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## Contest (1)

.bashrc	3 lines
alias c='g++ -Wall -Wconversion -Wfatal-errors -g - -fsanitize=undefined,address'	-std=c++17 \
xmodmap -e 'clear lock' -e 'keycode 66=less greate	r' # $caps = \Leftrightarrow$

hash.sh # Hashes a file, ignoring all whitespace and comments. Use for

# verifying that code was correctly typed.

To make executable, run the command: chmod +x hash.sh

To execute: ./hash.sh < file.cpp

cpp -dD -P -fpreprocessed | tr -d '[:space:]' | md5sum |cut -c-6

## Mathematics (2)

## 2.1 Equations

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

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

$$ax + by = e$$

$$cx + dy = f$$

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

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

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

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

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

### 2.2 Recurrences

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

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

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

## 2.3 Trigonometry

$$\sin(v + w) = \sin v \cos w + \cos v \sin w$$
$$\cos(v + w) = \cos v \cos w - \sin v \sin w$$

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

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

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

$$a\cos x + b\sin x = r\cos(x - \phi)$$
  
$$a\sin x + b\cos x = r\sin(x + \phi)$$

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

## 2.4 Derivatives/Integrals

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

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

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

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

Integration by parts:

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

### 2.5 Sums

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

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

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

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

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

#### 2.6 Series

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

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

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

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

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

### 2.7 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

Expectation is linear:

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

instead be integrals with  $p_X(x)$  replaced by  $f_X(x)$ .

For independent X and Y,

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

### 2.7.1 Discrete distributions Binomial distribution

The number of successes in n independent yes/no experiments, each which yields success with probability p is

Bin
$$(n, p)$$
,  $n = 1, 2, ..., 0 \le p \le 1$ .

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

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

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

### .bashrc hash OrderStatisticTree HashMap

### First success distribution

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

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

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

### Poisson distribution

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

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

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

### 2.7.2 Continuous distributions Uniform distribution

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

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

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

### Exponential distribution

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

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

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

### Normal distribution

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

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

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

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

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

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

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

A Markov chain is an A-chain if the states can be partitioned into two sets 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  $t_i = 1 + \sum_{k \in \mathbf{G}} p_{ki} t_k$ .

## Geometry

### 2.9.1 Triangles

Side lengths: a, b, c

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

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

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

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

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

Length of bisector (divides angles in two):

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

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

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

### 2.9.2 Quadrilaterals

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

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

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

### 2.9.3 Spherical coordinates



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

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

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

## Data structures (3)

OrderStatisticTree.h

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

#include <bits/extc++.h> using namespace \_\_gnu\_pbds; template<class T> using Tree = tree<T, null\_type, less<T>, rb\_tree\_tag, tree order statistics node update>; void example() { Tree<int> t, t2; t.insert(8); auto it = t.insert(10).first; assert(it == t.lower bound(9)); assert(t.order\_of\_key(10) == 1); assert(t.order\_of\_key(11) == 2); assert(\*t.find\_by\_order(0) == 8); t.join(t2); // assuming T < T2 or T > T2, merge t2 into t

### HashMap.h

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

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

```
__gnu_pbds::gp_hash_table<11,int,chash> h({},{},{},{},{1<<16});
```

### SegmentTree.h

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

Time:  $\mathcal{O}(\log N)$ 0f4bdb, 19 lines struct Tree { typedef int T; static constexpr T unit = INT MIN; T f(T a, T b) { return max(a, b); } // (any associative fn) vector<T> s; int n; Tree(int n = 0, T def = unit) : s(2\*n, def), n(n) {} void update(int pos, T val) { for (s[pos += n] = val; pos /= 2;) s[pos] = f(s[pos \* 2], s[pos \* 2 + 1]);T query (int b, int e) { // query [b, e)T ra = unit, rb = unit; for  $(b += n, e += n; b < e; b /= 2, e /= 2) {$ if (b % 2) ra = f(ra, s[b++]); if (e % 2) rb = f(s[--e], rb);

### LazySegmentTree.h

};

return f(ra, rb);

**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, 50 lines
const int inf = 1e9;
struct Node {
 Node *1 = 0, *r = 0;
 int lo, hi, mset = inf, madd = 0, val = -inf;
 Node (int lo, int hi):lo(lo), hi(hi){} // Large interval of -inf
 Node (vi& v, int lo, int hi) : lo(lo), hi(hi) {
   if (lo + 1 < hi) {
     int mid = 10 + (hi - 10)/2;
     1 = new Node(v, lo, mid); r = new Node(v, mid, hi);
     val = max(1->val, r->val);
    else val = v[lo];
  int query(int L, int R) {
    if (R <= lo || hi <= L) return -inf;
    if (L <= lo && hi <= R) return val;
   return max(1->query(L, R), r->query(L, R));
  void set(int L, int R, int x) {
    if (R <= lo || hi <= L) return;
    if (L <= lo && hi <= R) mset = val = x, madd = 0;
     push(), l\rightarrow set(L, R, x), r\rightarrow set(L, R, x);
     val = max(1->val, r->val);
  void add(int L, int R, int x) {
   if (R <= lo || hi <= L) return;
    if (L <= lo && hi <= R) {
     if (mset != inf) mset += x;
     else madd += x;
     val += x;
```

```
else {
    push(), 1->add(L, R, x), r->add(L, R, x);
    val = max(1->val, r->val);
}

void push() {
    if (!1) {
        int mid = lo + (hi - lo)/2;
        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;
}
};
```

#### Wavelet.h

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

```
11aee1, 38 lines
struct WaveletTree {
 int n; vvi bv; // vector<BitVector> bv;
 WaveletTree(vl a, ll max val):
   n(sz(a)), bv(1+__lq(max_val), \{\{\}\}) {
    vl nxt(n);
    for (int h = sz(bv); h--;) {
      vector<bool> b(n);
     rep(i, 0, n) b[i] = ((a[i] >> h) & 1); bv[h] = vi(n+1); // bv[h] = b;
      rep(i, 0, n) bv[h][i+1] = bv[h][i] + !b[i]; // delete
      array it{begin(nxt), begin(nxt) + bv[h][n]};
      rep(i, 0, n) * it[b[i]] ++ = a[i];
      swap(a, nxt);
 11 kth(int 1, int r, int k) {
    11 \text{ res} = 0;
    for (int h = sz(bv); h--:) {
      int 10 = bv[h][1], r0 = bv[h][r];
      if (k < r0 - 10) 1 = 10, r = r0;
        k -= r0 - 10, res |= 1ULL << h,
          1 += bv[h][n] - 10, r += bv[h][n] - r0;
    return res;
 int count(int 1, int r, 11 ub) {
    int res = 0:
    for (int h = sz(bv); h--;) {
     int 10 = bv[h][1], r0 = bv[h][r];
      if ((\sim ub >> h) \& 1) 1 = 10, r = r0;
       res += r0 - 10, 1 += bv[h][n] - 10,
          r += bv[h][n] - r0;
   return res;
};
```

### BitVector.h

**Description:** Given vector of bits, counts number of 0's in [0, r). Use with WaveletTree.h by using modifications in comments in that file and replacing bv[h][x] with bv[h].cnt0(x)

```
Time: \mathcal{O}(1) time
```

struct BitVector {

```
vector<pair<11, int>> b;
BitVector(vector<bool> a): b(sz(a) / 64 + 1) {
    rep(i, 0, sz(a))
       b[i >> 6].first |= ll(a[i]) << (i & 63);
    rep(i, 0, sz(b)-1)
       b[i + 1].second = __builtin_popcountll(b[i].first)
            + b[i].second;
}
int cnt0(int r) {
    auto [x, y] = b[r >> 6];
    return r - y
            - __builtin_popcountll(x & ((1ULL << (r & 63)) - 1));
}
};</pre>
```

#### PST.h

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

7ddad1, 41 lines

de4ad0, 21 lines

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

#### UnionFindRollback.h

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

```
Usage: int t = uf.time(); ...; uf.rollback(t); Time: O(\log(N))
```

```
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;
};
```

### MonoRange.h

```
\label{eq:Description: when cmp = less(): a[le[i]] < a[i] >= a[ri[i]] \\ \textbf{Usage: vi le = mono.st(a, less()),} \\ \textit{ri = mono.range(le);} \\ \textit{less_equal(), greater(), greater_equal()} \\ \textbf{Time: } \mathcal{O}\left(N\right).
```

191698, 16 lines

#### CountRect.h

 $\label{eq:Description: cnt[i][j] = number of times an i-by-j sub rectangle appears such that all i*j cells $ARE 1. cnt[i][0], cnt[0][j]$ are garbage}$ 

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

71b256, 22 lines

```
vector<vi> count_rectangles(
  const vector<vector<bool>>&grid)
    int n = sz(qrid), m = sz(qrid[0]);
   vector < vi > cnt(n + 1, vi(m + 1, 0));
    for ( const auto &row : grid) {
        transform(all(h), begin(row), begin(h),
        [](int a, bool g) { return g * (a + 1); });
       vi le ( mono_st(h,less())), r(mono_range(le));
        rep(j,0,m) {
            int cnt_1 = j - le[j] - 1, cnt_r = r[j] - j - 1;
            cnt[h[j]][cnt_l + cnt_r + 1]++;
            cnt[h[j]][cnt_l]--;
            cnt[h[j]][cnt_r]--;
    rep(i,1,n+1) rep(k,0,2) for (int j = m; j > 1; j--)
        cnt[i][j - 1] += cnt[i][j];
    for (int i = n ; i > 1; i--)
        rep(j, 1, m + 1) cnt[i - 1][j] += cnt[i][j];
```

```
return cnt;
```

### Lichao.h

**Description:** min Li-chao tree allows for range add of arbitary functions such that any two functions only occur atmost once.

```
Usage:
                inc-inc, implicit, works with negative indices,
O(log(n)) query
flip signs in update and modify query to change to max. 1eac23, 37 lines
struct func {
   11 A,B;
    func(11 A, 11 B): A(A), B(B) {}
    11 operator()(11 x) { return (A*x + B); }
const func idem = {0,(11)8e18}; //idempotent, change for max
struct node {
    int lo, md, hi;
    func f = idem;
    node *left = nullptr, *right = nullptr;
    node(int lo, int hi): lo(lo), hi(hi), md(lo+(hi-lo)/2) {}
    void check(){
        if(left) return;
        left = new node(lo,md);
        right = new node (md+1, hi);
    void update(func e) { //flip signs for max
        if(e(md) < f(md)) swap(e, f);
        if(lo == hi) return;
        if(e(lo) > f(lo) \&\& e(hi) > f(hi)) return;
        check();
        if(e(lo) < f(lo)) left->update(e);
        else right->update(e);
    void rangeUpdate(int L, int R, func e) { //[]
        if(R < lo || hi < L) return;
        if (L <= lo && hi <= R) return update(e);
        check();
        left->rangeUpdate(L, R, e);
        right->rangeUpdate(L, R,e);
    11 query(int x) { //change to max if needed
        if(lo == hi) return f(x); check();
        if(x <= md) return min(f(x), left->query(x));
        return min(f(x), right->query(x));
```

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

```
struct Line {
    mutable l1 k, m, p;
    bool operator<(const Line& o) const { return k < o.k; }
    bool operator<(l1 x) const { return p < x; }
};

struct LineContainer : multiset<Line, less<>> {
    // (for doubles, use inf = 1/.0, div(a,b) = a/b)
    static const l1 inf = LLONG_MAX;
    l1 div(l1 a, l1 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(11 k, 11 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(11 x) {
  assert(!empty());
  auto 1 = *lower_bound(x);
  return 1.k * x + 1.m;
}
};
```

#### Treap.h

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

635edf, 41 lines

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

### PQupdate.h

**Description:** T: value/update type. DS: Stores T. Same semantics as std::priority\_queue.

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

35a7d2, 36 lines

### FenwickTree FenwickTree2d RMQ MoQueries Point

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

### FenwickTree.h

**Description:** Computes partial sums a[0] + a[1] + ... + a[pos - 1], and updates single elements a[i], taking the difference between the old and new 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, 11 dif) { // a[pos] \neq = dif
    for (; pos < sz(s); pos |= pos + 1) s[pos] += dif;
  11 query(int pos) { // sum of values in [0, pos)
    11 \text{ res} = 0;
    for (; pos > 0; pos &= pos - 1) res += s[pos-1];
    return res;
  int lower_bound(ll sum) \{// min \ pos \ st \ sum \ of \ [0, \ pos] >= sum
    // Returns n if no sum is >= sum, or -1 if empty sum is.
    if (sum <= 0) return -1;
    int pos = 0;
    for (int pw = 1 << 25; pw; pw >>= 1) {
     if (pos + pw \le sz(s) \&\& s[pos + pw-1] \le sum)
        pos += pw, sum -= s[pos-1];
    return pos;
};
```

### FenwickTree2d.h

**Description:** Computes sums a[i,j] for all i<I, j<J, and increases single elements a[i,j]. Requires that the elements to be updated are known in advance (call fakeUpdate() before init()).

**Time:**  $\mathcal{O}(\log^2 N)$ . (Use persistent segment trees for  $\mathcal{O}(\log N)$ .)

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

### RMQ.h

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

Usage: RMQ rmq(values);

rmg.query(inclusive, exclusive);

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

```
510c32, 16 lines
template<class T>
struct RMO {
 vector<vector<T>> jmp;
 RMQ(const vector<T>& V) : jmp(1, V) {
   for (int pw = 1, k = 1; pw * 2 <= sz(V); pw *= 2, ++k) {
      jmp.emplace_back(sz(V) - pw * 2 + 1);
     rep(j,0,sz(jmp[k]))
        jmp[k][j] = min(jmp[k - 1][j], jmp[k - 1][j + pw]);
 T query(int a, int b) {
   assert (a < b); // or return inf if a == b
   int dep = 31 - __builtin_clz(b - a);
   return min(jmp[dep][a], jmp[dep][b - (1 << dep)]);</pre>
};
```

### MoQueries.h

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

```
a12ef4, 49 lines
void add(int ind, int end) { ... } // add a[ind] (end = 0 or 1)
void del(int ind, int end) { ... } // remove a[ind]
int calc() { ... } // compute current answer
vi mo(vector<pii> Q) {
 int L = 0, R = 0, blk = 350; // \sim N/sqrt(Q)
 vi s(sz(0)), res = s;
#define K(x) pii(x.first/blk, x.second ^ -(x.first/blk & 1))
 iota(all(s), 0);
```

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

## Geometry (4)

### Point.h

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

```
template \langle \text{class T} \rangle int \text{sgn}(\text{T x}) \{ \text{return } (\text{x} > 0) - (\text{x} < 0); \}
template<class T>
struct Point {
 typedef Point P;
 T x, v;
  explicit Point (T x=0, T y=0) : x(x), y(y) {}
  bool operator<(P p) const { return tie(x,y) < tie(p.x,p.y); }</pre>
  bool operator==(P p) const { return tie(x,y)==tie(p.x,p.y); }
  P operator+(P p) const { return P(x+p.x, y+p.y); }
  P operator-(P p) const { return P(x-p.x, y-p.y); }
  P operator*(T d) const { return P(x*d, y*d); }
 P operator/(T d) const { return P(x/d, y/d); }
 T dot(P p) const { return x*p.x + y*p.y; }
 T cross(P p) const { return x*p.y - y*p.x; }
 T cross(P a, P b) const { return (a-*this).cross(b-*this); }
 T dist2() const { return x*x + y*y; }
 double dist() const { return sqrt((double)dist2()); }
  // angle to x-axis in interval [-pi, pi]
  double angle() const { return atan2(y, x); }
 P unit() const { return *this/dist(); } // makes dist()=1
 P perp() const { return P(-y, x); } // rotates +90 degrees
```

1ff9f1, 11 lines

```
P normal() const { return perp().unit(); }
// returns point rotated 'a' radians ccw around the origin
P rotate(double a) const {
 return P(x*cos(a)-y*sin(a),x*sin(a)+y*cos(a)); }
friend ostream& operator<<(ostream& os, P p) {</pre>
  return os << "(" << p.x << "," << p.v << ")"; }
```

### 4.1 Lines and Segments

#### sideOf.h

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

```
Usage: bool left = sideOf(p1,p2,q)==1;
"Point.h"
                                                       3af81c, 9 lines
template<class P>
int sideOf(P s, P e, P p) { return sgn(s.cross(e, p)); }
template<class P>
int sideOf(const P& s, const P& e, const P& p, double eps) {
  auto a = (e-s).cross(p-s);
 double 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 (segDist(s,e,p) <=epsilon) instead when using Point < double >.

```
c597e8, 3 lines
template < class P > bool on Segment (P s, P e, P p) {
 return p.cross(s, e) == 0 \&\& (s - p).dot(e - p) <= 0;
```

### lineIntersection.h

#### Description:

If a unique intersection point of the lines going through s1,e1 and s2,e2 exists {1, point} is returned. If no intersection point exists  $\{0, (0,0)\}$  is returned and if infinitely many exists  $\{-1, e^{i}\}$ (0.0) is returned. The wrong position will be returned if P is Point<|l> and the intersection point does not have integer coordinates. Products of three coordinates are used in inter- 1 mediate steps so watch out for overflow if using int or ll. Usage: auto res = lineInter(s1,e1,s2,e2);



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

### SegmentIntersection.h

return  $\{1, (s1 * p + e1 * q) / d\};$ 

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

```
Usage: vector<P> inter = segInter(s1,e1,s2,e2);
if (sz(inter) == 1)
cout << "segments intersect at " << inter[0] << endl;</pre>
"Point.h", "OnSegment.h"
template < class P > vector < P > seqInter(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) };
  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)};
```

#### 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.

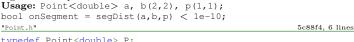


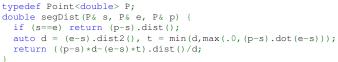
b4c5ca, 4 lines template<class P> double lineDist(const P& a, const P& b, const P& p) { return (b-a).cross(p-a)/(b-a).dist();

#### SegmentDistance.h

#### Description:

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





### 4.2 Polygons

#### PolygonArea.h

Description: Returns twice the signed area of a polygon. Clockwise enumeration gives negative area. Watch out for overflow if using int as T!

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

### InsidePolygon.h

"Point.h", "OnSegment.h"

**Description:** Returns 0 if the point is outside the polygon, 1 if it is strictly inside the polygon, and 2 if it is on the polygon.

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

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

### ConvexHull.h

#### Description:

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



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

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

#### HullDiameter.h

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

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

```
"Point.h"
                                                      c571b8, 12 lines
typedef Point<ll> P;
array<P, 2> hullDiameter(vector<P> S) {
 int n = sz(S), j = n < 2 ? 0 : 1;
 pair<11, array<P, 2>> res({0, {S[0], S[0]}});
  rep(i,0,j)
    for (;; j = (j + 1) % n) {
      res = \max(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;
```

### hullTangents.h

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

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

```
"Point.h"
                                                     53d067, 16 lines
#define cmp(i, j) p.cross(h[i], h[j == n ? 0 : j]) * (R ? 1 :
template<bool R, class P> int getTangent(vector<P>& h, P p) {
 int n = sz(h), lo = 0, hi = n - 1, md;
 if (cmp(0, 1) >= R && cmp(0, n - 1) >= !R) return 0;
  while (md = (lo + hi + 1) / 2, lo < hi) {
   auto a = cmp(md, md + 1), b = cmp(md, lo);
   if (a \ge R \&\& cmp(md, md - 1) \ge !R) return md;
   if (cmp(lo, lo + 1) < R)
     a < R\&\& b >= 0 ? lo = md : hi = md - 1;
   else a < R \mid | b <= 0 ? lo = md : hi = md - 1;
  return -1; // point strictly inside hull
template<class P> pii hullTangents(vector<P>& h, P p) {
  return {getTangent<0>(h, p), getTangent<1>(h, p)};
```

### inHull.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:  $\mathcal{O}(\log N)$ 

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

#### LineHullIntersection.h

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

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

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

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

### 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(polv)) {
   P cur = poly[i], prev = i ? poly[i-1] : poly.back();
    bool side = s.cross(e, cur) < 0;</pre>
    if (side != (s.cross(e, prev) < 0))</pre>
      res.push_back(lineInter(s, e, cur, prev).second);
    if (side)
      res.push back(cur);
```

### halfplaneIntersection.h

return res;

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

```
e9fe62, 42 lines
const double eps = 1e-8;
typedef Point < double > P;
struct HalfPlane {
 P s, e, d;
 HalfPlane(P s = P(), P e = P()): s(s), e(e), d(e - s) {}
 bool contains(P p) { return d.cross(p - s) > -eps; }
 bool operator<(HalfPlane hp) {</pre>
   if(abs(d.x) < eps && abs(hp.d.x) < eps)
     return d.y > 0 && hp.d.y < 0;
   bool side = d.x < eps \mid \mid (abs(d.x) <= eps && d.y > 0);
   bool sideHp = hp.d.x < eps || (abs(hp.d.x) \le eps \&\& hp.d.y
         > 0);
   if(side != sideHp) return side;
   return d.cross(hp.d) > 0;
```

```
P inter(HalfPlane hp) {
    auto p = hp.s.cross(e, hp.e), q = hp.s.cross(hp.e, s);
    return (s * p + e * q) / d.cross(hp.d);
};
vector<P> hpIntersection(vector<HalfPlane> hps) {
 sort(all(hps));
 int n = sz(hps), 1 = 1, r = 0;
 vector<HalfPlane> dq(n+1);
 rep(i, 0, n) {
    while (1 < r \&\& !hps[i].contains(dg[r].inter(dg[r-1]))) r--;
    while (1 < r \&\& !hps[i].contains(dq[l].inter(dq[l+1]))) l++;
    dq[++r] = hps[i];
    if(1 < r \&\& abs(dq[r].d.cross(dq[r-1].d)) < eps) {
      if(dq[r].d.dot(dq[r-1].d) < 0) return {};
      if(dq[--r].contains(hps[i].s)) dq[r] = hps[i];
  while (1 < r - 1 \&\& !dq[1].contains(dq[r].inter(dq[r-1]))) r
  while (1 < r - 1 \&\& !dq[r].contains(dq[l].inter(dq[l+1]))) 1
  if(1 > r - 2) return {};
  vector<P> poly;
  rep(i, 1, r)
   poly.push_back(dq[i].inter(dq[i+1]));
  poly.push_back(dq[r].inter(dq[l]));
  return poly;
```

#### centerOfMass.h

**Description:** Returns the center of mass for a polygon. Memory:  $\mathcal{O}(1)$ 

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

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

#### minkowskiSum.h

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

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

### 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:**  $O(N^2)$ , where N is the total number of points

```
"Point.h", "sideOf.h"
                                                      3931c<u>6</u>, 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), sqn(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);
    sort (all (segs));
    for (auto\& s : seqs) s.first = min(max(s.first, 0.0), 1.0);
    double sum = 0;
    int cnt = segs[0].second;
    rep(i,1,sz(segs)) {
     if (!cnt) sum += segs[j].first - segs[j - 1].first;
     cnt += seqs[j].second;
    ret += A.cross(B) * sum;
  return ret / 2;
```

### 4.3 Circles

# circumcircle.h Description:

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



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

### CircleIntersection.h

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

### CirclePolygonIntersection.h

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

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

```
"../../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, g) * r2;
    auto s = max(0., -a-sqrt(det)), t = min(1., -a+sqrt(det));
    if (t < 0 \mid | 1 \le s) return arg(p, q) * r2;
   Pu = p + d * s, v = p + d * t;
   return arg(p,u) * r2 + u.cross(v)/2 + arg(v,g) * r2;
  };
  rep(i, 0, sz(ps))
    sum += tri(ps[i] - c, ps[(i + 1) % sz(ps)] - c);
  return sum;
```

### 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.

```
out.push_back({c1 + v * r1, c2 + v * r2});
}
if (h2 == 0) out.pop_back();
return out;
}
```

### MinimumEnclosingCircle.h

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

```
recircumcircle.h"

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};
}
```

### 4.4 3D Geometry

### Point3D.h

**Description:** Class to handle points in 3D space. T can be e.g. double or long long.  ${}_{8058ae,\ 32\ lines}$ 

```
template<class T> struct Point3D {
 typedef Point3D P:
 typedef const P& R;
 T x, y, z;
 explicit Point3D(T x=0, T y=0, T z=0) : x(x), y(y), z(z) {}
 bool operator<(R p) const {
   return tie(x, y, z) < tie(p.x, p.y, p.z); }
 bool operator == (R p) const {
   return tie(x, y, z) == tie(p.x, p.y, p.z); }
 P operator+(R p) const { return P(x+p.x, y+p.y, z+p.z); }
 P operator-(R p) const { return P(x-p.x, y-p.y, z-p.z); }
 P operator*(T d) const { return P(x*d, y*d, z*d); }
 P operator/(T d) const { return P(x/d, y/d, z/d); }
 T dot(R p) const { return x*p.x + y*p.y + z*p.z; }
 P cross(R p) const {
   return P(y*p.z - z*p.y, z*p.x - x*p.z, x*p.y - y*p.x);
 T dist2() const { return x*x + y*y + z*z; }
 double dist() const { return sqrt((double)dist2()); }
  //Azimuthal angle (longitude) to x-axis in interval [-pi, pi]
 double phi() const { return atan2(y, x); }
  //Zenith angle (latitude) to the z-axis in interval [0, pi]
 double theta() const { return atan2(sgrt(x*x+y*y),z); }
 P unit() const { return *this/(T) dist(); } //makes dist()=1
  //returns unit vector normal to *this and p
 P normal(P p) const { return cross(p).unit(); }
  //returns point rotated 'angle' radians ccw around axis
 P rotate(double angle, P axis) const {
   double s = sin(angle), c = cos(angle); P u = axis.unit();
   return u*dot(u)*(1-c) + (*this)*c - cross(u)*s;
};
```

#### 3dHull.h

UCF

Description: Computes all faces of the 3-dimension hull of a point set. \*No four points must be coplanar\*, or else random results will be returned. All faces will point outwards.

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

```
"Point3D.h"
                                                     928b1f, 33 lines
typedef Point3D<double> P:
const double eps = 1e-6;
vector<array<int, 3>> convex_shell(vector<P> &p) {
 int n = sz(p);
  if(n < 3) return {};
  vector<array<int, 3>> faces;
  vvi active(n, vi(n, false));
  auto add_face = [&] (int a, int b, int c) -> void {
   faces.push_back({a, b, c});
   active[a][b] = active[b][c] = active[c][a] = true;
  add face (0, 1, 2);
  add_face(0, 2, 1);
  rep(i, 3, n) {
   vector<array<int, 3>> new_faces;
    for(auto [a, b, c]: faces)
     if((p[i] - p[a]).dot(p[a].cross(p[b], p[c])) > eps)
       active[a][b] = active[b][c] = active[c][a] = false;
      else new_faces.push_back({a, b, c});
    faces.clear();
    for(array<int, 3> f: new_faces)
     rep(j, 0, 3) if(!active[f[(j+1)%3]][f[j]])
       add_face(f[(j+1)%3], f[j], i);
    faces.insert(end(faces), all(new_faces));
 return faces;
```

### 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. 611f07, 8 lines

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

#### PolyhedronVolume.h

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

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

### 4.5 Miscellaneous

#### ClosestPair.h

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

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

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

#### FastDelaunav.h

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

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

```
typedef Point<11> P:
typedef struct Quad* Q;
typedef __int128_t lll; // (can be ll if coords are < 2e4)
P arb(LLONG_MAX, LLONG_MAX); // not equal to any other point
struct Ouad {
 Q rot, o; P p = arb; bool mark;
 P& F() { return r()->p; }
  Q& r() { return rot->rot; }
  O prev() { return rot->o->rot;
  Q next() { return r()->prev(); }
bool circ(P p, P a, P b, P c) { // is p in the circumcircle?
  111 p2 = p.dist2(), A = a.dist2()-p2,
      B = b.dist2()-p2, C = c.dist2()-p2;
  return p.cross(a,b) *C + p.cross(b,c) *A + p.cross(c,a) *B > 0;
Q makeEdge(P orig, P dest) {
  Q r = H ? H : new Quad{new Quad{new Quad{new Quad{0}}}};
  H = r -> 0; 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 = makeEdge(a->F(), b->p);
  splice(q, a->next());
  splice(q->r(), b);
  return q;
pair<Q,Q> rec(const vector<P>& s) {
 if (sz(s) \le 3)  {
```

```
Q = 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())) {
     0 t = e \rightarrow dir; \
     splice(e, e->prev()); \
     splice(e->r(), e->r()->prev()); \
     e->o = H; H = e; e = t; \setminus
 for (;;) {
   DEL(LC, base->r(), o); DEL(RC, base, prev());
   if (!valid(LC) && !valid(RC)) break;
   if (!valid(LC) || (valid(RC) && circ(H(RC), H(LC))))
     base = connect(RC, base->r());
     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 {};
 O 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); \
 g.push back(c\rightarrow r()); c = c\rightarrow next(); while (c != e);
 ADD; pts.clear();
 while (qi < sz(q)) if (!(e = q[qi++]) \rightarrow mark) ADD;
 return pts;
```

#### PlanarFaceExtraction.h

Description: Given a planar graph and where the points are, extract the set of faces that the graph makes

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

eefdf5, 88 lines

```
63f230, 39 lines
template<class P>
vector<vector<P>> extract faces(vvi adj, vector<P> pts) {
 int n = sz(pts);
 #define cmp(i) [&](int pi, int qi) -> bool { \
    P p = pts[pi] - pts[i], q = pts[qi] - pts[i]; \
    bool sideP = p.y < 0 || (p.y == 0 \&\& p.x < 0);
    bool sideQ = q.y < 0 \mid \mid (q.y == 0 && q.x < 0); 
    if(sideP != sideQ) return sideP; \
    return p.cross(q) > 0; }
  rep(i, 0, n)
    sort(all(adj[i]), cmp(i));
```

8b0e19, 21 lines

```
rep(i, 0, n) for(int j: adj[i])
 ed.emplace_back(i, j);
sort (all (ed));
auto get_idx = [&](int i, int j) -> int {
 return lower_bound(all(ed), pii(i, j))-begin(ed);
};
vector<vector<P>> faces;
vi used(sz(ed));
rep(i, 0, n) for(int j: adj[i]) {
 if (used[get_idx(i, j)])
   continue;
 used[get_idx(i, j)] = true;
  vector<P> face = {pts[i]};
  int prv = i, cur = j;
  while(cur != i) {
   face.push_back(pts[cur]);
   auto it = lower_bound(all(adj[cur]), prv, cmp(cur));
   if(it == begin(adj[cur]))
     it = end(adj[cur]);
   prv = cur, cur = *prev(it);
   used[get_idx(prv, cur)] = true;
  faces.push back(face);
#undef cmp
return faces;
```

## Graphs (5)

### 5.1 Network flow

MinCostMaxFlow.h

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

**Time:** Approximately  $\mathcal{O}\left(E^2\right)$ , actually  $\mathcal{O}\left(FS\right)$  where S is the time complexity of the SSSP alg used in find path (in this case SPFA)  $_{\text{e4f62e}, 56 \text{ lines}}$ 

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

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

### MinCostMaxFlowDijkstra.h

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

efdefd, 24 lines

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

#### Dinic h

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

```
struct Dinic {
  struct Edge {
    int to, rev;
    ll c, oc;
    ll flow() { return max(oc - c, OLL); } // if you need flows
```

```
vi lvl, ptr, q;
vector<vector<Edge>> adj;
Dinic(int n) : lvl(n), ptr(n), q(n), adj(n) {}
void addEdge(int a, int b, ll c, ll rcap = 0) {
  adj[a].push_back({b, sz(adj[b]), c, c});
  adj[b].push_back({a, sz(adj[a]) - 1, rcap, rcap});
11 dfs(int v, int t, 11 f) {
  if (v == t || !f) return f;
  for (int& i = ptr[v]; i < sz(adj[v]); i++) {</pre>
    Edge& e = adj[v][i];
    if (lvl[e.to] == lvl[v] + 1)
     if (ll p = dfs(e.to, t, min(f, e.c))) {
        e.c -= p, adj[e.to][e.rev].c += p;
        return p;
  return 0;
11 calc(int s, int t) {
  11 \text{ flow} = 0; q[0] = s;
  rep(L, 0, 31) do { // 'int L=30' maybe faster for random data
    lvl = ptr = vi(sz(q));
    int qi = 0, qe = lvl[s] = 1;
    while (qi < qe && !lvl[t]) {
      int v = q[qi++];
      for (Edge e : adj[v])
        if (!lvl[e.to] && e.c >> (30 - L))
          q[qe++] = e.to, lvl[e.to] = lvl[v] + 1;
    while (ll p = dfs(s, t, LLONG_MAX)) flow += p;
  } while (lvl[t]);
  return flow;
bool leftOfMinCut(int a) { return lvl[a] != 0; }
```

#### GlobalMinCut.h

**Description:** Find a global minimum cut in an undirected graph, as represented by an adjacency matrix. **Time:**  $\mathcal{O}\left(V^{3}\right)$ 

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

### GomoryHu.h

return best;

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

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

### MatroidIntersection.h

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

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

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

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

```
9812a7, 60 lines
"../data-structures/UnionFind.h"
struct ColorMat {
 vi cnt, clr;
 ColorMat(int n, vector<int> clr) : cnt(n), clr(clr) {}
 bool check(int x) { return !cnt[clr[x]]; }
  void add(int x) { cnt[clr[x]]++; }
  void clear() { fill(all(cnt), 0); }
struct GraphMat {
 UF uf:
  vector<array<int, 2>> e;
 GraphMat(int n, vector<array<int, 2>> e) : uf(n), e(e) {}
  bool check(int x) { return !uf.sameSet(e[x][0], e[x][1]); }
 void add(int x) { uf.join(e[x][0], e[x][1]); }
  void clear() { uf = UF(sz(uf.e)); }
template <class M1, class M2> struct MatroidIsect {
 int n:
 vector<char> iset;
 M1 m1; M2 m2;
 MatroidIsect(M1 m1, M2 m2, int n) : n(n), iset(n + 1), m1(m1)
      , m2(m2) {}
  vi solve() {
    rep(i,0,n) if (m1.check(i) && m2.check(i))
     iset[i] = true, m1.add(i), m2.add(i);
    while (augment());
   rep(i,0,n) if (iset[i]) ans.push_back(i);
   return ans;
  bool augment() {
   vector<int> frm(n, -1);
    queue<int> q({n}); // starts at dummy node
   auto fwdE = [&](int a) {
     vi ans;
     m1.clear();
     rep(v, 0, n) if (iset[v] \&\& v != a) ml.add(v);
     rep(b, 0, n) if (!iset[b] && frm[b] == -1 && m1.check(b))
```

```
ans.push back(b), frm[b] = a;
 return ans:
};
auto backE = [&](int b) {
 m2.clear();
 rep(cas, 0, 2) rep(v, 0, n)
   if ((v == b \mid | iset[v]) && (frm[v] == -1) == cas) {
     if (!m2.check(v))
       return cas ? q.push(v), frm[v] = b, v : -1;
     m2.add(v):
 return n;
};
while (!q.empty()) {
 int a = q.front(), c; q.pop();
  for (int b : fwdE(a))
   while((c = backE(b)) >= 0) if (c == n) {
     while (b != n) iset[b] ^= 1, b = frm[b];
     return true;
return false;
```

### 5.2 Matching

### hopcroftKarp.h

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

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

```
Time: \mathcal{O}\left(\sqrt{V}E\right)
                                                       f612e4, 42 lines
bool dfs(int a, int L, vector<vi>& q, vi& btoa, vi& A, vi& B) {
 if (A[a] != L) return 0;
 A[a] = -1;
 for (int b : g[a]) if (B[b] == L + 1) {
   B[b] = 0;
   if (btoa[b] == -1 \mid | dfs(btoa[b], L + 1, q, btoa, A, B))
      return btoa[b] = a, 1;
 return 0:
int hopcroftKarp(vector<vi>& g, vi& btoa) {
 int res = 0;
 vi A(q.size()), B(btoa.size()), cur, next;
 for (;;) {
   fill(all(A), 0);
   fill(all(B), 0);
    cur.clear();
    for (int a : btoa) if (a !=-1) A[a] = -1;
    rep(a, 0, sz(g)) if(A[a] == 0) cur.push_back(a);
    for (int lay = 1;; lay++) {
     bool islast = 0;
      next.clear();
      for (int a : cur) for (int b : 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); 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 : q[di])
   if (!vis[e] && find(e, g, btoa, vis)) {
     btoa[e] = di;
      return 1;
 return 0;
int dfsMatching(vector<vi>& g, vi& btoa) {
  rep(i, 0, sz(q)) {
    vis.assign(sz(btoa), 0);
    for (int j : q[i])
      if (find(j, g, btoa, vis)) {
        btoa[i] = i;
        break:
 return sz(btoa) - (int)count(all(btoa), -1);
```

#### MinimumVertexCover.h

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

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

### WeightedMatching.h

**Description:** Given a weighted bipartite graph, matches every node on the left with a node on the right such that no nodes are in two matchings and the sum of the edge weights is minimal. Takes  $\cos[N][M]$ , where  $\cos[i][j] = \cos f$  for L[i] to be matched with R[j] and returns (min cost, match), where L[i] is matched with R[match[i]]. Negate costs for max cost. Requires  $N \leq M$ . **Time:**  $\mathcal{O}(N^2M)$ 

```
pair<int, vi> hungarian(const vector<vi> &a) {
  if (a.empty()) return {0, {}};
  int n = sz(a) + 1, m = sz(a[0]) + 1;
  vi u(n), v(m), p(m), ans(n-1);
  rep(i,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[i0] = true;
     int i0 = p[j0], j1, delta = INT_MAX;
     rep(j,1,m) if (!done[j]) {
       auto cur = a[i0 - 1][j - 1] - u[i0] - v[j];
       if (cur < dist[j]) dist[j] = cur, pre[j] = j0;</pre>
       if (dist[j] < delta) delta = dist[j], j1 = j;</pre>
     rep(j,0,m) {
       if (done[j]) u[p[j]] += delta, v[j] -= delta;
       else dist[j] -= delta;
     j0 = j1;
    } while (p[j0]);
    while (j0) { // update alternating path
     int j1 = pre[j0];
     p[j0] = p[j1], j0 = j1;
  rep(j,1,m) if (p[j]) ans[p[j] - 1] = j - 1;
 return {-v[0], ans}; // min cost
```

#### GeneralMatching.h

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

```
"../numerical/MatrixInverse-mod.h"
vector<pii> generalMatching(int N, vector<pii>& ed) {
  vector<vector<ll>> mat(N, vector<ll>(N)), A;
  for (pii pa : ed) {
   int a = pa.first, b = pa.second, r = rand() % mod;
   mat[a][b] = r, mat[b][a] = (mod - r) % mod;
  int r = matInv(A = mat), M = 2*N - r, fi, 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(it,0,M/2) {
   rep(i,0,M) if (has[i])
     rep(j,i+1,M) if (A[i][j] && mat[i][j]) {
```

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

### 5.3 DFS algorithms

#### SCC h

```
Description: Finds strogly connected components in a directed graph. Usage: auto [num_sccs, scc_id] = sccs(adj);
```

scc.id[v] = id,  $0 <= id < num_sccs$  for each edge  $u \rightarrow v$ : scc.id[u] >= scc.id[v]Time:  $\mathcal{O}(E+V)$ 

```
auto sccs(const vector<vi>adj) {
    int n = sz(adj), num_sccs = 0, q = 0, s = 0;
    vi scc_id(n, -1), tin(n), st(n);
    auto dfs = [&] (auto&& self, int v) -> int {
        int low = tin[v] = ++q; st[s++] = v;
        for (int u : adj[v]) if (scc_id[u] < 0)
            low = min(low, tin[u] ?: self(self, u));
        if (tin[v] == low) {
            while (scc_id[v] < 0) scc_id[st[--s]] = num_sccs;
            num_sccs++;
        }
        return low;
    };
    rep(i,0,n) if (!tin[i]) dfs(dfs, i);
    return pair{num_sccs, scc_id};
```

### 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([a](const vi& edgelist) {...});
```

Time:  $\mathcal{O}(E+V)$ 2965e5, 33 lines vi num, st; vector<vector<pii>> ed; int Time; template<class F> int dfs(int at, int par, F& f) { int me = num[at] = ++Time, e, y, top = me; for (auto pa : ed[at]) if (pa.second != par) { tie(y, e) = pa;if (num[y]) { top = min(top, num[y]); if (num[y] < me)</pre> st.push\_back(e); } else { int si = sz(st);int up = dfs(y, e, f);

```
top = min(top, up);
   if (up == me) {
        st.push_back(e);
        f(vi(st.begin() + si, st.end()));
        st.resize(si);
    }
    else if (up < me) st.push_back(e);
    else { /* e is a bridge */ }
   }
}
return top;
}

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

#### blockvertextree.h

**Description:** articulation points and block-vertex tree self edges not allowed adj[u] += v, i; adj[v] += u, i; iscut[v] = 1 iff cut node bccid[edge id] = id, 0 <= id < numbccs returns numbccs, bccid, iscut Assumes the root node points to itself.

```
auto cuts(const auto& adj, int m) {
 int n = ssize(adj), num_bccs = 0, q = 0, s = 0;
 vector\langle int \rangle bcc id(m, -1), is cut(n), tin(n), st(m);
 auto dfs = [&](auto&& self, int v, int p) -> int {
   int low = tin[v] = ++q;
   for (auto [u, e] : adj[v]) {
   assert (v != u);
   if (e == p) continue;
   if (tin[u] < tin[v]) st[s++] = e;
   int lu = -1:
   low = min(low, tin[u] ?: (lu = self(self, u, e)));
   if (lu >= tin[v]) {
     is\_cut[v] = p >= 0 || tin[v] + 1 < tin[u];
     while (bcc_id[e] < 0) bcc_id[st[--s]] = num_bccs;</pre>
     num_bccs++;
   return low:
 for (int i = 0; i < n; i++)
   if (!tin[i]) dfs(dfs, i, -1);
  return tuple{num_bccs, bcc_id, is_cut};
  //! vector<vector<pii>>> adj(n);
  //! auto [num\_bccs, bcc\_id, is\_cut] = cuts(adj, m);
  //! auto bvt = block\_vertex\_tree(adj,
  //! num_bccs, bcc_id);
  //! vector<br/>
basic\_string < array < int, \gg >> adj(n);
  //! auto [num\_bccs, bcc\_id, is\_cut] = cuts(adj, m);
  //! auto bvt = block\_vertex\_tree(adj, num\_bccs, bcc\_id);
  //! //to loop over each unique bcc containing a node u:
  //! for (int bccid : bvt[v]) {
         bccid = n;
  //! }
  //! //to loop over each unique node inside a bcc:
  //! for (int v : bvt/bccid + n) {}
  //! [0, n) are original nodes
  //! [n, n + num_bccs) are BCC nodes
  //! @time O(n + m)
  //! @time O(n)
 auto block_vertex_tree(const auto& adj, int num_bccs,
```

### bridgetree 2sat EulerWalk DominatorTree

```
const vector<int>& bcc_id) {
  int n = ssize(adj);
  vector<basic_string<int>> bvt(n + num_bccs);
  vector<bool> vis(num_bccs);
  for (int i = 0; i < n; i++) {
    for (auto [_, e_id] : adj[i]) {
      int bccid = bcc_id[e_id];
      if (!vis[bccid]) {
        vis[bccid] = 1;
        bvt[i] += bccid + n;
        bvt[bccid + n] += i;
    }
    for (int bccid : bvt[i]) vis[bccid - n] = 0;
}
return bvt;
}</pre>
```

auto bridges (const auto& adj, int m) {

#### bridgetree.h

**Description:** bridges adj[u] += v, i; adj[v] += u, i; iscut[v] = 1 iff cut node brid[v] = id, 0 <= id < numbccs returns numbccs, bccid, iscut Assumes the root node points to itself.

int n = ssize(adj), num ccs = 0, q = 0, s = 0; vector $\langle int \rangle$  br id(n, -1), is br(m), tin(n), st(n); auto dfs = [&] (auto&& self, int v, int p) -> int { int low = tin[v] = ++q;st[s++] = v;for (auto [u, e] : adj[v]) if (e != p && br id[u] < 0)</pre> low = min(low, tin[u] ?: self(self, u, e));  $if (tin[v] == low) {$ if  $(p != -1) is_br[p] = 1;$ while (br\_id[v] < 0) br\_id[st[--s]] = num\_ccs;</pre> num ccs++; return low; for (int i = 0; i < n; i++) if (!tin[i]) dfs(dfs, i, -1); return tuple{num\_ccs, br\_id, is\_br}; //! @code

vector < vector < pii >> adj(n);

auto bridge\_tree(const auto& adj, int num\_ccs,

if (is\_br[e\_id]) tree[br\_id[i]] += br\_id[u];

vector<basic\_string<int>> tree(num\_ccs);

for (int i = 0; i < ssize(adj); i++)</pre>

for (auto [u, e\_id] : adj[i])

## 2sat.h

//! @endcode

return tree;

//! @time O(n + m)

//! @space O(n)

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

auto  $[num\_ccs, br\_id, is\_br] = bridges(adj, m);$ 

 $vector < basic\_string < array < int, \gg >> adj(n);$ 

const vector<int>& br\_id, const vector<int>& is\_br) {

auto  $[num\_ccs, br\_id, is\_br] = bridges(adj, m);$ 

//! auto  $bt = bridge\_tree(adj, num\_ccs, br\_id, is\_br);$ 

 $auto bt = bridge\_tree(adj, num\_ccs, br\_id, is\_br);$ 

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

```
struct TwoSat {
 int N:
 vector<vi> gr;
 vi values; // 0 = false, 1 = true
 TwoSat(int n = 0): N(n), gr(2*n) {}
 int addVar() { // (optional)
   gr.emplace_back();
   gr.emplace back();
   return N++;
 void either(int f, int j) {
   f = max(2*f, -1-2*f);
   j = \max(2*j, -1-2*j);
   gr[f].push_back(j^1);
   gr[j].push_back(f^1);
 void setValue(int x) { either(x, x); }
 void atMostOne(const vi& li) { // (optional)
   if (sz(li) <= 1) return;</pre>
   int cur = \simli[0];
   rep(i,2,sz(li)) {
     int next = addVar();
     either(cur, ~li[i]);
     either(cur, next);
     either(~li[i], next);
     cur = ~next;
    either(cur, ~li[1]);
 vi val, comp, z; int time = 0;
 int dfs(int i) {
   int low = val[i] = ++time, x; z.push_back(i);
    for(int e : gr[i]) if (!comp[e])
     low = min(low, val[e] ?: dfs(e));
    if (low == val[i]) do {
     x = z.back(); z.pop_back();
     comp[x] = low;
     if (values[x>>1] == -1)
       values[x>>1] = x&1;
    } while (x != i);
    return val[i] = low;
 bool solve() {
   values.assign(N, -1);
   val.assign(2*N, 0); comp = val;
   rep(i,0,2*N) if (!comp[i]) dfs(i);
   rep(i,0,N) if (comp[2*i] == comp[2*i+1]) return 0;
   return 1;
```

EulerWalk.h

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

vi eulerWalk (vector<vector<pii>>>& gr, int nedges, int src=0) {
 int n = sz(gr);
 vi D(n), its(n), eu(nedges), ret, s = {src};
 D[src]++; // to allow Euler paths, not just cycles
 while (!s.empty()) {
 int x = s.back(), y, e, &it = its[x], end = sz(gr[x]);
 if (it == end) { ret.push\_back(x); s.pop\_back(); continue; }
 tie(y, e) = gr[x][it++];
 if (!eu[e]) {
 D[x]--, D[y]++;
 eu[e] = 1; s.push\_back(y);
 }
 for (int x : D) if (x < 0 || sz(ret) != nedges+1) return {};
 return {ret.rbegin(), ret.rend()};</pre>

### DominatorTree.h

vi dom(n);

rep(i, 1, t) {

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

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

```
if(idom[i] != sdom[i]) idom[i] = idom[idom[i]];
 dom[revLabel[i]] = revLabel[idom[i]];
return dom;
```

## 5.4 Coloring

### EdgeColoring.h

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

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

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

### 5.5 Heuristics

#### MaximalCliques.h

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

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

b0d5b1, 12 lines

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

### MaximumClique.h

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

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

```
typedef vector<br/>bitset<200>> vb;
struct Maxclique {
 double limit=0.025, pk=0;
 struct Vertex { int i, d=0; };
 typedef vector<Vertex> vv;
 vv V;
 vector<vi> C;
 vi qmax, q, S, old;
 void init(vv& r) {
    for (auto\& v : r) v.d = 0;
    for (auto& v : r) for (auto j : r) v.d += e[v.i][j.i];
   sort(all(r), [](auto a, auto b) { return a.d > b.d; });
   int mxD = r[0].d;
   rep(i, 0, sz(r)) r[i].d = min(i, mxD) + 1;
 void expand(vv& R, int lev = 1) {
   S[lev] += S[lev - 1] - old[lev];
   old[lev] = S[lev - 1];
   while (sz(R)) {
     if (sz(q) + R.back().d <= sz(qmax)) return;</pre>
     g.push_back(R.back().i);
     vv T;
      for(auto v:R) if (e[R.back().i][v.i]) T.push_back({v.i});
     if (sz(T)) {
       if (S[lev]++ / ++pk < limit) init(T);</pre>
       int j = 0, mxk = 1, mnk = max(sz(qmax) - sz(q) + 1, 1);
       C[1].clear(), C[2].clear();
       for (auto v : T) {
         int k = 1:
         auto f = [&](int i) { return e[v.i][i]; };
         while (any_of(all(C[k]), f)) k++;
         if (k > mxk) mxk = k, C[mxk + 1].clear();
         if (k < mnk) T[j++].i = v.i;
         C[k].push_back(v.i);
       if (j > 0) T[j - 1].d = 0;
       rep(k, mnk, mxk + 1) for (int i : C[k])
         T[j].i = i, T[j++].d = k;
       expand(T, lev + 1);
      else if (sz(q) > sz(qmax)) qmax = q;
     q.pop_back(), R.pop_back();
 vi maxClique() { init(V), expand(V); return qmax; }
 Maxclique(vb conn): e(conn), C(sz(e)+1), S(sz(C)), old(S) {
    rep(i,0,sz(e)) V.push_back({i});
};
```

#### MaximumIndependentSet.h

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

### 5.6 Trees

#### BinaryLifting.h

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

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

bfce85, 25 lines

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

#### LCA.h

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

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

```
"../data-structures/RMQ.h"
                                                       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[rmg.query(a, b)];
  //dist(a,b){return depth[a] + depth[b] - 2*depth[lca(a,b)];}
```

#### CentroidDecomp.h

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

2c9a06, 33 lines

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

```
template <class F> struct centroid {
 vector<vi> adj;
 F f;
 vi sub sz, par;
 centroid(const vector<vi>& a_adj, F a_f)
   : adj(a_adj), f(a_f), sub_sz(sz(adj), -1), par(sz(adj), -1)
    rep(i, 0, sz(adj))
     if (sub sz[i] == -1) dfs(i);
```

### EdgeCD CompressTree HLD LinkCutTree

```
void calc_sz(int u, int p) {
    sub_sz[u] = 1;
    for (int v : adj[u])
     if (v != p)
       calc_sz(v, u), sub_sz[u] += sub_sz[v];
  int dfs(int u) {
    calc_sz(u, -1);
    for (int p = -1, sz\_root = sub\_sz[u];;) {
      auto big_ch = find_if(begin(adj[u]), end(adj[u]), [&](int
        return v != p && 2 * sub_sz[v] > sz_root;
     if (big_ch == end(adj[u])) break;
     p = u, u = *biq_ch;
    f(adj, u);
    for (int v : adj[u]) {
     iter_swap(find(begin(adj[v]), end(adj[v]), u), rbegin(adj
          [v]));
     adj[v].pop_back();
     par[dfs(v)] = u;
    return u;
};
```

### EdgeCD.h

```
Time: \mathcal{O}(n \log n)
                                                      fe3ded, 35 lines
template <class F> struct edge_cd {
 vector<vector<int>> adi;
  vector<int> sub sz:
  edge_cd(const vector<vector<int>>& a_adj, F a_f) : adj(a_adj)
      , f(a_f), sub_sz((int)size(adj)) {
   dfs(0, (int)size(adj));
  int find_cent(int u, int p, int siz) {
   sub_sz[u] = 1;
   for (int v : adj[u])
     if (v != p) {
       int cent = find_cent(v, u, siz);
       if (cent != -1) return cent;
       sub_sz[u] += sub_sz[v];
   if (p == -1) return u;
   return 2 * sub_sz[u] >= siz ? sub_sz[p] = siz - sub_sz[u],
        u : -1;
  void dfs(int u, int siz) {
   if (siz <= 2) return;
   u = find_cent(u, -1, siz);
   int sum = 0;
    auto it = partition(begin(adj[u]), end(adj[u]), [&](int v)
     bool ret = 2 * sum + sub\_sz[v] < siz - 1 && 3 * (sum + sub\_sz[v])
           sub sz[v]) \le 2 * (siz - 1);
     if (ret) sum += sub_sz[v];
     return ret;
    f(adj, u, it - begin(adj[u]));
    vector<int> oth(it, end(adj[u]));
   adj[u].erase(it, end(adj[u]));
   dfs(u, sum + 1);
   swap(adj[u], oth);
    dfs(u, siz - sum);
```

### CompressTree.h

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

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

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

### HLD.h

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

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

```
"../data-structures/LazySegmentTree.h"
                                                     6f34db, 46 lines
template <bool VALS_EDGES> struct HLD {
 int N, tim = 0;
 vector<vi> adj;
 vi par, siz, depth, rt, pos;
 Node *tree;
 HLD(vector<vi> adj )
   : N(sz(adj_)), adj(adj_), par(N, -1), siz(N, 1), depth(N),
     rt(N),pos(N),tree(new Node(0, N)) { dfsSz(0); dfsHld(0); }
 void dfsSz(int v) {
   if (par[v] != -1) adj[v].erase(find(all(adj[v]), par[v]));
   for (int& u : adj[v]) {
     par[u] = v, depth[u] = depth[v] + 1;
     dfsSz(u);
     siz[v] += siz[u];
     if (siz[u] > siz[adj[v][0]]) swap(u, adj[v][0]);
 void dfsHld(int v) {
   pos[v] = tim++;
   for (int u : adj[v]) {
     rt[u] = (u == adj[v][0] ? rt[v] : u);
     dfsHld(u);
 template <class B> void process(int u, int v, B op) {
   for (; rt[u] != rt[v]; v = par[rt[v]]) {
     if (depth[rt[u]] > depth[rt[v]]) swap(u, v);
```

```
op(pos[rt[v]], pos[v] + 1);
   if (depth[u] > depth[v]) swap(u, v);
   op(pos[u] + VALS_EDGES, pos[v] + 1);
 void modifyPath(int u, int v, int val) {
   process(u, v, [&](int 1, int r) { tree->add(1, r, val); });
 int queryPath(int u, int v) { // Modify depending on problem
   int res = -1e9:
   process(u, v, [&](int l, int r) {
       res = max(res, tree->query(1, r));
   return res;
 int querySubtree(int v) { // modifySubtree is similar
    return tree->query(pos[v] + VALS_EDGES, pos[v] + siz[v]);
};
```

#### LinkCutTree.h

Description: Represents a forest of unrooted trees. Nodes are 1-indexed. You can add and remove edges (as long as the result is still a forest). You can also do path sum, subtree sum, and LCA queries, which depend on the current root.

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

9aa6da, 105 lines

```
struct SplayTree {
 struct Node {
    int ch[2] = \{0, 0\}, p = 0;
                                   // Path aggregates
   11 \text{ self} = 0, \text{ path} = 0;
    11 \text{ sub} = 0, \text{ vir} = 0;
                                   // Subtree aggregates
                                           // Lazy tags
    bool flip = 0;
  vector<Node> T:
  SplayTree(int n) : T(n + 1) {}
  void push(int x) {
   if (!x || !T[x].flip) return;
    int 1 = T[x].ch[0], r = T[x].ch[1];
    T[1].flip ^= 1, T[r].flip ^= 1;
    swap(T[x].ch[0], T[x].ch[1]);
    T[x].flip = 0;
  void pull(int x) {
    int l = T[x].ch[0], r = T[x].ch[1]; push(l); push(r);
    T[x].path = T[1].path + T[x].self + T[r].path;
    T[x].sub = T[x].vir + T[1].sub + T[r].sub + T[x].self;
  void set(int x, int d, int y) {
    T[x].ch[d] = y; T[y].p = x; pull(x);
 void splay(int x) {
    auto dir = [&](int x) {
      int p = T[x].p; if (!p) return -1;
      return T[p].ch[0] == x ? 0 : T[p].ch[1] == x ? 1 : -1;
    auto rotate = [&](int x) {
      int y = T[x].p, z = T[y].p, dx = dir(x), dy = dir(y);
      set(y, dx, T[x].ch[!dx]);
      set(x, !dx, y);
      if (\sim dy) set (z, dy, x);
      T[x].p = z;
```

### DirectedMST Polynomial PolyRoots PolyInterpolate

```
for (push(x); \sim dir(x);) {
     int y = T[x].p, z = T[y].p;
     push(z); push(y); push(x);
     int dx = dir(x), dy = dir(y);
     if (\sim dy) rotate(dx != dy ? x : y);
      rotate(x);
};
struct LinkCut : SplayTree {
 LinkCut(int n) : SplayTree(n) {}
  int access(int x) {
   int u = x, v = 0;
    for (; u; v = u, u = T[u].p) {
     splay(u);
     int & ov = T[u].ch[1];
     T[u].vir += T[ov].sub;
     T[u].vir -= T[v].sub;
     ov = v; pull(u);
    return splay(x), v;
  void reroot(int x) {
   access(x); T[x].flip ^= 1; push(x);
  void Link(int u, int v) {
   reroot(u); access(v);
   T[v].vir += T[u].sub;
   T[u].p = v; pull(v);
  void Cut(int u, int v) {
    reroot(u); access(v);
   T[v].ch[0] = T[u].p = 0; pull(v);
  // Rooted tree LCA. Returns 0 if u and v arent connected.
  int LCA(int u, int v) {
   if (u == v) return u;
    access(u); int ret = access(v);
    return T[u].p ? ret : 0;
  // Query subtree of u where v is outside the subtree.
  11 Subtree(int u, int v) {
    reroot(v); access(u); return T[u].vir + T[u].self;
  // Query path [u..v]
  ll Path(int u, int v) {
    reroot(u); access(v); return T[v].path;
  // Update vertex u with value v
  void Update(int u, ll v) {
    access(u); T[u].self = v; pull(u);
};
```

### DirectedMST.h

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

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

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

39e620, 60 lines

```
struct Edge { int a, b; ll w; };
struct Node {
 Edge key;
 Node *1, *r;
 11 delta;
  void prop()
   kev.w += delta;
   if (1) 1->delta += delta;
   if (r) r->delta += delta;
   delta = 0;
 Edge top() { prop(); return key; }
Node *merge(Node *a, Node *b) {
 if (!a || !b) return a ?: b;
 a->prop(), b->prop();
 if (a->key.w > b->key.w) swap(a, b);
 swap(a->1, (a->r = merge(b, a->r)));
 return a;
void pop(Node*& a) { a\rightarrow prop(); a = merge(a\rightarrow 1, a\rightarrow r); }
pair<ll, vi> dmst(int n, int r, vector<Edge>& g) {
 RollbackUF uf(n);
 vector<Node*> heap(n);
 for (Edge e : g) heap[e.b] = merge(heap[e.b], new Node(e));
 11 \text{ res} = 0;
 vi seen(n, -1), path(n), par(n);
 seen[r] = r;
 vector<Edge> Q(n), in(n, \{-1,-1\}), comp;
  deque<tuple<int, int, vector<Edge>>> cycs;
 rep(s,0,n) {
   int u = s, qi = 0, w;
    while (seen[u] < 0) {</pre>
      if (!heap[u]) return {-1,{}};
      Edge e = heap[u]->top();
      heap[u]->delta -= e.w, pop(heap[u]);
      Q[qi] = e, path[qi++] = u, seen[u] = s;
      res += e.w, u = uf.find(e.a);
      if (seen[u] == s) {
        Node \star cyc = 0;
        int end = qi, time = uf.time();
        do cyc = merge(cyc, heap[w = path[--qi]]);
        while (uf.join(u, w));
        u = uf.find(u), heap[u] = cyc, seen[u] = -1;
        cycs.push_front({u, time, {&Q[qi], &Q[end]}});
    rep(i, 0, qi) in[uf.find(Q[i].b)] = Q[i];
  for (auto& [u,t,comp] : cycs) { // restore sol (optional)
   uf.rollback(t);
    Edge inEdge = in[u];
    for (auto& e : comp) in[uf.find(e.b)] = e;
    in[uf.find(inEdge.b)] = inEdge;
 rep(i,0,n) par[i] = in[i].a;
 return {res, par};
```

## Numerical Methods (6)

### 6.1 Polynomials and recurrences

### Polynomial.h

c9b7b0, 17 lines

```
struct Poly {
  vector<double> a;
  double operator() (double x) const {
    double val = 0;
    for (int i = sz(a); i--;) (val *= x) += a[i];
    return val;
}

void diff() {
    rep(i,1,sz(a)) a[i-1] = i*a[i];
    a.pop_back();
}

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

### PolyRoots.h

**Description:** Finds the real roots to a polynomial.

Usage: polyRoots({{2,-3,1}},-1e9,1e9) // solve  $x^2-3x+2=0$  | Time:  $\mathcal{O}(n^2 \log(1/\epsilon))$ 

"Polynomial.h" vector<double> polyRoots(Poly p, double xmin, double xmax) { if  $(sz(p.a) == 2) \{ return \{-p.a[0]/p.a[1]\}; \}$ vector<double> ret; Poly der = p; der.diff(); auto dr = polyRoots(der, xmin, xmax); dr.push back(xmin-1); dr.push\_back(xmax+1); sort(all(dr)); rep(i, 0, sz(dr) -1) { double l = dr[i], h = dr[i+1]; bool sign = p(1) > 0;if  $(sign ^ (p(h) > 0)) {$ rep(it, 0, 60) { // while (h - l > 1e-8)double m = (1 + h) / 2, f = p(m);  $if ((f <= 0) ^ sign) 1 = m;$ else h = m: ret.push\_back((1 + h) / 2); return ret:

#### PolyInterpolate.h

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

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

```
}
return res;
```

### BerlekampMassey.h

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

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

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

### LinearRecurrence.h

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

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

```
f4e444, 26 lines
typedef vector<ll> Polv:
11 linearRec(Poly S, Poly tr, 11 k) {
  int n = sz(tr);
  auto combine = [&] (Poly a, Poly b) {
   Poly res(n * 2 + 1);
   rep(i, 0, n+1) rep(j, 0, n+1)
     res[i + j] = (res[i + j] + a[i] * b[j]) % mod;
    for (int i = 2 * n; i > n; --i) rep(j,0,n)
     res[i - 1 - j] = (res[i - 1 - j] + res[i] * tr[j]) % mod;
   res.resize(n + 1);
   return res;
 Poly pol(n + 1), e(pol);
 pol[0] = e[1] = 1;
  for (++k; k; k /= 2) {
   if (k % 2) pol = combine(pol, e);
   e = combine(e, e);
 11 \text{ res} = 0;
 rep(i, 0, n) res = (res + pol[i + 1] * S[i]) % mod;
  return res;
```

## 6.2 Optimization

### GoldenSectionSearch.h

**Description:** Finds the argument minimizing the function f in the interval [a,b] assuming f is unimodal on the interval, i.e. has only one local minimum. The maximum error in the result is eps. Works equally well for maximization with a small change in the code. See TernarySearch.h in the Various chapter for a discrete version.

```
Usage: double func (double x) { return 4+x+.3*x*x; } double xmin = gss(-1000,1000,func);

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

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

### HillClimbing.h

Description: Poor man's optimization for unimodal functions

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

template<class F> pair<double, P> hillClimb(P start, F f) {
  pair<double, P> cur(f(start), start);
  for (double jmp = le9; jmp > le-20; jmp /= 2) {
    rep(j,0,100) rep(dx,-1,2) rep(dy,-1,2) {
        P p = cur.second;
        p[0] += dx*jmp;
        p[1] += dy*jmp;
        cur = min(cur, make_pair(f(p), p));
    }
  }
  return cur;
}
```

### IntegrateAdaptiveTyler.h

Description: Gets area under a curve

e7beba, 17 lines

### RungeKutta4.h

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

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

### Simplex.h

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

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

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

```
typedef double T; // long double, Rational, double + modeP>...
typedef vector<T> vd;
typedef vector<vd> vvd:
const T eps = 1e-8, inf = 1/.0;
#define MP make pair
#define ltj(X) if(s == -1 \mid \mid MP(X[j], N[j]) < MP(X[s], N[s])) s=j
struct LPSolver {
 int m, n;
 vi N. B:
 vvd D;
 LPSolver (const vvd& A, const vd& b, const vd& c) :
    m(sz(b)), n(sz(c)), N(n+1), B(m), D(m+2, vd(n+2)) {
      rep(i, 0, m) rep(j, 0, n) D[i][j] = A[i][j];
      rep(i,0,m) \{ B[i] = n+i; D[i][n] = -1; D[i][n+1] = b[i]; \}
      rep(j, 0, n) \{ N[j] = j; D[m][j] = -c[j]; \}
      N[n] = -1; D[m+1][n] = 1;
  void pivot(int r, int s) {
   T *a = D[r].data(), inv = 1 / a[s];
    rep(i, 0, m+2) if (i != r \&\& abs(D[i][s]) > eps) {
     T *b = D[i].data(), inv2 = b[s] * inv;
      rep(j, 0, n+2) b[j] -= a[j] * inv2;
      b[s] = a[s] * inv2;
```

rep(j,0,n+2) if (j != s) D[r][j] \*= inv; rep(i,0,m+2) if (i != r) D[i][s] \*= -inv;

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

D[r][s] = inv;

for (;;) {

swap(B[r], N[s]);

bool simplex(int phase)

int x = m + phase - 1;

```
if (D[x][s] >= -eps) return true;
      int r = -1:
      rep(i,0,m) {
       if (D[i][s] <= eps) continue;</pre>
       if (r == -1 || MP(D[i][n+1] / D[i][s], B[i])
                     < MP(D[r][n+1] / D[r][s], B[r])) r = i;
     if (r == -1) return false:
     pivot(r, s);
 T solve(vd &x) {
   int r = 0;
    rep(i,1,m) if (D[i][n+1] < D[r][n+1]) r = i;
   if (D[r][n+1] < -eps) {
     pivot(r, n);
     if (!simplex(2) || D[m+1][n+1] < -eps) return -inf;
     rep(i, 0, m) if (B[i] == -1) {
       int s = 0;
       rep(j,1,n+1) ltj(D[i]);
       pivot(i, s);
   bool ok = simplex(1); x = vd(n);
   rep(i, 0, m) if (B[i] < n) x[B[i]] = D[i][n+1];
    return ok ? D[m][n+1] : inf;
};
```

### 6.3 Matrices

#### Determinant.h

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

```
Time: \mathcal{O}\left(N^3\right) bd5cec, 15 lines double det (vector<vector<double>>& a) { int n = sz(a); double res = 1; rep(i,0,n) { int b = i; rep(j,i+1,n) if (fabs(a[j][i]) > fabs(a[b][i])) b = j; if (i != b) swap(a[i], a[b]), res *= -1; res *= a[i][i]; if (res == 0) return 0; rep(j,i+1,n) { double v = a[j][i] / a[i][i]; if (v != 0) rep(k,i+1,n) a[j][k] -= v * a[i][k]; } } return res; }
```

#### IntDeterminant.h

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

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

3313dc, 18 lines

```
const ll mod = 12345;
ll det(vector<vector<1l>>& 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:**  $\mathcal{O}\left(n^2m\right)$ 

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

#### SolveLinear2.h

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

### SolveLinearBinary.h

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

typedef bitset<1000> bs;

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

#### MatrixInverse.h

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

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

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

00ced6, 35 lines

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

#### MatrixInverse-mod.h

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

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

```
a6f68f, 36 lines
"../number-theory/ModPow.h"
int matInv(vector<vector<11>>& A) {
 int n = sz(A); vi col(n);
 vector<vector<ll>> tmp(n, vector<ll>(n));
  rep(i, 0, n) tmp[i][i] = 1, col[i] = i;
  rep(i,0,n) {
   int r = i, c = i;
   rep(j,i,n) rep(k,i,n) if (A[j][k]) {
    r = j; c = k; goto found;
   return i;
   A[i].swap(A[r]); tmp[i].swap(tmp[r]);
    rep(j,0,n) swap(A[j][i], A[j][c]), swap(tmp[j][i], tmp[j][c
    swap(col[i], col[c]);
    11 v = modpow(A[i][i], mod - 2);
    rep(j,i+1,n)
     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) {
   11 v = A[j][i];
   rep(k,0,n) tmp[j][k] = (tmp[j][k] - v*tmp[i][k]) % mod;
  rep(i,0,n) rep(j,0,n)
   A[col[i]][col[j]] = tmp[i][j] % mod + (tmp[i][j] < 0 ? mod
       : 0);
 return n;
```

### Tridiagonal.h

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

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

This is useful for solving problems on the type

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

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

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

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

8f9fa8, 26 lines

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

#### JacobianMatrix.h

Description: Makes Jacobian Matrix using finite differences 75dc90, 15 lines

```
template<class F, class T>
vector<vector<T>> makeJacobian(F &f, vector<T> &x) {
   int n = sz(x);
   vector<vector<T>> J(n, vector<T>(n));
   vector<T> fX0 = f(x);
   rep(i, 0, n) {
      x[i] += eps;
   vector<T> fX1 = f(x);
   rep(j, 0, n) {
      J[j][i] = (fX1[j] - fX0[j]) / eps;
   }
   x[i] -= eps;
   return J;
}
```

### NewtonsMethod.h

Description: Solves a system on non-linear equations

```
jacobianMatrix.h

faf945, 10 lines

template<class F, class T>

void solveNonlinear(F f, vector<T> &x) {
  int n = sz(x);
  rep(iter, 0, 100) {
    vector<vector<T>> J = makeJacobian(f, x);
    matInv(J);
    vector<T> dx = J * f(x);
    x = x - dx;
  }
}
```

### 6.4 Fourier transforms

FastFourierTransform.h

**Description:** fft(a) computes  $\hat{f}(k) = \sum_x a[x] \exp(2\pi i \cdot kx/N)$  for all k. N must be a power of 2. Useful for convolution: conv (a, b) = c, where  $c[x] = \sum_i a[i]b[x-i]$ . 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  $(\sum_i a_i^2 + \sum_i b_i^2) \log_2 N < 9 \cdot 10^{14}$  (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 \neq 2) {
    R.resize(n); rt.resize(n);
    auto x = polar(1.0L, acos(-1.0L) / k);
    rep(i,k,2*k) rt[i] = R[i] = i&1 ? R[i/2] * x : R[i/2];
  vi rev(n);
  rep(i, 0, n) \ rev[i] = (rev[i / 2] | (i \& 1) << L) / 2;
  rep(i,0,n) if (i < rev[i]) swap(a[i], a[rev[i]]);
  for (int k = 1; k < n; k *= 2)
    for (int i = 0; i < n; i += 2 * k) rep(j, 0, k) {
     Cz = rt[j+k] * a[i+j+k]; // (25\% faster if hand-rolled)
     a[i + j + k] = a[i + j] - z;
      a[i + j] += z;
vd conv(const vd& a, const vd& b) {
  if (a.empty() || b.empty()) return {};
  vd res(sz(a) + sz(b) - 1);
  int L = 32 - __builtin_clz(sz(res)), n = 1 << L;</pre>
  vector<C> in(n), out(n);
  copy(all(a), begin(in));
  rep(i,0,sz(b)) in[i].imag(b[i]);
  fft(in);
  for (C\& x : in) x *= x;
  rep(i, 0, n) out[i] = in[-i & (n - 1)] - conj(in[i]);
  fft(out);
  rep(i, 0, sz(res)) res[i] = imag(out[i]) / (4 * n);
  return res;
```

#### FastFourierTransformMod.h

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

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

```
"FastFourierTransform.h"
                                                     b82773, 22 lines
typedef vector<ll> vl;
template<int M> vl convMod(const vl &a, const vl &b) {
 if (a.empty() || b.empty()) return {};
 vl res(sz(a) + sz(b) - 1);
 int B=32-__builtin_clz(sz(res)), n=1<<B, cut=int(sqrt(M));</pre>
  vector<C> L(n), R(n), outs(n), outl(n);
  rep(i, 0, sz(a)) L[i] = C((int)a[i] / cut, (int)a[i] % cut);
  rep(i, 0, sz(b)) R[i] = C((int)b[i] / cut, (int)b[i] % cut);
 fft(L), fft(R);
 rep(i,0,n) {
   int j = -i \& (n - 1);
   outl[j] = (L[i] + conj(L[j])) * R[i] / (2.0 * n);
    outs[j] = (L[i] - conj(L[j])) * R[i] / (2.0 * n) / 1i;
 fft (outl), fft (outs);
 rep(i,0,sz(res)) {
```

```
11 \text{ av} = 11(\text{real}(\text{outl}[i]) + .5), \text{ cv} = 11(\text{imag}(\text{outs}[i]) + .5);
  11 bv = 11(imag(out1[i])+.5) + 11(real(outs[i])+.5);
  res[i] = ((av % M * cut + bv) % M * cut + cv) % M;
return res;
```

### NumberTheoreticTransform.h

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

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

```
"../number-theory/ModPow.h"
const 11 mod = (119 << 23) + 1, root = 62; // = 998244353
// For p < 2^30 there is also e.g. 5 << 25, 7 << 26, 479 << 21
// and 483 \ll 21 (same root). The last two are > 10^9.
typedef vector<ll> vl;
void ntt(vl &a) {
  int n = sz(a), L = 31 - _builtin_clz(n);
  static vl rt(2, 1);
  for (static int k = 2, s = 2; k < n; k *= 2, s++) {
   rt.resize(n);
   11 z[] = {1, modpow(root, mod >> s)};
   rep(i,k,2*k) rt[i] = rt[i / 2] * z[i & 1] % mod;
  vi rev(n);
  rep(i,0,n) \ rev[i] = (rev[i / 2] | (i \& 1) << L) / 2;
  rep(i,0,n) if (i < rev[i]) swap(a[i], a[rev[i]]);
  for (int k = 1; k < n; k *= 2)
    for (int i = 0; i < n; i += 2 * k) rep(j,0,k) {
     11 z = rt[j + k] * a[i + j + k] % mod, &ai = a[i + j];
     a[i + j + k] = ai - z + (z > ai ? mod : 0);
     ai += (ai + z >= mod ? z - mod : z);
vl conv(const vl &a, const vl &b) {
 if (a.empty() || b.empty()) return {};
  int s = sz(a) + sz(b) - 1, B = 32 - _builtin_clz(s), n = 1
  int inv = modpow(n, mod - 2);
  vl L(a), R(b), out(n);
  L.resize(n), R.resize(n);
  ntt(L), ntt(R);
  rep(i, 0, n) out [-i \& (n - 1)] = (11) L[i] * R[i] % mod * inv %
  return {out.begin(), out.begin() + s};
```

#### FastSubsetTransform.h

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

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

```
464cf3, 16 lines
void FST(vi& a, bool inv) {
  for (int n = sz(a), step = 1; step < n; step *= 2) {
    for (int i = 0; i < n; i += 2 * step) rep(j,i,i+step) {
      int \&u = a[j], \&v = a[j + step]; tie(u, v) =
        inv ? pii(v - u, u) : pii(v, u + v); // AND
       inv ? pii(v, u - v) : pii(u + v, u); // OR
                                              // XOR
        pii(u + v, u - v);
```

```
if (inv) for (int& x : a) x /= sz(a); // XOR only
vi conv(vi a, vi b) {
 FST(a, 0); FST(b, 0);
 rep(i, 0, sz(a)) a[i] *= b[i];
 FST(a, 1); return a;
```

### Minconv.h

Description: @param convex, arbitrary arrays where convex satisfies convex[i+1]- $convex[i] \le convex[i+2]$ -convex[i+1] @returns array 'res' where 'res[k]' = the min of (a[i]+b[j]) for all pairs (i,j) where i+j== $k_{633806,\ 26\ lines}$ 

```
vector<int> min plus(const vector<int>& convex,
 const vector<int>& arbitrary) {
 int n = ssize(convex);
 int m = ssize(arbitrary);
 vector<int> res(n + m - 1, INT MAX);
 auto dnc = [&](auto&& self, int res_le, int res_ri,
        int arb_le, int arb_ri) -> void {
   if (res_le >= res_ri) return;
   int mid_res = (res_le + res_ri) / 2;
   int op arb = arb le;
   for (int i = arb_le; i < min(mid_res + 1, arb_ri);</pre>
   i++) {
   int j = mid_res - i;
   if (j >= n) continue;
   if (res[mid_res] > convex[j] + arbitrary[i]) {
     res[mid_res] = convex[j] + arbitrary[i];
     op_arb = i;
   self(self, res_le, mid_res, arb_le,
   min(arb_ri, op_arb + 1));
   self(self, mid_res + 1, res_ri, op_arb, arb_ri);
 dnc(dnc, 0, n + m - 1, 0, m);
 return res;
```

### gcdconv.h

**Description:** ssize(a) = ssize(b) gcdconv[k] = sum of (a[i]\*b[j]) for all pairs(i,j) where gcd(i,j)==kTime:  $\mathcal{O}(N \log N)$ 

```
const int mod = 998'244'353;
vector<int> gcd_convolution(const vector<int>& a,
 const vector<int>& b) {
 int n = ssize(a);
 vector<int> c(n);
 for (int q = n - 1; q >= 1; q--) {
   int64_t sum_a = 0, sum_b = 0;
   for (int i = q; i < n; i += q) {
     sum_a += a[i], sum_b += b[i];
     if ((c[g] -= c[i]) < 0) c[g] += mod;
   sum_a %= mod, sum_b %= mod;
```

 $c[g] = (c[g] + sum_a * sum_b) % mod;$ 

#### lcmconv.h

return c:

**Description:** ssize(a)==ssize(b) lcmconv[k] = sum of (a[i]\*b[i]) for all pairs (i,j) where lcm(i,j)==kee1440, 16 lines

```
const int mod = 998'244'353;
vector<int> lcm_convolution(const vector<int>& a,
 const vector<int>& b) {
```

```
int n = ssize(a);
vector<int64_t> sum_a(n), sum_b(n);
vector<int> c(n);
for (int i = 1; i < n; i++) {
  for (int j = i; j < n; j += i)
   sum_a[j] += a[i], sum_b[j] += b[i];
  sum_a[i] %= mod, sum_b[i] %= mod;
  c[i] = (c[i] + sum_a[i] * sum_b[i]) % mod;
  for (int j = i + i; j < n; j += i)
    if ((c[j] -= c[i]) < 0) c[j] += mod;
return c;
```

## Number theory (7)

### 7.1 Modular arithmetic

#### ModInverse.h

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

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

#### ModLog.h

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

Time:  $\mathcal{O}(\sqrt{m})$ c040b8, 11 lines

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

### ModSum.h

2dfb20, 16 lines

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

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

5c5bc5, 16 lines

```
typedef unsigned long long ull;
ull sumsq(ull to) { return to / 2 * ((to-1) | 1); }
ull divsum(ull to, ull c, ull k, ull m) {
 ull res = k / m * sumsq(to) + c / m * to;
 k %= m; c %= m;
 if (!k) return res;
 ull to2 = (to * k + c) / m;
 return res + (to - 1) * to2 - divsum(to2, m-1 - c, m, k);
11 modsum(ull to, 11 c, 11 k, 11 m) {
 c = ((c \% m) + m) \% m;
 k = ((k % m) + m) % m;
 return to * c + k * sumsq(to) - m * divsum(to, c, k, m);
```

#### ModMulLL.h

**Description:** Calculate  $a \cdot b \mod c$  (or  $a^b \mod c$ ) for  $0 < a, b < c < 7.2 \cdot 10^{18}$ . **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 >= (11)M);
}
ull modpow(ull b, ull e, ull mod) {
    ull ans = 1;
    for (; e; b = modmul(b, b, mod), e /= 2)
        if (e & 1) ans = modmul(ans, b, mod);
    return ans;
}
```

### ModSqrt.h

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

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

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

## 7.2 Primality

FastEratosthenes.h

**Description:** Prime sieve for generating all primes smaller than LIM. **Time:** LIM=1e9  $\approx 1.5$ s

```
6b2912, 20 lines
const int LIM = 1e6;
bitset<LIM> isPrime;
vi eratosthenes() {
  const int S = (int)round(sqrt(LIM)), R = LIM / 2;
  vi pr = {2}, sieve(S+1); pr.reserve(int(LIM/log(LIM)*1.1));
  vector<pii> cp;
  for (int i = 3; i <= S; i += 2) if (!sieve[i]) {
    cp.push_back(\{i, i * i / 2\});
    for (int j = i * i; j \le S; j += 2 * i) sieve[j] = 1;
  for (int L = 1; L \le R; L += S) {
    array<bool, S> block{};
    for (auto &[p, idx] : cp)
     for (int i=idx; i < S+L; idx = (i+=p)) block[i-L] = 1;
    rep(i, 0, min(S, R - L))
     if (!block[i]) pr.push_back((L + i) * 2 + 1);
  for (int i : pr) isPrime[i] = 1;
  return pr;
```

### LinearSieve.h

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

```
const int LIM = 1000000;
vi lp(LIM+1), primes;

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

### MillerRabin.h

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

**Time:** 7 times the complexity of  $a^b \mod c$ .

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

#### Factor.h

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

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

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

#### GetFactors.h

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

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

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

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

## 7.3 Divisibility

euclid.h

32eeca, 8 lines

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

```
11 euclid(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  $0 \le x < \operatorname{lcm}(m, n)$ . Assumes  $mn < 2^{62}$ .

Time:  $\log(n)$ 

### 7.3.1 Bézout's identity

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

$$ax + by = d$$

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

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

phiFunction.h

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

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

### 7.4 Fractions

ContinuedFractions.h

### FracBinarySearch Fraction IntPerm

**Description:** Given N and a real number x > 0, finds the closest rational approximation p/q with p, q < N. It will obey |p/q - x| < 1/qN. For consecutive convergents,  $p_{k+1}q_k - q_{k+1}p_k = (-1)^k$ .  $(p_k/q_k \text{ alternates between }$ > x and < x.) If x is rational, y eventually becomes  $\infty$ ; 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 \sim 1e7; long double for N \sim 1e9
pair<11, 11> approximate(d x, 11 N) {
  11 LP = 0, LQ = 1, P = 1, Q = 0, inf = LLONG_MAX; d y = x;
    ll lim = min(P ? (N-LP) / P : inf, Q ? (N-LQ) / Q : inf),
       a = (ll) floor(y), b = min(a, lim),
       NP = b*P + LP, NO = b*O + LO;
    if (a > b) {
      // If b > a/2, we have a semi-convergent that gives us a
      // better approximation; if b = a/2, we *may* have one.
      // Return {P, Q} here for a more canonical approximation.
      return (abs(x - (d)NP / (d)NO) < abs(x - (d)P / (d)O)) ?
        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 < N. You may want to throw an exception from f if it finds an exact solution, in which case N can be removed.

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

```
struct Frac { ll p, q; };
template<class F>
Frac fracBS(F f, 11 N) {
  bool dir = 1, A = 1, B = 1;
  Frac lo{0, 1}, hi{1, 1}; // Set hi to 1/0 to search (0, N]
  if (f(lo)) return lo;
  assert(f(hi));
  while (A | | B) {
   11 adv = 0, step = 1; // move hi if dir, else lo
    for (int si = 0; step; (step *= 2) >>= si) {
      adv += step;
     Frac mid{lo.p * adv + hi.p, lo.q * adv + hi.q};
     if (abs(mid.p) > N || mid.q > N || dir == !f(mid)) {
       adv -= step; si = 2;
   hi.p += lo.p * adv;
   hi.q += lo.q * adv;
   dir = !dir;
   swap(lo, hi);
   A = B; B = !!adv;
  return dir ? hi : lo;
```

#### Fraction.h

**Description:** Safer struct for representing fractions/rationals. Comparison is 100% overflow safe; other ops are safer but can still overflow. All ops are  $O(\log N)$ .

8ff7f8, 32 lines

template<class T> struct QO { Ta, b;

QO(T p, T q = 1) { T q = qcd(p, q);a = p / g;b = q / g;if (b < 0) a = -a, b = -b; } T gcd(T x, T y) const { return \_\_gcd(x, y); } QO operator+(const QO& o) const { T g = gcd(b, o.b), bb = b / g, obb = o.b / g;return {a \* obb + o.a \* bb, b \* obb}; } QO operator-(const QO& o) const return \*this + QO(-o.a, o.b); 00 operator\*(const 00% o) const T g1 = gcd(a, o.b), g2 = gcd(o.a, b);return { (a / g1) \* (o.a / g2), (b / g2) \* (o.b / g1) }; } QO operator/(const QO& o) const { return \*this \* QO(o.b, o.a); QO recip() const { return {b, a}; } int signum() const { return (a > 0) - (a < 0); } static bool lessThan(T a, T b, T x, T y) { if (a / b != x / y) return a / b < x / y; if (x % y == 0) return false; if (a % b == 0) return true; return lessThan(y, x % y, b, a % b); } bool operator<(const QO& o) const { if (this->signum() != o.signum() || a == 0) return a < o.a; if (a < 0) return lessThan(abs(o.a), o.b, abs(a), b); else return lessThan(a, b, o.a, o.b); } friend ostream& operator << (ostream& cout, const QO& o) { return cout << o.a << "/" << o.b; } };

### 7.5 Pythagorean Triples

The Pythagorean triples are uniquely generated by

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

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

### 7.6 Primes

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  $\mathbb{Z}_2 \times \mathbb{Z}_{2^{a-2}}$ .

### 7.7 Estimates

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

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

### Mobius Function

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

Mobius Inversion:

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

Other useful formulas/forms:

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

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

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

## Combinatorial (8)

### Permutations

### 8.1.1 Factorial

n	1 2 3	4	5 6	7	8	9	10	
n!	1 2 6	24 1	20 720	5040	40320	362880	3628800	
n	11	12	13	14	15	16	17	
n!	4.0e7	4.8e	8.6.2e	9 8.7el	10 1.3e	$12 \ 2.1e1$	3 3.6e14	
n	20	25	30	40	50 10	00 - 150	3628800 17 3 3.6e14 0 171	
n!	2e18	2e25	3e32 8	8e47 3	e64 9e	157  6e26	$62$ >DBL_M.	AX

### IntPerm.h

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

Time:  $\mathcal{O}(n)$ 044568, 6 lines int permToInt(vi& v) { int use = 0, i = 0, r = 0; for(int x:v)  $r = r * ++i + \underline{\quad}$  builtin\_popcount(use & -(1<<x)), use |= 1 << x;// (note: minus, not  $\sim$ !) return r:

### **8.1.2** Cycles

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

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

### 8.1.3 Derangements

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

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

### 8.1.4 Burnside's lemma

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

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

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

### multinomial KMP Zfunc Manacher

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

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

### Partitions and subsets

### 8.2.1 Partition function

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

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

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

#### 8.2.2 Lucas' Theorem

Let n, m be non-negative integers and p a prime. Write  $n = n_k p^k + ... + n_1 p + n_0$  and  $m = m_k p^k + ... + m_1 p + m_0$ . Then  $\binom{n}{m} \equiv \prod_{i=0}^{k} \binom{n_i}{m_i} \pmod{p}.$  **8.2.3** Binomials

multinomial.h

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

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

### General purpose numbers

#### 8.3.1 Bernoulli numbers

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

Sums of powers:

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

Euler-Maclaurin formula for infinite sums:

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

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

### 8.3.2 Stirling numbers of the first kind

Number of permutations on n items with k cycles.

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

c(8, k) = 8, 0, 5040, 13068, 13132, 6769, 1960, 322, 28, 1 $c(n, 2) = 0, 0, 1, 3, 11, 50, 274, 1764, 13068, 109584, \dots$ 

### 8.3.3 Eulerian numbers

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

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

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

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

### 8.3.4 Stirling numbers of the second kind

Partitions of n distinct elements into exactly k groups.

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

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

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

#### 8.3.5 Bell numbers

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

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

#### 8.3.6 Labeled unrooted trees

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

#### 8.3.7 Catalan numbers

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

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

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

- sub-diagonal monotone paths in an  $n \times n$  grid.
- strings with n pairs of parenthesis, correctly nested.
- binary trees with with n+1 leaves (0 or 2 children).

- ordered trees with n+1 vertices.
- ways a convex polygon with n+2 sides can be cut into triangles by connecting vertices with straight lines.
- permutations of [n] with no 3-term increasing subseq.

# Strings (9)

### KMP.h

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

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

```
vi pi(const string& s) {
 vi p(sz(s));
 rep(i, 1, sz(s))
    int q = p[i-1];
    while (g \&\& s[i] != s[g]) g = p[g-1];
    p[i] = g + (s[i] == s[g]);
 return p;
vi match (const string& s, const string& pat) {
 vi p = pi(pat + ' \setminus 0' + s), res;
 rep(i,sz(p)-sz(s),sz(p))
    if (p[i] == sz(pat)) res.push_back(i - 2 * sz(pat));
 return res;
```

### Zfunc.h

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

```
Time: \mathcal{O}(n)
                                                        3ae526, 12 lines
vi Z(string S) {
 vi z(sz(S));
 int 1 = -1, r = -1;
 rep(i,1,sz(S)) {
    z[i] = i >= r ? 0 : min(r - i, z[i - 1]);
    while (i + z[i] < sz(S) \&\& S[i + z[i]] == S[z[i]])
    if (i + z[i] > r)
      1 = i, r = i + z[i];
 return z:
```

#### Manacher.h

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

```
Time: \mathcal{O}(N)
                                                         e7ad79, 13 lines
array<vi, 2> manacher(const string& s) {
  array < vi, 2 > p = {vi(n+1), vi(n)};
  rep(z,0,2) for (int i=0,1=0,r=0; i < n; i++) {
    int t = r-i+!z;
    if (i<r) p[z][i] = min(t, p[z][l+t]);</pre>
    int L = i-p[z][i], R = i+p[z][i]-!z;
    while (L>=1 \&\& R+1< n \&\& s[L-1] == s[R+1])
      p[z][i]++, L--, R++;
    if (R>r) l=L, r=R;
  return p;
```

#### Eertree.h

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

```
Time: \mathcal{O}(|S|) 2fc643, 24 lines struct Eertree {
  vi slink = {0, 0}, len = {-1, 0};
  vvi down;
  int cur = 0, t = 0;
  Eertree(string &str) : down(2, vi(26, -1)) {
    for (int i = 0; i < sz(str); i++) {
      char c = str[i]; int ci = c - 'a';
      while (t <= 0 || str[t-1] != c)
      t = i - len[cur = slink[cur]];
    if (down[cur][ci] == -1) {
```

down[cur][ci] = sz(slink);

down.emplace back(26, -1);

} else slink.push back(1);

} else cur = down[cur][ci];

if (len.back() > 1) {

cur = sz(slink) - 1;

t = i - len[cur] + 1;

len.push\_back(len[cur] + 2);

do t = i - len[cur = slink[cur]];

while  $(t \le 0 \mid | str[t-1] != c);$ 

slink.push\_back(down[cur][ci]);

### MinRotation.h

};

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

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

#### SuffixArrav.h

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

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

```
for (int i = 0, j; i < n - 1; lcp[rank[i++]] = k)
    for (k && k--, j = sa[rank[i] - 1];
        s[i + k] == s[j + k]; k++);
};</pre>
```

#### SuffixAutomaton.h

**Description:** Creates a partial DFA (DAG) that accepts all suffixes, with suffix links. One-to-one map between a path from the root and a substring len is the longest-length substring ending here. pos is the first index in the string matching here. term is whether this node is a terminal (aka a suffix) **Time:** construction takes  $\mathcal{O}(N \log K)$ , where  $K = \text{Alphabet Size}_{1914a9, 22 \text{ lines}}$ 

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

#### SuffixTree.h

**Description:** Ukkonen's algorithm for online suffix tree construction. Each node contains indices [l, r) into the string, and a list of child nodes. Suffixes are given by traversals of this tree, joining [l, r) substrings. The root is 0 (has l = -1, r = 0), non-existent children are -1. To get a complete tree, append a dummy symbol – otherwise it may contain an incomplete path (still useful for substring matching, though). **Time:**  $\mathcal{O}(26N)$ 

```
aae0b8, 50 lines
struct SuffixTree {
 enum { N = 200010, ALPHA = 26 }; // N \sim 2*maxlen+10
 int toi(char c) { return c - 'a'; }
 string a; //v = cur \ node, q = cur \ position
 int t[N][ALPHA], 1[N], r[N], p[N], s[N], v=0, q=0, m=2;
 void ukkadd(int i, int c) { suff:
   if (r[v] \le q) {
     if (t[v][c]==-1) { t[v][c]=m; l[m]=i;
       p[m++]=v; v=s[v]; q=r[v]; goto suff; }
     v=t[v][c]; q=l[v];
   if (q==-1 || c==toi(a[q])) q++; else {
     l[m+1]=i; p[m+1]=m; l[m]=l[v]; r[m]=q;
     p[m]=p[v]; t[m][c]=m+1; t[m][toi(a[q])]=v;
     l[v]=q; p[v]=m; t[p[m]][toi(a[l[m]])]=m;
     v=s[p[m]]; q=l[m];
     while (q < r[m]) \{ v = t[v][toi(a[q])]; q + = r[v] - l[v]; \}
     if (q==r[m]) s[m]=v; else s[m]=m+2;
     q=r[v]-(q-r[m]); m+=2; qoto 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] = l[1] = -1; r[0] = r[1] = p[0] = p[1] = 0;
    rep(i,0,sz(a)) ukkadd(i, toi(a[i]));
  // example: find longest common substring (uses ALPHA = 28)
  pii best;
  int lcs(int node, int i1, int i2, int olen) {
   if (l[node] <= i1 && i1 < r[node]) return 1;</pre>
    if (1[node] <= i2 && i2 < r[node]) return 2;</pre>
    int mask = 0, len = node ? olen + (r[node] - 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.

4b8fa:

```
// Arithmetic mod 2^64-1. 2x slower than mod 2^64 and more
// code, but works on evil test data (e.g. Thue-Morse, where
// ABBA... and BAAB... of length 2^10 hash the same mod 2^64).
// "typedef ull H;" instead if you think test data is random,
// or work mod 10^9+7 if the Birthday paradox is not a problem.
typedef uint64 t ull;
struct H {
 ull x; H(ull x=0) : x(x) {}
  H operator+(H o) { return x + o.x + (x + o.x < x); }
  H operator-(H o) { return *this + ~o.x; }
  H 	ext{ operator} * (H 	ext{ o}) { auto } m = (\underline{\text{uint128}} t) x * o.x;
   return H((ull)m) + (ull)(m >> 64); }
  ull get() const { return x + !~x; }
  bool operator==(H o) const { return get() == o.get(); }
  bool operator<(H o) const { return get() < o.get(); }</pre>
static const H C = (11)1e11+3; // (order \sim 3e9; random also ok)
H hashString(string& s){H h{}; for(char c:s) h=h*C+c; return h;}
```

### HashInterval.h

**Description:** Various self-explanatory methods for string hashing.

```
"Hashing.h" 122649, 12 lines
struct HashInterval {
  vector<H> ha, pw;
  HashInterval(string& str) : ha(sz(str)+1), pw(ha) {
    pw[0] = 1;
    rep(i,0,sz(str))
    ha[i+1] = ha[i] * C + str[i],
    pw[i+1] = pw[i] * C;
}
H hashInterval(int a, int b) { // hash [a, b)
    return ha[b] - ha[a] * pw[b - a];
};
```

753a4c, 19 lines

2932a0, 17 lines

#### Wildcard.h

Description: string matching with wildcards, returns boolean vector of size s-p+1 representing if a match occurs at this start position, wild cards are repsented by 0 and can be in s,p or both.

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

```
vector<vl> make powers(const v1& v) {
   int n = sz(v);
   vector < vl > pws(3, vl(n)); pws[0] = v;
    rep(k, 1, 3) rep(i, 0, n) //mod?
       pws[k][i] = pws[k-1][i] * v[i];
    return pws;
vector<bool> wildcard pattern matching(const vl& s,
   const v1& p) {
   int n = sz(s), m = sz(p);
   auto s_pws = make_powers(s), p_pws = make_powers(p);
   for (auto& p_pw : p_pws) reverse(all(p_pw));
   vector<vl> res(3);
   rep(pw_hay, 0, 3) //ntt
       res[pw_hay] = conv(s_pws[pw_hay], p_pws[2 - pw_hay]);
   vector < bool > mtch(n - m + 1);
    rep(i,0,sz(mtch)){
       int id = i + m - 1;
       auto num = res[0][id] - 2 * res[1][id] + res[2][id];
       mtch[i] = !num; //num == 0
   return mtch;
```

### AhoCorasick-Tyler.h

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

**Time:** construction takes  $\mathcal{O}(26N)$ , where N = sum of length of patterns. find(x) is  $\mathcal{O}(N)$ , where N = length of x. findAll is  $\mathcal{O}(NM)$ . 647ca9, 47 lines

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

```
trie[cur].ids.push_back(id);
int getLink(int cur);
int calc(int cur, char c) {
 int idx = c - 'a';
 auto &ret = trie[cur].linkMemo[idx];
 if (ret != -1) return ret;
 if(trie[cur].nxt[idx] != -1)
   return ret = trie[cur].nxt[idx];
 return ret = cur == 0 ? 0 : calc(getLink(cur), c);
int getLink(int cur) {
 auto &ret = trie[cur].link;
 if (ret != -1) return ret;
 if(cur == 0 || trie[cur].prv == 0) return ret = 0;
 return ret = calc(getLink(trie[cur].prv), trie[cur].c);
```

## Various (10)

### 10.1 Intervals

IntervalContainer.h

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

```
Time: \mathcal{O}(\log N)
set<pii>::iterator addInterval(set<pii>& is, int L, int R) {
  if (L == R) return is.end();
 auto it = is.lower_bound({L, R}), before = it;
  while (it != is.end() && it->first <= R) {
   R = max(R, it->second);
   before = it = is.erase(it);
 if (it != is.begin() && (--it)->second >= L) {
   L = min(L, it->first);
   R = max(R, it->second);
   is.erase(it);
 return is.insert(before, {L,R});
void removeInterval(set<pii>& is, int L, int R) {
 if (L == R) return;
 auto it = addInterval(is, L, R);
 auto r2 = it->second;
 if (it->first == L) is.erase(it);
 else (int&)it->second = L;
 if (R != r2) is.emplace(R, r2);
```

#### IntervalCover.h

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

```
9e9d8d, 19 lines
template<class T>
vi cover(pair<T, T> G, vector<pair<T, T>> I) {
 vi S(sz(I)), R;
 iota(all(S), 0);
 sort(all(S), [&](int a, int b) { return I[a] < I[b]; });</pre>
```

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

#### ConstantIntervals.h

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

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

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

#### Misc. algorithms 10.2

LIS.h

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

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

### FastKnapsack.h

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

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

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

## 10.3 Dynamic programming

### KnuthDP.h

**Description:** When doing DP on intervals:  $a[i][j] = \min_{i < k < j} (a[i][k] + a[i][k])$ a[k][j] + f(i,j), where the (minimal) optimal k increases with both i and j, one can solve intervals in increasing order of length, and search k = p[i][j] for a[i][j] only between p[i][j-1] and p[i+1][j]. This is known as Knuth DP. Sufficient criteria for this are if  $f(b,c) \leq f(a,d)$  and  $f(a,c) + f(b,d) \le f(a,d) + f(b,c)$  for all  $a \le b \le c \le d$ . Consider also: LineContainer (ch. Data structures), monotone queues, ternary search. Time:  $\mathcal{O}(N^2)$ 

### DivideAndConquerDP.h

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

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

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

### 10.4 Debugging tricks

• signal(SIGSEGV, [](int) { \_Exit(0); }); converts segfaults into Wrong Answers. Similarly one can catch SIGABRT (assertion failures) and SIGFPE (zero divisions). \_GLIBCXX\_DEBUG failures generate SIGABRT (or SIGSEGV on gcc 5.4.0 apparently).

• feenableexcept (29); kills the program on NaNs (1), 0-divs (4), infinities (8) and denormals (16).

### 10.5 Optimization tricks

\_\_builtin\_ia32\_ldmxcsr(40896); disables denormals (which make floats 20x slower near their minimum value).

### 10.5.1 Bit hacks

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

### 10.5.2 Pragmas

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

#### FastMod.h

**Description:** Compute a%b about 5 times faster than usual, where b is constant but not known at compile time. Returns a value congruent to a (mod 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</pre>

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

```
7b3c70, 17 lines
inline char gc() { // like getchar()
 static char buf[1 << 16];
 static size_t bc, be;
 if (bc >= be) {
   buf[0] = 0, bc = 0;
   be = fread(buf, 1, sizeof(buf), stdin);
 return buf[bc++]; // returns 0 on EOF
int readInt() {
 int a, c;
 while ((a = gc()) < 40);
```

```
if (a == '-') return -readInt();
while ((c = qc()) >= 48) a = a * 10 + c - 480;
return a - 48;
```

### BumpAllocator.h

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

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

### SmallPtr.h

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

```
"BumpAllocator.h"
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<vector<int, small<int>>> ed(N);
```

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

#### SIMD.h

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

551b82, 43 lines

bb66d4, 14 lines

```
#pragma GCC target ("avx2") // or sse4.1
#include "immintrin.h"
typedef __m256i mi;
#define L(x) _mm256_loadu_si256((mi*)&(x))
```

```
// High-level/specific methods:
// load(u)?_si256, store(u)?_si256, setzero_si256, \_mm\_malloc
// blendv_{-}(epi8|ps|pd) (z?y:x), movemask_{-}epi8 (hibits of bytes)
// i32gather_epi32(addr, x, 4): map addr[] over 32-b parts of x
// sad_epu8: sum of absolute differences of u8, outputs 4xi64
// maddubs_epi16: dot product of unsigned i7's, outputs 16xi15
// madd_epi16: dot product of signed i16's, outputs 8xi32
// extractf128_si256(, i) (256->128), cvtsi128_si32 (128->lo32)
// permute2f128_si256(x,x,1) swaps 128-bit lanes
// shuffle_epi32(x, 3*64+2*16+1*4+0) = x for each lane
// shuffle_epi8(x, y) takes a vector instead of an imm
// Methods that work with most data types (append e.g. _epi32):
// set1, blend (i8?x:y), add, adds (sat.), mullo, sub, and/or,
// and not, abs, min, max, sign(1,x), cmp(gt|eq), unpack(lo|hi)
int sumi32(mi m) { union {int v[8]; mi m;} u; u.m = m;
 int ret = 0; rep(i,0,8) ret += u.v[i]; return ret; }
mi zero() { return _mm256_setzero_si256(); }
mi one() { return _mm256_set1_epi32(-1); }
bool all_zero(mi m) { return _mm256_testz_si256(m, m); }
bool all_one(mi m) { return _mm256_testc_si256(m, one()); }
11 example_filteredDotProduct(int n, short* a, short* b) {
  int i = 0; 11 r = 0;
  mi zero = _mm256_setzero_si256(), acc = zero;
  while (i + 16 \le n) {
    mi \ va = L(a[i]), \ vb = L(b[i]); \ i += 16;
    va = _mm256_and_si256(_mm256_cmpgt_epi16(vb, va), va);
    mi vp = _mm256_madd_epi16(va, vb);
    acc = _mm256_add_epi64(_mm256_unpacklo_epi32(vp, zero),
      _mm256_add_epi64(acc, _mm256_unpackhi_epi32(vp, zero)));
  union {ll v[4]; mi m;} u; u.m = acc; rep(i,0,4) r += u.v[i];
  for (;i \le n;++i) if (a[i] \le b[i]) r += a[i] *b[i]; // <- equiv
  return r;
```

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