$ `"Point.h"` c571b8, 12 lines

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

## PointInsideHull.h

**Description:** Determine whether a point t lies inside a convex hull (CCW order, with no collinear points). Returns true if point lies within the hull. If strict is true, points on the boundary aren't included.**Time:**  $\mathcal{O}(\log N)$ `"Point.h", "sideOf.h", "OnSegment.h"` 71446b, 14 lines

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

```

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

```

## LineHullIntersection.h

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

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

```
"Point.h"
7cf45b, 39 lines
#define cmp(i, j) sgn(dir.perp().cross(poly[(i)%n]-poly[(j)%n]))
#define extr(i) cmp(i + 1, i) >= 0 && cmp(i, i - 1 + n) < 0
template <class P> int extrVertex(vector<P>& poly, P dir) {
    int n = sz(poly), lo = 0, hi = n;
    if (extr(0)) return 0;
    while (lo + 1 < hi) {
        int m = (lo + hi) / 2;
        if (extr(m)) return m;
        int ls = cmp(lo + 1, lo), ms = cmp(m + 1, m);
        (ls < ms || (ls == ms && ls == cmp(lo, m)) ? hi : lo) = m;
    }
    return lo;
}

#define cmpL(i) sgn(a.cross(poly[i], b))
template <class P>
array<int, 2> lineHull(P a, P b, vector<P>& poly) {
    int endA = extrVertex(poly, (a - b).perp());
    int endB = extrVertex(poly, (b - a).perp());
    if (cmpL(endA) < 0 || cmpL(endB) > 0)
        return {-1, -1};
    array<int, 2> res;
    rep(i, 0, 2) {
        int lo = endB, hi = endA, n = sz(poly);
        while ((lo + 1) % n != hi) {
            int m = ((lo + hi + (lo < hi ? 0 : n)) / 2) % n;
            (cmpL(m) == cmpL(endB) ? lo : hi) = m;
        }
        res[i] = (lo + !cmpL(hi)) % n;
        swap(endA, endB);
    }
    if (res[0] == res[1]) return {res[0], -1};
    if (!cmpL(res[0]) && !cmpL(res[1])) {
        switch ((res[0] - res[1] + sz(poly) + 1) % sz(poly)) {
            case 0: return {res[0], res[0]};
            case 2: return {res[1], res[1]};
        }
    }
    return res;
}
```

## 10.4 Misc. Point Set Problems

## ClosestPair.h

**Description:** Finds the closest pair of points.

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

```
"Point.h"
ac41a6, 17 lines
typedef Point<ll> P;
pair<P, P> closest(vector<P> v) {
    assert(sz(v) > 1);
    set<P> S;
```

```

sort(all(v), [](P a, P b) { return a.y < b.y; });
pair<ll, pair<P, P>> ret{LLONG_MAX, {P(), P()}};
int j = 0;
for (P p : v) {
    P d{1 + (ll)sqrt(ret.first), 0};
    while (v[j].y <= p.y - d.x) S.erase(v[j++]);
    auto lo = S.lower_bound(p - d), hi = S.upper_bound(p + d);
    for (; lo != hi; ++lo)
        ret = min(ret, {(*lo - p).dist2(), {*lo, p}});
    S.insert(p);
}
return ret.second;
}
```

## ManhattanMST.h

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

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

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

## kdTree.h

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

```
"Point.h"
bac5b0, 63 lines
typedef long long T;
typedef Point<T> P;
const T INF = numeric_limits<T>::max();

bool on_x(const P& a, const P& b) { return a.x < b.x; }
bool on_y(const P& a, const P& b) { return a.y < b.y; }

struct Node {
    P pt; // if this is a leaf, the single point in it
    T x0 = INF, xl = -INF, y0 = INF, yl = -INF; // bounds
    Node *first = 0, *second = 0;

    T distance(const P& p) { // min squared distance to a point
        T x = (p.x < x0 ? x0 : p.x > xl ? xl : p.x);
        T y = (p.y < y0 ? y0 : p.y > yl ? yl : p.y);
        return (P(x, y) - p).dist2();
    }

    Node(vector<P>&& vp) : pt(vp[0]) {
        for (P p : vp) {
```

```

        x0 = min(x0, p.x); xl = max(xl, p.x);
        y0 = min(y0, p.y); yl = max(yl, p.y);
    }
    if (vp.size() > 1) {
        // split on x if width >= height (not ideal...)
        sort(all(vp), xl - x0 >= yl - y0 ? on_x : on_y);
        // divide by taking half the array for each child (not
        // best performance with many duplicates in the middle)
        int half = sz(vp)/2;
        first = new Node({vp.begin(), vp.begin() + half});
        second = new Node({vp.begin() + half, vp.end()});
    }
}
```

## struct KDTTree {

```
Node* root;
KDTTree(const vector<P>& vp) : root(new Node({all(vp)})) {}
```

```
pair<T, P> search(Node *node, const P& p) {
    if (!node->first) {
        // uncomment if we should not find the point itself:
        // if (p == node->pt) return {INF, P()};
        return make_pair((p - node->pt).dist2(), node->pt);
    }
```

```
Node *f = node->first, *s = node->second;
T bfirst = f->distance(p), bsec = s->distance(p);
if (bfirst > bsec) swap(bsec, bfirst), swap(f, s);
```

```
// search closest side first, other side if needed
auto best = search(f, p);
if (bsec < best.first)
    best = min(best, search(s, p));
return best;
}
```

```
// find nearest point to a point, and its squared distance
// (requires an arbitrary operator< for Point)
pair<T, P> nearest(const P& p) {
    return search(root, p);
}
```

## DelaunayTriangulation.h

**Description:** Computes the Delaunay triangulation of a set of points. Each circumcircle contains none of the input points. If any three points are collinear or any four are on the same circle, behavior is undefined.

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

```
"Point.h", "3dHull.h"
c0e7bc, 10 lines
template <class P, class F>
void delaunay(vector<P>& ps, F trifun) {
    if (sz(ps) == 3) { int d = (ps[0].cross(ps[1], ps[2]) < 0);
        trifun(0, 1+d, 2-d); }
    vector<P> p3;
    for (P p : ps) p3.emplace_back(p.x, p.y, p.dist2());
    if (sz(ps) > 3) for (auto t:hull3d(p3)) if ((p3[t.b]-p3[t.a]).cross(p3[t.c]-p3[t.a]).dot(P3(0,0,1)) < 0)
        trifun(t.a, t.c, t.b);
}
```

## FastDelaunay.h

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

```

Time:  $\mathcal{O}(n \log n)$ 
"Point.h" eefdf5, 88 lines
typedef Point<ll> P;
typedef struct Quad* Q;
typedef __int128_t lll; // (can be ll if coords are < 2e4)
P arb(LLONG_MAX,LLONG_MAX); // not equal to any other point

struct Quad {
    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(); }
} *H;

bool circ(P p, P a, P b, P c) { // is p in the circumcircle?
    lll 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(new Quad{0})));
    H = r->o; r->r()>r() = r;
    rep(i,0,4) r = r->rot, r->p = arb, r->o = i & 1 ? r : r->r();
    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 q = makeEdge(a->F(), b->p);
    splice(q, a->next());
    splice(q->r(), b);
    return q;
}

pair<Q,Q> rec(const vector<P>& s) {
    if (sz(s) <= 3) {
        Q a = makeEdge(s[0], s[1]), b = makeEdge(s[1], s.back());
        if (sz(s) == 2) return { a, a->r() };
        splice(a->r(), b);
        auto side = s[0].cross(s[1], s[2]);
        Q c = side ? connect(b, a) : 0;
        return {side < 0 ? c->r() : a, side < 0 ? c : b->r() };
    }

    #define H(e) e->F(), e->p
    #define valid(e) (e->F().cross(H(base)) > 0)
    Q 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;
            splice(e, e->prev()); \
            splice(e->r(), e->r()>prev()); \
            e->o = H; H = e; e = t; \
        }
    for (;;) {
        DEL(LC, base->r(), o); DEL(RC, base, prev());
        if (!valid(LC) && !valid(RC)) break;
    }
}

```

## PolyhedronVolume Point3D 3dHull sphericalDistance

```

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++])->mark) ADD;
    return pts;
}

10.5 3D
PolyhedronVolume.h
Description: Magic formula for the volume of a polyhedron. Faces should point outwards.
Time:  $\mathcal{O}(n^2)$ 
"Point3D.h" 5b45fc, 49 lines

```

```

typedef Point3D<double> P3;
struct PR {
    void ins(int x) { (a == -1 ? a : b) = x; }
    void rem(int x) { (a == x ? a : b) = -1; }
    int cnt() { return (a != -1) + (b != -1); }
    int a, b;
};

struct F { P3 q; int a, b, c; };

vector<F> hull3d(const vector<P3>& A) {
    assert(sz(A) >= 4);
    vector<vector<PR>> E(sz(A)), vector<PR>(sz(A), {-1, -1}));
#define E(x,y) E[f.x][f.y]
    vector<F> FS;
    auto mf = [&](int i, int j, int k, int l) {
        P3 q = (A[j] - A[i]).cross((A[k] - A[i]));
        if (q.dot(A[l]) > q.dot(A[i]))
            q = q * -1;
        F f{q, i, j, k};
        E(a,b).ins(k); E(a,c).ins(j); E(b,c).ins(i);
        FS.push_back(f);
    };
    rep(i,0,4) rep(j,i+1,4) rep(k,j+1,4)
        mf(i, j, k, 6 - i - j - k);

    rep(i,4,sz(A)) {
        rep(j,0,sz(FS)) {
            F f = FS[j];
            if (f.q.dot(A[i]) > f.q.dot(A[f.a])) {
                E(a,b).rem(f.c);
                E(a,c).rem(f.b);
                E(b,c).rem(f.a);
                swap(FS[j--], FS.back());
                FS.pop_back();
            }
        }
        int nw = sz(FS);
        rep(j,0,nw) {
            F f = FS[j];
#define C(a, b, c) if (E(a,b).cnt() != 2) mf(f.a, f.b, i, f.c);
            C(a, b, c); C(a, c, b); C(b, c, a);
        }
    }
    for (F& it : FS) if ((A[it.b] - A[it.a]).cross(
        A[it.c] - A[it.a]).dot(it.q) <= 0) swap(it.c, it.b);
    return FS;
}

```

## sphericalDistance.h

```

Description: Returns the shortest distance on the sphere with radius radius between the points with azimuthal angles (longitude) f1 ( $\phi_1$ ) and f2 ( $\phi_2$ ) from x axis and zenith angles (latitude) t1 ( $\theta_1$ ) and t2 ( $\theta_2$ ) from z axis (0 = north pole). All angles measured in radians. The algorithm starts by converting the spherical coordinates to cartesian coordinates so if that is what you have you can use only the two last rows. dx*radius is then the difference between the two points in the x direction and d*radius is the total distance between the points.
Time:  $\mathcal{O}(1)$ 
"sphericalDistance.h" 611f07, 8 lines
double sphericalDistance(double f1, double t1,

```

```

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

```

## Strings (11)

### 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  $\rightarrow$  0010123). Can be used to find all occurrences of a string.

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

d4375c, 16 lines

```

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

vi match(const string& s, const string& pat) {
    vi p = pi(pat + '\0' + s), res;
    rep(i,sz(p)-sz(s),sz(p))
        if (p[i] == sz(pat)) res.push_back(i - 2 * sz(pat));
    return res;
}

```

### Zfunc.h

**Description:**  $z[i]$  computes the length of the longest common prefix of  $s[i:]$  and  $s$ , except  $z[0] = 0$ . (abacaba  $\rightarrow$  0010301)

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

ee09e2, 12 lines

```

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

```

### Manacher.h

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

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

e7ad79, 13 lines

```

array<vi, 2> manacher(const string& s) {
    int n = sz(s);
    array<vi, 2> p = {vi(n+1), vi(n)};
    rep(z,0,2) for (int i=0, l=0, r=0; i < n; i++) {
        int t = r-i+z;
        if (i < r) p[2][i] = min(t, p[2][l+t]);
        int L = i-p[2][i], R = i+p[2][i]-z;
        while (L>=1 && R+1<n && s[L-1] == s[R+1])
            p[2][i]++, L--, R++;
        if (R>r) l=L, r=R;
    }
}

```

```

    return p;
}

```

### MinRotation.h

**Description:** Finds the lexicographically smallest rotation of a string.

**Usage:** `rotate(v.begin(), v.begin() + minRotation(v), v.end())`

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

d07aa42, 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 || s[a+k] < s[b+k]) {b += max(0, k-1); break;}
        if (s[a+k] > s[b+k]) {a = b; break;}
    }
    return a;
}

```

### SuffixArray.h

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

bc716b, 22 lines

```

struct SuffixArray {
    vi sa, lcp;
    SuffixArray(string& s, int lim=256) { // or basic_string<int>
        int n = sz(s) + 1, k = 0, a, b;
        vi x(all(s)), y(n), ws(max(n, lim));
        x.push_back(0), 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) a = sa[i - 1], b = sa[i], x[b] =
                (y[a] == y[b] && y[a + j] == y[b + j]) ? p - 1 : p++;
        }
        for (int i = 0, j; i < n - 1; lcp[x[i++]] = k)
            for (k && k--, j = sa[x[i] - 1];
                 sa[i + k] == s[j + k]; k++);
    }
};

```

### SuffixTree.h

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

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

aae0b8, 50 lines

```

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

```

```

if (q===-1 || c==toi(a[q])) q++; else {
    l[m+1]=i; p[m+1]=m; l[m]=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]-1[v]; }
    if (q==r[m]) s[m]=v; else s[m]=m+2;
    q=r[v]-(q-r[m]); m+=2; goto suff;
}
}

```

### SuffixTree(string a) : a(a)

```

    fill(r,r+N,sz(a));
    memset(s, 0, sizeof s);
    memset(t, -1, sizeof t);
    fill(t[1],t[1]+ALPHA,0);
    s[0] = 1; l[0] = 1[l] = -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  $\text{ALPHA} = 28$ )

```

pii best;
int lcs(int node, int i1, int i2, int olen) {
    if (l[node] <= i1 && i1 < r[node]) return 1;
    if (l[node] <= i2 && i2 < r[node]) return 2;
    int mask = 0, len = node ? olen + (r[node] - l[node]) : 0;
    rep(c,0,ALPHA) if (t[node][c] != -1)
        mask |= lcs(t[node][c], i1, i2, len);
    if (mask == 3)
        best = max(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.

2d2a67, 44 lines

```

// Arithmetic mod  $2^{64}-1$ . 2x slower than mod  $2^{64}$  and more
// code, but works on evil test data (e.g. Thue-Morse, where
// ABBA... and BAAB... of length  $2^{10}$  hash the same mod  $2^{64}$ ).
// "typedef ull H;" instead if you think test data is random,
// or work mod  $10^{9}+7$  if the Birthday paradox is not a problem.
typedef uint64_t ull;
struct H {
    ull x; H(ull x=0) : x(x) {}
    H operator+(H o) { return x + o.x + (x + o.x < x); }
    H operator-(H o) { return *this + ~o.x; }
    H operator*(H o) { auto m = (_uint128_t)x * o.x;
        return H((ull)m + (ull)(m >> 64)); }
    ull get() const { return x + !~x; }
    bool operator==(H o) const { return get() == o.get(); }
    bool operator<(H o) const { return get() < o.get(); }
};
static const H C = (ll)1e11+3; // (order ~ 3e9; random also ok)

struct HashInterval {
    vector<H> ha, pw;
    HashInterval(string& str) : ha(sz(str)+1), pw(ha) {
        pw[0] = 1;
        rep(i,0,sz(str))
            ha[i+1] = ha[i] * C + str[i],
            pw[i+1] = pw[i] * C;
    }
    H hashInterval(int a, int b) { // hash [a, b)
}

```

```

        return ha[b] - ha[a] * pw[b - a];
    }

vector<H> getHashes(string& str, int length) {
    if (sz(str) < length) return {};
    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;}
```

**Trie.h**  
**Description:** Trie implementation for binary strings  
**Time:**  $\mathcal{O}(N)$  per operation where N is string length

```
e67de7, 48 lines
```

```

struct TrieNode {
    TrieNode* child[2];
    int cnt, id;

    TrieNode() {
        cnt = id = 0;
        for(int i = 0; i < 2; ++i) child[i] = nullptr;
    }
};

TrieNode* root = new TrieNode();

void add_string(const string &s, int id) {
    TrieNode* p = root;
    for(auto c: s) {
        int nxt = c - '0';
        if(p->child[nxt] == nullptr) p->child[nxt] = new TrieNode();
        p = p->child[nxt];
        p->cnt++;
    }
    p->id = id;
}

bool find_string(const string &s) {
    TrieNode* p = root;
    for(auto c: s) {
        int nxt = c - '0';
        if(p->child[nxt] == nullptr) return false;
        p = p->child[nxt];
    }
    return true;
}

bool del_string(TrieNode* p, const string &s, int pos) {
    if (pos != (int)s.size()) {
        int c = s[pos] - '0';
        bool is_deleted = del_string(p->child[c], s, pos + 1);
        if(is_deleted) p->child[c] = nullptr;
    }
    if(p != root) {
        p->cnt--;
        if(p->cnt == 0) {
            delete(p);
            return true;
        }
    }
}
```

```

        return false;
    }



## Various (12)



### 12.1 Intervals



IntervalContainer.h



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



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



```
edce47, 23 lines
```



```

set<pii>::iterator addInterval(set<pii>& is, int L, int R) {
    if (L == R) return is.end();
    auto it = is.lower_bound({L, R}), before = it;
    while (it != is.end() && it->first <= R) {
        R = max(R, it->second);
        before = it = is.erase(it);
    }
    if (it != is.begin() && (--it)->second >= L) {
        L = min(L, it->first);
        R = max(R, it->second);
        is.erase(it);
    }
    return is.insert(before, {L, R});
}

void removeInterval(set<pii>& is, int L, int R) {
    if (L == R) return;
    auto it = addInterval(is, L, R);
    auto r2 = it->second;
    if (it->first == L) is.erase(it);
    else (int)it->second = L;
    if (R != r2) is.emplace(R, r2);
}
```



IntervalCover.h



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



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



```
9e9d8d, 19 lines
```



```

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



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];});  
**Time:**  $\mathcal{O}(k \log \frac{n}{k})$

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

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

### 12.2 Misc. algorithms

**TernarySearch.h**

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

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

**9155b4, 11 lines**

```

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

**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  $[0, 2b)$ .

**751a02, 8 lines**

```

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

**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 = gc()) >= 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*) {}

```

**SmallPtr.h**

**Description:** A 32-bit pointer that points into BumpAllocator memory.

"BumpAllocator.h" 2dd6e9, 10 lines

```

template<class T> struct ptr {
    unsigned ind;
    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&lt;vector&lt;int, small&lt;int&gt;&gt;&gt; ed(N); bb66d4, 14 lines

```

char buf[450 << 20] alignas(16);
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) {}
};

```

**Unrolling.h**

520e76, 5 lines

```

#define F {...; ++i;}
int i = from;
while (i&3 && i < to) F // for alignment, if needed
while (i + 4 <= to) { F F F F }
while (i < to) F

```

**SIMD.h**

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

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/pd|pd)(z?y:x), movemask epi8 (hibits of bytes)
// i32gather epi32(addr, x, 4): map addr[] over 32-b parts of x
// sad epi8: 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, 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_epli8(x, y) takes a vector instead of an imm

```

// Methods that work with most data types (append e.g. `_epi32`):  
// set1, blend (i8?x:y), add, adds (sat.), mullo, sub, and/or,  
// andnot, abs, min, max, sign(1,x), cmp(gt/eq), unpack(lo/hi)

```

int sumi32(mi m) { union {int v[8]; mi m;} u; u.m = m;
    int ret = 0; rep(i,0,8) ret += u.v[i]; return ret; }
mi zero() { return _mm256_setzero_si256(); }
mi one() { return _mm256_set1_epi32(-1); }
bool all_zero(mi m) { return _mm256_testz_si256(m, m); }
bool all_one(mi m) { return _mm256_testc_si256(m, one()); }

```

```

ll example_filteredDotProduct(int n, short* a, short* b) {
    int i = 0; ll r = 0;
    mi zero = _mm256_setzero_si256(), acc = zero;
    while (i + 16 <= n) {
        mi va = L(a[i]), vb = L(b[i]); i += 16;
        va = _mm256_and_si256(_mm256_cmplt_epi16(vb, va), va);
        mi vp = _mm256_madd_epi16(va, vb);
        acc = _mm256_add_epi64(_mm256_unpacklo_epi32(vp, zero),
            _mm256_add_epi64(acc, _mm256_unpackhi_epi32(vp, zero)));
    }
    union {ll v[4]; mi m;} u; u.m = acc; rep(i,0,4) r += u.v[i];
    for (;i<n;++i) if (a[i] < b[i]) r += a[i]*b[i]; // <- equiv
    return r;
}

```

# Techniques (A)

## techniques.txt

159 lines

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

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

Knuth-Morris-Pratt  
 Tries  
 Rolling polynomial hashes  
 Suffix array  
 Suffix tree  
 Aho-Corasick  
 Manacher's algorithm  
 Letter position lists  
 Combinatorial search  
     Meet in the middle  
     Brute-force with pruning  
     Best-first (A\*)  
     Bidirectional search  
     Iterative deepening DFS / A\*

Data structures  
     LCA ( $2^k$ -jumps in trees in general)  
     Pull/push-technique on trees  
     Heavy-light decomposition  
     Centroid decomposition  
     Lazy propagation  
     Self-balancing trees  
     Convex hull trick ([wcipeg.com/wiki/Convex\\_hull\\_trick](http://wcipeg.com/wiki/Convex_hull_trick))  
     Monotone queues / monotone stacks / sliding queues  
     Sliding queue using 2 stacks  
     Persistent segment tree