

# University of New South Wales

# idk

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C	$\frac{\text{Contest}}{\text{contest}}$ (1)	
e:	mplate.cpp	14 lines
is	14 filles	
	efine rep(i, a, b) for(int i = a; i < (b); ++i)	

#define sz(x) (int)(x).size() typedef long long 11; typedef pair<int, int> pii; typedef vector<int> vi; int main() { cin.tie(0)->sync\_with\_stdio(0); cin.exceptions(cin.failbit);

# .bashrc

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

#### hash.sh

# Hashes a file, ignoring all whitespace and comments. Use for # verifying that code was correctly typed. cpp -dD -P -fpreprocessed | tr -d '[:space:]' | md5sum |cut -c-6

# troubleshoot.txt

Write a few simple test cases if sample is not enough. Are time limits close? If so, generate max cases. Is the memory usage fine? Could anything overflow? Make sure to submit the right file.

Print your solution! Print debug output, as well.

```
Can your algorithm handle the whole range of input?
Read the full problem statement again.
Do you handle all corner cases correctly?
Have you understood the problem correctly?
Any uninitialized variables?
Any overflows?
Confusing N and M, i and j, etc.?
Are you sure your algorithm works?
What special cases have you not thought of?
Are you sure the STL functions you use work as you think?
Add some assertions, maybe resubmit.
Create some testcases to run your algorithm on.
Go through the algorithm for a simple case.
Go through this list again.
Explain your algorithm to a teammate.
Ask the teammate to look at your code.
Go for a small walk, e.g. to the toilet.
Is your output format correct? (including whitespace)
Rewrite your solution from the start or let a teammate do it.
```

Are you clearing all data structures between test cases?

# Runtime error:

Have you tested all corner cases locally? Any uninitialized variables? Are you reading or writing outside the range of any vector? Any assertions that might fail? Any possible division by 0? (mod 0 for example) Any possible infinite recursion? Invalidated pointers or iterators? Are you using too much memory? Debug with resubmits (e.g. remapped signals, see Various).

Time limit exceeded:

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

Memory limit exceeded:

What is the max amount of memory your algorithm should need? Are you clearing all data structures between test cases?

# Mathematics (2)

# 2.1 Equations

$$ax + by = e$$

$$cx + dy = f$$

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

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

# 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 Sums

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

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

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

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

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

# Data structures (3)

#### OrderStatisticTree.h

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

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

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

# HashMap.h

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

```
#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;
   ll operator()(ll x) const { return __builtin_bswap64(x*C);
__gnu_pbds::gp_hash_table<11, int, chash> h({}, {}, {}, {}, {1}
```

#### LazySegtree.h

**Description:** Generic lazy segment tree

Usage: Implement +, upd and id for Node object; += and id for Lazy object.

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

683133, 49 lines struct LazySeg { int n: vt<Node> seq; vt<Lazy> lazy; void init(int \_n) { for  $(n = 1; n < _n; n *= 2);$ seq.resize(2 \* n, nid); lazy.resize(2 \* n, lid);

```
void pull(int i) {
    seg[i] = seg[2 * i] + seg[2 * i + 1];
void push(int i, int l, int r) {
    seg[i].upd(lazy[i], 1, r);
   if (r - 1 > 1) FOR (j, 2) lazy[2 * i + j] += lazy[i];
   lazv[i] = lid;
void build() {
    for (int i = n - 1; i > 0; i--) pull(i);
void upd(int lo, int hi, Lazy val) { upd(lo, hi, val, 1, 0,
void upd(int lo, int hi, Lazy val, int i, int l, int r) {
   if (r == -1) r = n;
   push(i, 1, r);
   if (r <= lo || l >= hi) return;
   if (lo <= 1 && r <= hi) {
        lazy[i] += val;
        push(i, 1, r);
        return;
    int m = (1 + r) / 2;
   upd(lo, hi, val, 2 * i, 1, m);
   upd(lo, hi, val, 2 * i + 1, m, r);
   pull(i);
Node query() { return query(0, n, 1, 0, n); }
Node query(int lo, int hi) { return query(lo, hi, 1, 0, n);
Node query(int lo, int hi, int i, int l, int r) {
   push(i, 1, r);
   if (r <= lo || l >= hi) return nid;
   if (lo <= 1 && r <= hi) return seg[i];
   int m = (1 + r) / 2;
    return query(lo, hi, 2 * i, 1, m)
        + query(lo, hi, 2 * i + 1, m, r);
Node& operator[](int i) {
    return seq[i + n];
```

#### SparseSegtree.h

};

**Description:** Generic-ish sparse segment tree (point update, range query). Usage: Choose appropriate identity element and merge function. Time:  $\mathcal{O}(\log N)$ .

```
d51c9b, 28 lines
using ptr = struct Node*;
const int sz = 1 \ll 30;
struct Node {
    #define func(a, b) min(a, b)
    #define ID INF
   ll val;
   ptr lc, rc;
   ptr get(ptr& p) { return p ? p : p = new Node {ID}; }
    ll querv(int lo, int hi, int l = 0, int r = sz) {
       if (lo >= r || hi <= 1) return ID;
       if (lo <= l && r <= hi) return val;
       int m = (1 + r) / 2;
        return func(get(lc)->query(lo, hi, l, m),
            get(rc)->query(lo, hi, m, r));
   ll upd(int i, ll nval, int l = 0, int r = sz) {
       if (r - 1 == 1) return val = nval;
       int m = (1 + r) / 2;
```

```
if (i < m) get(lc)->upd(i, nval, l, m);
       else get(rc)->upd(i, nval, m, r);
       return val = func(get(lc)->val, get(rc)->val);
    #undef ID
    #undef func
PersistentSegtree.h
Description: Generic-ish persistent segment tree (point update, range
```

Usage: Choose appropriate identity element and merge function. Time:  $\mathcal{O}(\log N)$ . f155de, 30 lines

```
using ptr = struct Node*;
const int sz = 1 << 18;
struct Node {
    #define func(a, b) min(a, b)
    #define ID inf
   int v:
   ptr lc, rc;
   ptr pull(ptr lc, ptr rc) {
        return new Node {func(lc->v, rc->v), lc, rc};
   ptr upd(int i, int nv, int l = 0, int r = sz) {
       if (r - 1 == 1) return new Node {nv};
       int m = (1 + r) / 2;
       if (i < m) return pull(lc->upd(i, nv, l, m), rc);
       else return pull(lc, rc->upd(i, nv, m, r));
   int query(int lo, int hi, int l = 0, int r = sz) {
       if (lo >= r || hi <= l) return ID;
       if (lo <= 1 && r <= hi) return v;
       int m = (1 + r) / 2;
       return func(lc->query(lo, hi, l, m),
           rc->query(lo, hi, m, r));
    #undef id
    #undef func
```

#### LiChaoTree.h

Description: LiChao tree

Usage: self explanatory i think

Time:  $\mathcal{O}(\log N)$ . 00d540, 36 lines

```
struct Line {
   11 m, c;
    11 operator()(11 x) {
        return m * x + c;
};
const 11 sz = 111 << 30;
using ptr = struct Node*;
struct Node {
   ptr lc, rc;
   Line line:
    Node(Line _line) {
       line = _line;
        1c = rc = 0;
};
```

```
// min tree (flip signs for max)
void add(ptr& n, Line loser, 11 1 = 0, 11 r = sz) {
    if (n ? 0 : n = new Node(loser)) return;
    11 \text{ m} = (1 + r) / 2;
    if (loser(m) < n->line(m)) swap(loser, n->line);
    if (r - 1 == 1) return;
    if (loser(1) < n->line(1)) add(n->lc, loser, 1, m);
    else add(n->rc, loser, m, r);
ll query (ptr n, ll x, ll l = 0, ll r = sz) {
    if (!n) return sz:
    11 m = (1 + r) / 2;
    if (x < m) return min(n->line(x), query(n->lc, x, 1, m));
    else return min(n->line(x), query(n->rc, x, m, r));
```

# SparseTable.h

**Description:** Generic sparse table for idempotent operations.

Usage: Define the desired operation

**Time:**  $\mathcal{O}(N \log N)$  build,  $\mathcal{O}(1)$  query.

```
3e1539, 18 lines
```

```
template<class T> struct RMQ {
    #define func min
    vt<vt<T>> dp;
    void init(const vt<T>& v) {
        dp.resize(32 - __builtin_clz(size(v)), vt<T>(size(v)));
        copy(all(v), begin(dp[0]));
        for (int j = 1; 1 << j <= size(v); ++j) {
            for (int i = 0; i < size(v) - (1 << j) + 1; i++)
                dp[j][i] = func(dp[j-1][i],
                    dp[j-1][i+(1<<(j-1))]);
    T query(int 1, int r) {
        int d = 31 - \underline{\quad} builtin_clz(r - 1);
        return func(dp[d][1], dp[d][r - (1 << d)]);
    #undef func
};
```

# StaticRangeQuery.h

Description: Generic static range query for associative operations.

Usage: Define the desired operation

**Time:**  $\mathcal{O}(N \log N)$  build,  $\mathcal{O}(1)$  query.

82b9ca, 34 lines

```
template<class T> struct RangeQuery {
    \#define\ comb(a,\ b)\ (a)\ +\ (b)
    #define id 0
   int lg, n;
    vt<vt<T>> stor;
    vt<T> a;
    void fill(int 1, int r, int ind) {
       if (ind < 0) return;
       int m = (1 + r) / 2;
       T prod = id;
        FOR (i, m, r) stor[i][ind] = prod = comb(prod, a[i]);
        prod = id;
        ROF (i, 1, m) stor[i][ind] = prod = comb(a[i], prod);
        fill(1, m, ind - 1);
        fill(m, r, ind - 1);
    template <typename It>
    void build(It 1, It r) {
       lg = 1;
       while ((1 << lg) < r - l) lg++;
        n = 1 << 1q;
        a.resize(n, id);
```

```
stor.resize(n, vt<T>(32 - __builtin_clz(n)));
        fill(0, n, lq - 1);
    T query(int 1, int r) {
        if (1 == r) return a[1];
        int t = 31 - \underline{\quad} builtin_clz(r ^ 1);
        return comb(stor[1][t], stor[r][t]);
    #undef id
    #undef comb
};
CaterpillowTree.h
Description: 64-ary set
Usage: bruh
Time: \mathcal{O}(\log_{64} N).
                                                        426eac, 89 lines
using ull = unsigned long long;
const int depth = 3;
const int sz = 1 \ll (depth * 6);
struct Tree {
    vt<ull> seg[depth];
    Tree() {
        FOR (i, depth) seg[i].resize(1 << (6 * i));
    void insert(int x) {
        ROF (d, 0, depth) {
            seq[d][x >> 6] = 1ull << (x & 63);
            x >>= 6;
    void erase(int x) {
        ull b = 0:
        ROF (d, 0, depth) {
            seg[d][x >> 6] &= \sim(1ull << (x & 63));
            seg[d][x >> 6] = b << (x & 63);
            x >>= 6;
            b = bool(seq[d][x]);
    int next(int x) {
        if (x \ge sz) return sz;
        x = std::max(x, 0);
        int d = depth - 1;
        while (true) {
            if (ull m = seq[d][x >> 6] >> (x & 63)) {
                 x += __builtin_ctzll(m);
                break;
            x = (x >> 6) + 1;
            if (d == 0 | | x >= (1 << (6 * d))) return sz;
            d--;
        while (++d < depth) {
            x = (x \ll 6) + \underline{\underline{}}builtin_ctzll(seg[d][x]);
        return x;
    int prev(int x) {
        if (x < 0) return -1;
        x = std::min(x, sz - 1);
        int d = depth - 1;
```

for (It i = 1; i != r; i++) a[i - 1] = \*i;

```
while (true) {
            if (ull m = seg[d][x >> 6] << (63 - (x & 63))) {
                x -= __builtin_clzll(m);
                break;
            x = (x >> 6) - 1;
            if (d == 0 | | x == -1) return -1;
            d--;
        while (++d < depth) {
            x = (x << 6) + 63 - \underline{builtin_clzll(seq[d][x])};
        return x;
    }
    int min() {
        if (empty()) return sz;
        int ans = 0;
        FOR (d, depth) {
            ans <<= 6;
            ans += __builtin_ctzll(seq[d][ans >> 6]);
        return ans;
    }
    int max() {
        if (empty()) return -1;
        int ans = 0;
        FOR (d, depth) {
            ans <<= 6;
            ans += 63 - __builtin_clzll(seg[d][ans >> 6]);
        return ans;
    inline bool empty() { return !seg[0][0]; }
    inline int operator[](int i) { return 1 & (seq[depth - 1][i
          >> 6] >> (i & 63)); }
};
Treap.h
Description: Treap with too many operations
Time: \mathcal{O}(\log N)
                                                     817442, 218 lines
using K = 11;
random_device rd;
mt19937 mt(rd());
struct Lazy {
   11 v;
  bool inc, rev;
    void operator+=(const Lazy &b) {
```

```
Description: Treap with too many operations
Time: O(\log N)

using K = ll;
random_device rd;
mt19937 mt(rd());

struct Lazy {
    ll v;
    bool inc, rev;
    void operator+=(const Lazy &b) {
        if (b.inc) v += b.v;
        else v = b.v, inc = false;
        rev ^= b.rev;
    }
};

struct Value {
    ll mx, sum;
    void upd(const Lazy &b, int sz) {
        if (!b.inc) mx = sum = 0;
        mx += b.v, sum += b.v * sz;
    }
    Value operator+(const Value &b) const {
        return {max(mx, b.mx), sum + b.sum};
    }
};
```

```
const Lazy LID = {0, true, false};
const Value VID = {INF, 0};
using ptr = struct Node*;
struct Node {
    int pri;
    K key;
    ptr 1, r;
    int sz:
    Value val, agg;
    Lazy lazy;
    Node(K key, Value val) : key(key), val(val), agg(val) {
        sz = 1;
        pri = mt();
        1 = r = 0;
        lazy = LID;
    ~Node() {
        delete 1;
        delete r;
};
int sz(ptr n) { return n ? n->sz : 0; }
Value val(ptr n) { return n ? n->val : VID; }
Value agg(ptr n) { return n ? n->agg : VID; }
ptr push(ptr n) {
    if (!n) return n;
    if (n->lazy.rev) swap(n->l, n->r);
    ptr 1 = n->1, r = n->r;
    n->val.upd(n->lazy, 1);
    n->agg.upd(n->lazy, n->sz);
    if (1) n\rightarrow 1\rightarrow 1azy += n\rightarrow 1azy;
    if (r) n\rightarrow r\rightarrow lazv += n\rightarrow lazv;
    n->lazy = LID;
    return n;
ptr pull(ptr n) {
    ptr 1 = n->1, r = n->r;
    push(1), push(r);
    n->sz = sz(1) + 1 + sz(r);
    n->agg = agg(1) + n->val + agg(r);
    return n;
pair<ptr, ptr> split(ptr n, K k) {
    if (!n) return {n, n};
    push(n);
    if (k \le n->key) {
        auto [1, r] = split(n->1, k);
        n->1 = r:
        return {1, pull(n)};
    } else {
        auto [1, r] = split(n->r, k);
        n->r = 1;
        return {pull(n), r};
pair<ptr, ptr> spliti(ptr n, int i) {
    if (!n) return {n, n};
    push(n);
    if (i \le sz(n->1)) {
```

```
auto [1, r] = spliti(n->1, i);
       n->1 = r:
        return {1, pull(n)};
        auto [1, r] = spliti(n->r, i - sz(n->1) - 1);
       n->r = 1;
        return {pull(n), r};
ptr merge(ptr 1, ptr r) {
    if (!1 || !r) return 1 ? 1 : r;
    push(1), push(r);
   ptr t;
    if (1->pri > r->pri) 1->r = merge(1->r, r), t = 1;
    else r->1 = merge(1, r->1), t = r;
    return pull(t);
ptr ins(ptr n, K k, Value val) { // insert k
  auto [1, r] = split(n, k);
  return merge(1, merge(new Node(k, val), r));
ptr insi(ptr n, int i, K k, Value val) { // insert before i
  auto [1, r] = spliti(n, i);
  return merge(l, merge(new Node(k, val), r));
ptr del(ptr n, K k) { // delete k
 auto a = split(n, k), b = spliti(a.s, 1);
  return merge(a.f, b.s);
ptr deli(ptr n, int i) {
 auto b = spliti(n, i + 1), a = spliti(b.f, i);
  return merge(a.f, b.s);
ptr find(ptr n, K k) {
  if (!n || n->key == k) return n;
  if (k < n->key) return find(n->1, k);
  else return find(n->r, k);
ptr findi(ptr n, int i) {
  if (!n \mid | i == sz(n->1)) return n;
  if (i < sz(n->1)) return find(n->1, i);
  else return find(n->r, i);
ptr upd(ptr n, K lo, K hi, Lazy nv) {
  if (lo > hi) return n;
  auto [lhs, r] = split(n, hi + 1);
  auto [1, m] = split(lhs, lo);
 m->lazv += nv;
  return merge(l, merge(m, r));
ptr updi(ptr n, int lo, int hi, Lazy nv) {
 if (lo > hi) return n;
  auto [lm, r] = spliti(n, hi + 1);
  auto [1, m] = spliti(lm, lo);
 m->lazy += nv;
  return merge(l, merge(m, r));
```

```
Value query(ptr &n, K lo, K hi) {
  auto [lm, r] = split(n, hi + 1);
  auto [1, m] = split(lm, lo);
  Value res = agg(m);
 n = merge(1, merge(m, r));
  return res:
Value queryi(ptr &n, int lo, int hi) {
 auto [lm, r] = spliti(n, hi + 1);
  auto [1, m] = spliti(lm, lo);
  Value res = agg(m);
  n = merge(1, merge(m, r));
  return res;
int mn(ptr n) {
    assert(n);
    push(n);
    if (n->1) return mn(n->1);
    else return n->key;
ptr unite(ptr 1, ptr r) {
    if (!1 || !r) return 1 ? 1 : r;
    // l has the smallest key
    if (mn(1) > mn(r)) swap(1, r);
    ptr res = 0;
    while (r) {
        auto [lt, rt] = split(1, mn(r) + 1);
        res = merge(res, lt);
        tie(l, r) = make_pair(r, rt);
    return merge(res, 1);
void heapify(ptr n) {
 if (!n) return;
 ptr mx = n;
  if (n->1 && n->1->pri > mx->pri) mx = n->1;
 if (n->r && n->r->pri > mx->pri) mx = n->r;
 if (mx != n) swap(n->pri, mx->pri), heapify(mx);
ptr build(int 1, int r, vt<ptr>& ns) {
 if (1 > r) return nullptr;
 if (1 == r) return ns[1];
 int m = (r + 1) / 2;
 ns[m] -> 1 = build(1, m - 1, ns);
  ns[m] \rightarrow r = build(m + 1, r, ns);
  heapify(ns[m]);
  return pull(ns[m]);
Node* tree;
```

# Numerical (4)

# 4.1 Polynomials and recurrences

# Polynomial.h

```
c9b7b0, 17 lines
```

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

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

# PolyRoots.h

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

```
Usage: polyRoots(\{\{2,-3,1\}\},-1e9,1e9\}) // solve x^2-3x+2=0
Time: \mathcal{O}\left(n^2\log(1/\epsilon)\right)
"Polynomial.h"
                                                                           b00bfe, 23 lines
```

```
vector<double> polyRoots(Poly p, double xmin, double xmax) {
 if (sz(p.a) == 2) \{ return \{-p.a[0]/p.a[1]\}; \}
 vector<double> ret;
 Polv 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:  $\mathcal{O}(n^2)$ 

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

# BerlekampMassev.h

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

```
Usage: berlekampMassey(\{\hat{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<11> C(n), B(n), T;
 C[0] = B[0] = 1;
 11 b = 1;
  rep(i, 0, n) \{ ++m;
   ll d = s[i] % mod;
   rep(j, 1, L+1) d = (d + C[j] * s[i - j]) % mod;
   if (!d) continue;
   T = C; 11 coef = d * modpow(b, mod-2) % mod;
   rep(j,m,n) C[j] = (C[j] - coef * B[j - m]) % mod;
   if (2 * L > i) continue;
   L = i + 1 - L; B = T; b = d; m = 0;
 C.resize(L + 1); C.erase(C.begin());
  for (11& x : C) x = (mod - x) % mod;
  return C;
```

# LinearRecurrence.h

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

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

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

# 4.2 Optimization

# GoldenSectionSearch.h

**Description:** Finds the argument minimizing the function f in the interval [a,b] assuming f is unimodal on the interval, i.e. has only one local minimum and no local maximum. The maximum error in the result is eps. Works equally well for maximization with a small change in the code. See Ternary-Search.h in the Various chapter for a discrete version.

```
\label{eq:Usage: double func(double x) { return 4+x+.3*x*x; } double xmin = gss(-1000,1000,func); \\ \mathbf{Time:} \ \mathcal{O}\left(\log((b-a)/\epsilon)\right) \\ \phantom{\mathbf{\mathbf{31d45b}}} 31d45b, 14 \ \mathrm{lines}
```

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

# HillClimbing.h

Description: Poor man's optimization for unimodal functions<sub>8eeeaf, 14 lines</sub>

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

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

#### Integrate.h

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

4756fc. 7 lines

```
template < class F >
double quad(double a, double b, F f, const int n = 1000) {
  double h = (b - a) / 2 / n, v = f(a) + f(b);
  rep(i,1,n*2)
  v += f(a + i*h) * (i&l ? 4 : 2);
  return v * h / 3;
}
```

# IntegrateAdaptive.h

```
Description: Fast integration using an adaptive Simpson's rule.

Usage: double sphereVolume = quad(-1, 1, [] (double x) {
return quad(-1, 1, [&] (double y) {
return quad(-1, 1, [&] (double z) {
return x*x + y*y + z*z < 1; });});

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

template <class F>
d rec(F& f, d a, d b, d eps, d S) {
```

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

# Simplex.h

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

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

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

```
typedef double T; // long double, Rational, double + mod<P>...
typedef vector<T> vd;
typedef vector<vd> vvd;
const T eps = 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;
    D[r][s] = inv;
    swap(B[r], N[s]);
 bool simplex(int phase) {
    int x = m + phase - 1;
    for (;;) {
      int s = -1;
      rep(j, 0, n+1) if (N[j] != -phase) ltj(D[x]);
      if (D[x][s] >= -eps) return true;
      int r = -1;
      rep(i,0,m) {
       if (D[i][s] <= eps) continue;</pre>
        if (r == -1 \mid | MP(D[i][n+1] / D[i][s], B[i])
                     < MP(D[r][n+1] / D[r][s], B[r])) r = i;
      if (r == -1) return false;
      pivot(r, s);
 T solve(vd &x) {
   int r = 0;
    rep(i,1,m) if (D[i][n+1] < D[r][n+1]) r = i;
    if (D[r][n+1] < -eps) {
      pivot(r, n);
```

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

# Matrices

#### Determinant.h

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

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

#### IntDeterminant.h

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

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

3313dc, 18 lines

```
const 11 mod = 12345;
ll det(vector<vector<ll>>& a) {
 int n = sz(a); ll ans = 1;
 rep(i,0,n) {
    rep(j,i+1,n) {
     while (a[j][i] != 0) { // gcd step }
       ll t = a[i][i] / a[j][i];
       if (t) rep(k,i,n)
         a[i][k] = (a[i][k] - a[j][k] * t) % mod;
       swap(a[i], a[j]);
       ans *=-1;
   ans = ans * a[i][i] % mod;
   if (!ans) return 0;
  return (ans + mod) % mod;
```

## SolveLinear.h

**Description:** Solves A \* x = b. If there are multiple solutions, an arbitrary one is returned. Returns rank, or -1 if no solutions. Data in A and b is lost. Time:  $\mathcal{O}\left(n^2m\right)$ 44c9ab, 38 lines

```
typedef vector<double> vd;
const double eps = 1e-12;
int solveLinear(vector<vd>& A, vd& b, vd& x) {
 int n = sz(A), m = sz(x), rank = 0, br, bc;
 if (n) assert(sz(A[0]) == m);
```

```
vi col(m); iota(all(col), 0);
rep(i,0,n) {
  double v, bv = 0;
  rep(r,i,n) rep(c,i,m)
   if ((v = fabs(A[r][c])) > bv)
      br = r, bc = c, bv = v;
  if (bv <= eps) {
   rep(j,i,n) if (fabs(b[j]) > eps) return -1;
   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:

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

## SolveLinearBinary.h

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

```
typedef bitset<1000> bs:
int solveLinear(vector<bs>& A, vi& b, bs& x, int m) {
 int n = sz(A), rank = 0, br;
 assert(m \le sz(x));
 vi col(m); iota(all(col), 0);
 rep(i,0,n) {
   for (br=i; br<n; ++br) if (A[br].any()) break;
   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:  $\mathcal{O}\left(n^3\right)$ ebfff6, 35 lines int matInv(vector<vector<double>>& A) { int n = sz(A); vi col(n); vector<vector<double>> tmp(n, vector<double>(n)); rep(i, 0, n) tmp[i][i] = 1, col[i] = i;rep(i,0,n) { int r = i, c = i; rep(j,i,n) rep(k,i,n)if (fabs(A[j][k]) > fabs(A[r][c])) r = j, c = k;if (fabs(A[r][c]) < 1e-12) return i; A[i].swap(A[r]); tmp[i].swap(tmp[r]); rep(j,0,n) swap(A[j][i], A[j][c]), swap(tmp[j][i], tmp[j][c]); swap(col[i], col[c]); double v = A[i][i]; rep(j,i+1,n) { double f = A[j][i] / v; A[j][i] = 0;rep(k,i+1,n) A[j][k] -= f\*A[i][k];rep(k,0,n) tmp[j][k] -= f\*tmp[i][k];rep(j,i+1,n) A[i][j] /= v; rep(j,0,n) tmp[i][j] /= v; A[i][i] = 1;for (int i = n-1; i > 0; --i) rep(j,0,i) { double v = A[j][i]; rep(k,0,n) tmp[j][k] -= v\*tmp[i][k];rep(i,0,n) rep(j,0,n) A[col[i]][col[j]] = tmp[i][j]; return n;

# Tridiagonal.h

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

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

This is useful for solving problems on the type

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

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

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

Fails if the solution is not unique.

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

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

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

# 4.4 Fourier transforms

FastFourierTransform.h

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

**Time:**  $O(N \log N)$  with  $N = |A| + |B| (\sim 1s \text{ for } N = 2^{22})$ 00ced6, 35 lines

```
typedef complex<double> C;
typedef vector<double> vd;
void fft(vector<C>& a) {
 int n = sz(a), L = 31 - \underline{builtin_clz(n)};
  static vector<complex<long double>> R(2, 1);
  static vector<C> rt(2, 1); // (^ 10% faster if double)
  for (static int k = 2; k < n; k *= 2) {
   R.resize(n); rt.resize(n);
   auto x = polar(1.0L, acos(-1.0L) / k);
   rep(i,k,2*k) rt[i] = R[i] = i&1 ? R[i/2] * x : R[i/2];
  rep(i, 0, n) \ rev[i] = (rev[i / 2] | (i & 1) << L) / 2;
  rep(i,0,n) if (i < rev[i]) swap(a[i], a[rev[i]]);
  for (int k = 1; k < n; k *= 2)
    for (int i = 0; i < n; i += 2 * k) rep(j,0,k) {
     Cz = rt[j+k] * a[i+j+k]; // (25\% faster if hand-rolled)
     a[i + j + k] = a[i + j] - z;
     a[i + j] += z;
```

```
vd conv(const vd& a, const vd& b) {
 if (a.empty() || b.empty()) return {};
 vd res(sz(a) + sz(b) - 1);
 int L = 32 - \underline{\text{builtin\_clz}(\text{sz(res)})}, n = 1 << L;
 vector<C> in(n), out(n);
  copy(all(a), begin(in));
  rep(i,0,sz(b)) in[i].imag(b[i]);
  fft(in);
  for (C& x : in) x *= x;
  rep(i, 0, n) out[i] = in[-i & (n - 1)] - conj(in[i]);
  fft(out);
  rep(i, 0, sz(res)) res[i] = imag(out[i]) / (4 * n);
 return res;
```

# FastFourierTransformMod.h

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

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

```
"FastFourierTransform.h"
typedef vector<ll> vl;
template<int M> vl convMod(const vl &a, const vl &b) {
 if (a.empty() || b.empty()) return {};
 vl res(sz(a) + sz(b) - 1);
 int B=32-__builtin_clz(sz(res)), n=1<<B, cut=int(sqrt(M));</pre>
 vector < C > L(n), R(n), outs(n), outl(n);
 rep(i, 0, sz(a)) L[i] = C((int)a[i] / cut, (int)a[i] % cut);
 rep(i, 0, sz(b)) R[i] = C((int)b[i] / cut, (int)b[i] % cut);
 fft(L), fft(R);
 rep(i,0,n) {
   int j = -i \& (n - 1);
   outl[j] = (L[i] + conj(L[j])) * R[i] / (2.0 * n);
   outs[j] = (L[i] - conj(L[j])) * R[i] / (2.0 * n) / 1i;
 fft(outl), fft(outs);
 rep(i, 0, sz(res)) {
   11 \text{ av} = 11(\text{real}(\text{outl}[i]) + .5), \text{ cv} = 11(\text{imag}(\text{outs}[i]) + .5);
   11 bv = 11(imag(out1[i])+.5) + 11(real(outs[i])+.5);
    res[i] = ((av % M * cut + bv) % M * cut + cv) % M;
 return res;
```

# NumberTheoreticTransform.h

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

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

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

```
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 - \underline{builtin\_clz(s)},
     n = 1 << B;
  int inv = modpow(n, mod - 2);
  vl L(a), R(b), out(n);
  L.resize(n), R.resize(n);
  ntt(L), ntt(R);
  rep(i,0,n)
    out[-i \& (n - 1)] = (l1)L[i] * R[i] % mod * inv % mod;
  return {out.begin(), out.begin() + s};
```

#### FastSubsetTransform.h

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

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

464cf3, 16 lines

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

# Number theory (5)

# 5.1 Modular arithmetic

ModInverse.h

**Description:** Pre-computation of modular inverses. Assumes LIM < mod and that mod is a prime. e403d4, 3 lines

```
const 11 mod = 1000000007, LIM = 200000;
11* inv = new l1[LIM] - 1; inv[1] = 1;
FOR (i, 2, LIM) inv[i] = mod - (mod / i) * inv[mod % i] % mod;
```

#### ModPow.h

b83e45, 8 lines

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

# ModLog.h

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

11 modLog(ll a, ll b, ll m) { ll n = (ll) sqrt(m) + 1, e = 1, f = 1, j = 1;unordered\_map<11, 11> A; while  $(j \le n \&\& (e = f = e * a % m) != b % m)$ A[e \* b % m] = j++;if (e == b % m) return j;  $if (\underline{gcd}(m, e) == \underline{gcd}(m, b))$ rep(i,2,n+2) if (A.count(e = e \* f % m))

## ModSum.h

return -1;

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

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

return n \* i - A[e];

5c5bc5, 16 lines

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

# ModMulLL.h

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

```
typedef unsigned long long ull:
ull modmul(ull a, ull b, ull M) {
 11 \text{ 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) {
  for (; e; b = modmul(b, b, mod), e /= 2)
   if (e & 1) ans = modmul(ans, b, mod);
  return ans:
```

# ModSqrt.h

"ModPow.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

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

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

# 5.2 Primality

FastEratosthenes.h

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

Time: LIM=1e9  $\approx 1.5s$ 6b2912, 20 lines

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

## MillerRabin.h

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

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

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

# Factor.h

19a793, 24 lines

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.
```

```
d8d98d, 18 lines
ull pollard(ull n) {
 ull x = 0, y = 0, t = 30, prd = 2, i = 1, q;
  auto f = [\&](ull x) \{ return modmul(x, x, n) + i; \};
  while (t++ % 40 || _gcd(prd, n) == 1) {
    if (x == y) x = ++i, y = f(x);
    if ((q = modmul(prd, max(x,y) - min(x,y), n))) prd = q;
    x = f(x), y = f(f(y));
 return __gcd(prd, n);
vector<ull> factor(ull n) {
 if (n == 1) return {};
 if (isPrime(n)) return {n};
 ull x = pollard(n);
 auto 1 = factor(x), r = factor(n / x);
 1.insert(l.end(), all(r));
 return 1:
```

# 5.3 Divisibility

euclid.h

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

```
ll euclid(ll a, ll b, ll &x, ll &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 < x < lcm(m, n). Assumes  $mn < 2^{62}$ . Time:  $\log(n)$ 

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

# 5.3.1 Bézout's identity

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

$$ax + by = d$$

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

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

# phiFunction.h

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

cf7d6d, 8 lines

```
const int LIM = 5000000;
int phi[LIM];

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

# 5.4 Pythagorean Triples

The Pythagorean triples are uniquely generated by

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

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

# 5.5 Primes

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

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

# 5.6 Estimates

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

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

# 5.7 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 (6)

# 6.1 Permutations

# 6.1.1 Factorial

n	1 2 3	4	5 6	7	8	9	10	
$\overline{n!}$	1 2 6	24 1	20 720	5040	40320	362880	3628800	_
						5 16		
n!	4.0e7	4.8e	8 6.2e	9 8.7e	e10 1.3e	e12 2.1e	13 3.6e14	-
n	20	25	30	40	50 1	.00 15	0 171	
n!	2e18	2e25	3e32	8e47 :	3e64 9e	e157 6e2	62 >DBL_N	ЛАХ

#### IntPerm.h

**Description:** Permutation -> integer conversion. (Not order preserving.) Integer -> permutation can use a lookup table. **Time:**  $\mathcal{O}(n)$ 

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

# **6.1.2** Cycles

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

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

# 6.1.3 Derangements

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

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

## 6.1.4 Burnside's lemma

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

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

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

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

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

# 6.2 Partitions and subsets

# **6.2.1** Partition function

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

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

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

# 6.2.2 Lucas' Theorem

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

# 6.2.3 Binomials

multinomial.h

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

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

# 6.3 General purpose numbers

# 6.3.1 Bernoulli numbers

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

Sums of powers:

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

Euler-Maclaurin formula for infinite sums:

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

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

# 6.3.2 Stirling numbers of the first kind

Number of permutations on n items with k cycles.

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

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

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

# BellmanFord FloydWarshall FordFulkerson EdmondsKarp Dinic

# 6.3.3 Eulerian numbers

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

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

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

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

# 6.3.4 Stirling numbers of the second kind

Partitions of n distinct elements into exactly k groups.

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

# 6.3.5 Bell numbers

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

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

# 6.3.6 Labeled unrooted trees

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

# 6.3.7 Catalan numbers

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

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

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

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

# Graph (7)

# 7.1 Fundamentals

# BellmanFord.h

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

```
const ll inf = LLONG MAX;
struct Ed { int a, b, w, s() { return a < b ? a : -a; }};
struct Node { ll dist = inf; int prev = -1; };
void bellmanFord(vt<Node>& nodes, vt<Ed>& eds, int s) {
 nodes[s].dist = 0;
 sort(all(eds), [](Ed a, Ed b) { return a.s() < b.s(); });</pre>
 int lim = size(nodes) / 2 + 2; // /3+100 with shuffled
 FOR (i, lim) for (Ed ed : eds) {
   Node cur = nodes[ed.a], &dest = nodes[ed.b];
   if (abs(cur.dist) == inf) continue;
   11 d = cur.dist + ed.w;
   if (d < dest.dist) {
     dest.prev = ed.a;
     dest.dist = (i < lim-1 ? d : -inf);
 FOR (i, lim) for (Ed e : eds) {
   if (nodes[e.a].dist == -inf)
     nodes[e.b].dist = -inf;
```

#### FlovdWarshall.h

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

# 7.2 Network flow

# FordFulkerson.h

**Description:** Short algo for computing maximum flows with a bounded answer.

```
int flow = 0;
while (tim++, dfs(dfs, s)) flow++;
return flow;
}
```

#### EdmondsKarp.h

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

```
template<class T> T edmondsKarp(vector<unordered map<int, T>>&
    adj, int s, int t) {
  assert(s != t);
 T flow = 0;
 vt<int> par(size(adj)), q = par;
    while (1) {
    fill(all(par), -1);
    int ptr = 1;
    q[0] = s, par[s] = 0;
       FOR (i, ptr) {
      int u = q[i];
            for (auto &[v, c] : adj[u]) {
       if (par[v] == -1 \&\& c) {
         par[v] = u, q[ptr++] = v;
          if (v == t) goto out;
        return flow;
   T inc = numeric limits<T>::max();
    for (int y = t; y != s; y = par[y])
     inc = min(inc, adj[par[y]][y]);
    flow += inc;
    for (int y = t; y != s; y = par[y]) {
     int p = par[y];
     if ((adj[p][y] -= inc) <= 0) adj[p].erase(y);</pre>
     adj[y][p] += inc;
```

## 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;
       11 c, oc;
       ll flow() { return max(oc - c, OLL); } // if you need
            flows
    vi lvl, ptr, q;
   vt<vt<Edge>> adj;
    void init(int n) {
       lvl = ptr = q = vi(n);
        adj.resize(n);
    void ae(int a, int b, ll c, ll rcap = 0) {
        adj[a].pb({b, size(adj[b]), c, c});
       adj[b].pb({a, size(adj[a]) - 1, rcap, rcap});
   11 dfs(int v, int t, ll f) {
       if (v == t || !f) return f;
```

11

# PushRelabel MinCostMaxFlow MinCut GlobalMinCut

```
for (int& i = ptr[v]; i < size(adj[v]); i++) {</pre>
        auto &[to, rev, c, _] = adj[v][i];
        if (lvl[to] == lvl[v] + 1)
            if (ll p = dfs(to, t, min(f, c))) {
                c -= p, adj[to][rev].c += p;
                return p;
    return 0;
11 calc(int s, int t) {
   11 \text{ flow} = 0; q[0] = s;
   FOR (L, 31) do { // int L=30 maybe faster for
        random data
        lvl = ptr = vi(size(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 left_of_min_cut(int a) { return lvl[a] != 0; }
```

#### PushRelabel.h

Description: Push-relabel using the highest label selection rule and the gap heuristic. Quite fast in practice. To obtain the actual flow, look at positive values only.

Time:  $\mathcal{O}\left(V^2\sqrt{E}\right)$ 

```
fc59f0, 66 lines
template<typename flow_t = long long>
struct PushRelabel {
    struct Edge {
       int to, rev;
        flow_t f, c;
    };
   vt<vt<Edge>> q;
   vt<flow t> ec:
   vt<Edge*> cur:
   vt<vt<int>> hs;
   vt<int> h;
    void init(int n) {
       q.resize(n);
       ec.resize(n);
       cur.resize(n);
       hs.resize(2 * n);
       h.resize(n);
    void ae(int s, int t, flow_t cap, flow_t rcap = 0) {
       if (s == t) return;
        g[s].push_back({t, size(g[t]), 0, cap});
        g[t].push_back({s, size(g[s]) - 1, 0, rcap});
    void add_flow(Edge& e, flow_t f) {
        Edge &back = g[e.to][e.rev];
        if (!ec[e.to] && f)
           hs[h[e.to]].push_back(e.to);
        e.f += f; e.c -= f;
       ec[e.to] += f;
       back.f -= f; back.c += f;
        ec[back.to] -= f;
```

```
flow t calc(int s, int t) {
    int v = size(q);
   h[s] = v;
    ec[t] = 1;
    vt < int > co(2 * v);
    co[0] = v - 1;
FOR (i, v) cur[i] = g[i].data();
    for(auto &e : g[s]) add_flow(e, e.c);
    if (size(hs[0]))
    for (int hi = 0; hi >= 0;) {
        int u = hs[hi].back();
       hs[hi].pop_back();
        while (ec[u] > 0) // discharge u
            if (cur[u] == g[u].data() + size(g[u])) {
                h[u] = 1e9;
                for (auto &e : q[u])
                    if (e.c && h[u] > h[e.to] + 1)
                        h[u] = h[e.to] + 1, cur[u] = &e;
                if (++co[h[u]], !--co[hi] && hi < v)</pre>
        FOR (i, v)
                        if (hi < h[i] && h[i] < v)</pre>
                             --co[h[i]], h[i] = v + 1;
                hi = h[u];
            } else if (cur[u]->c && h[u] == h[cur[u]->to] +
                add_flow(*cur[u], min(ec[u], cur[u]->c));
            else ++cur[u];
        while (hi \geq= 0 && hs[hi].empty()) --hi;
    return -ec[s];
bool leftOfMinCut(int a) { return h[a] >= size(q); }
```

## MinCostMaxFlow.h

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

**Time:**  $\mathcal{O}(FE\log(V))$  where F is max flow.  $\mathcal{O}(VE)$  for setpi. (Cee40, 79) lines

```
#include <bits/extc++.h>
struct MCMF {
   struct edge {
       int from, to, rev;
       11 cap, cost, flow;
   };
   int N;
   vt<vt<edge>> ed;
   vt<int> seen;
   vt<ll> dist, pi;
   vt<edge*> par;
   MCMF(int N) : N(N), ed(N), seen(N), dist(N), pi(N), par(N)
        {}
   void ae(int from, int to, ll cap, ll cost) {
       if (from == to) return;
       ed[from].push back(edge{ from, to, size(ed[to]),
            cap, cost, 0 });
       ed[to].push_back(edge{ to, from, size(ed[from]) - 1,
            0, -cost, 0 });
   void path(int s) {
       fill(all(seen), 0);
       fill(all(dist), INF);
```

dist[s] = 0; ll di;

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

#### MinCut.h

};

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

# GlobalMinCut.h

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

```
d4d36f, 21 lines
pair<int, vt<int>> globalMinCut(vt<vt<int>> mat) {
 pair<int, vt<int>> best = {INT_MAX, {}};
 int n = size(mat);
```

# hopcroftKarp WeightedMatching Blossom SCC

```
vt<vt<int>> co(n);
FOR (i, n) co[i] = \{i\};
FOR(ph, 1, n) {
 vt < int > w = mat[0];
  size_t s = 0, t = 0;
 FOR (it, 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();
   FOR (i, 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]));
 FOR (i, n) mat[s][i] += mat[t][i];
 FOR (i, n) mat[i][s] = mat[s][i];
 mat[0][t] = INT_MIN;
return best;
```

# 7.3 Matching

hopcroftKarp.h

**Description:** Fast incremental bipartite matching. Zero-indexed.

Usage: {operator[]} for the pair of right node i, {n} is the size of the rhs, {add(v)} to add adjacency list of node on lhs

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

0821c1, 39 lines

```
struct Matching : vt<int> {
    vt<vt<int>> adj;
    vt<int> rank, low, pos, vis, seen;
    int k{0};
    // n = size \ of \ rhs
    Matching(int n) : vt < int > (n, -1), rank(n) {}
    bool add(vt<int> vec) {
        adj.pb(std::move(vec));
        low.pb(0); pos.pb(0); vis.pb(0);
        if (!adj.back().empty()) {
            int i = k;
        nxt:
            seen.clear();
            if (dfs(size(adj)-1, ++k-i)) return 1;
            for (auto &v : seen) for (auto &e : adj[v])
                if (rank[e] < 1e9 && vis[at(e)] < k)</pre>
                    goto nxt;
            for (auto &v : seen) {
                low[v] = 1e9;
        for (auto &w : adj[v]) rank[w] = 1e9;
        return 0;
    bool dfs(int v, int q) {
        if (vis[v] < k) vis[v] = k, seen.pb(v);
        while (low[v] < q) {
            int e = adj[v][pos[v]];
            if (at(e) != v && low[v] == rank[e]) {
                rank[e]++;
                if (at(e) == -1 || dfs(at(e), rank[e]))
                    return at(e) = v, 1;
            } else if (++pos[v] == size(adj[v])) {
                pos[v] = 0; low[v]++;
        return 0;
};
```

```
WeightedMatching.h
```

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

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

2e27e7, 31 lines pair<int, vi> hungarian(const vt<vi> &a) { if (a.empty()) return {0, {}}; int n = size(a) + 1, m = size(a[0]) + 1; vi u(n), v(m), p(m), ans(n-1);FOR (i, 1, n) { p[0] = i;int j0 = 0;vi dist(m, INT\_MAX), pre(m, -1); vector<bool> done(m + 1); done[j0] = true; int i0 = p[j0], j1, delta = INT\_MAX; FOR (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> FOR (j, m) { if (done[j]) u[p[j]] += delta, v[j] -= delta; else dist[i] -= delta; j0 = j1;} while (p[j0]); while (j0) { // update alternating path int j1 = pre[j0]; p[j0] = p[j1], j0 = j1;FOR (j, 1, m) if (p[j]) ans[p[j] - 1] = j - 1; return {-v[0], ans}; // min cost

#### Blossom.h

Description: Matching for general graphs. 1-indexed!

Time:  $\mathcal{O}(NM)$ 0b8afb, 58 lines struct Blossom { int n, h, t, cnt; vpi edges; vi vis, q, mate, col, fa, pre, he; void ae(int u, int v) { assert (u && v); edges.pb( $\{he[u], v\}$ ); he[u] = size(edges) - 1; edges.pb( $\{he[v], u\}$ ); he[v] = size(edges) - 1; inline int get(int u) { return fa[u] == u ? u : fa[u] = get (fa[u]); } void aug(int u, int v) { for (int p; u; u = p, v = pre[p]) p = mate[v], mate[mate[u] = v] = u; void init(int \_n) {  $n = _n;$ vis = q = mate = col = fa = pre = he = vi(n + 1);int lca(int u, int v) { for (cnt++;; u = pre[mate[u]]) { if (v) swap(u, v); if (vis[u = get(u)] == cnt) return u; vis[u] = cnt;

```
void blo(int u, int v, int f) {
        for (int p; get(u) != f; v = p, u = pre[p]) {
            p = mate[u]; pre[u] = v; fa[u] = fa[p] = f;
            if (col[p] != 1) col[q[++t] = p] = 1;
   bool bfs(int u) {
       FOR (i, 1, n + 1) col[i] = 0, fa[i] = i;
       h = 0; q[t = 1] = u; col[u] = 1;
       while (h != t) {
            int x = q[++h];
            for (int i = he[x]; i; i = edges[i].f) {
                int y = edges[i].s;
                if (!col[y]) {
                    if (!mate[y]) { aug(y, x); return 1; }
                    pre[y] = x;
                    col[y] = 2;
                    col[q[++t] = mate[y]] = 1;
                } else if (col[y] == 1 && get(x) != get(y)) {
                    int p = lca(x, y);
                    blo(x, y, p);
                    blo(y, x, p);
       }
       return 0;
    int solve() {
       int ans = 0;
       FOR (i, 1, n + 1) if (!mate[i]) ans += bfs(i);
        return ans;
};
```

# 7.4 DFS algorithms

Description: Finds strongly connected components in a directed graph. comps is top-sorted.

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

12

```
ae000b, 30 lines
struct SCC {
    vt<vt<int>> adj, radj;
    vt<int> todo, seen, comp, comps; // comps is top sorted
    void init(int _n) {
        n = _n;
        adj = radj = vt<vt<int>>(n);
        comp.resize(n, -1);
        seen.resize(n);
    void ae(int u, int v) {
        adj[u].pb(v);
        radj[v].pb(u);
    void dfs(int u) {
        if (seen[u]++) return;
        for (int v : adj[u]) dfs(v);
        todo.pb(u);
    void rdfs(int u, int w) {
        comp[u] = w;
        for (int v : radj[u]) if (comp[v] == -1) rdfs(v, w);
    void gen() {
        FOR (i, n) dfs(i);
        reverse(all(todo));
        for (int u : todo) if (comp[u] == -1)
            rdfs(u, u), comps.pb(u);
```

# BiconnectedComponents.h

**Description:** Finds all biconnected components in an undirected graph. 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. Note that degree 0 nodes are not considered components.

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

fb417e, 47 lines

```
struct BCC {
   int n, m, t;
   vt<vt<pair<int, int>>> adj;
  vector<pair<int, int>> edges;
   vt<vt<int>> comps; // lists of edges of bcc
   vt<int> tin, stk, is_art, is_bridge;
    void init(int _n, vt<pair<int, int>> &_edges) {
       n = _n;
    edges = _edges;
    m = size(edges);
       adj.resize(n);
    FOR (i, m) {
     auto [u, v] = edges[i];
     adj[u].pb({v, i});
     adj[v].pb({u, i});
   t = 0;
        tin = is_art = vt<int>(n);
    is_bridge.resize(m);
       FOR (u, n) if (!tin[u]) dfs(u, -1);
    // if we include bridges as 2-node bcc
   FOR (i, m) if (is_bridge[i]) comps.pb({i});
    int dfs(int u, int par) {
        int me = tin[u] = ++t, dp = me;
        for (auto [v, ei] : adj[u]) if (ei != par) {
            if (tin[v]) {
                dp = min(dp, tin[v]);
                if (tin[v] < me)</pre>
                    stk.push_back(ei);
            } else {
                int si = size(stk), up = dfs(v, ei);
                dp = min(dp, up);
                if (up == me) {
          is art[u] = 1;
                    stk.pb(ei);
                    comps.pb({si + all(stk)});
                    stk.resize(si);
                } else if (up < me) stk.push back(ei);
                else { is_bridge[ei] = 1; }
        return dp;
};
```

# BlockCutTree.h

Description: First, use BiconnectedComponents to locate VERTEX components (including degree 0 nodes). To build a block cut tree, make a bipartite graph: Put all the normal nodes on the left, and make a new node for each bcc on the right. Draw edges from normal nodes to BCC that contain them. Note that the graph may be disconnected.

2sat.h

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

Usage: TwoSat ts(number of boolean variables); ts.either(0,  $\sim$ 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 = 0;
   vt<pi> edges;
    void init(int _n) { n = _n; }
   int add() { return n++; }
   void either(int x, int y) { // x \mid y
       x = max(2 * x, -1 - 2 * x); // \sim (2 * x)
       y = \max(2 * y, -1 - 2 * y); // \sim (2 * y)
        edges.pb({x, y});
   void implies (int x, int y) { either (\sim x, y); }
   void force(int x) { either(x, x); }
   void exactly_one(int x, int y) {
        either (x, y), either (\sim x, \sim y);
   void tie(int x, int y) {
        implies (x, y), implies (\sim x, \sim y);
   void nand(int x, int y) { either(\sim x, \sim y); }
   void at most one(const vt<int>& li) {
       if (size(li) <= 1) return;
       int cur = \simli[0]:
       FOR (i, 2, size(li)) {
            int next = add();
            either(cur, ~li[i]);
            either(cur,next);
            either(~li[i], next);
            cur = ~next;
        either(cur, ~li[1]);
   vt<bool> solve() {
       SCC scc;
        scc.init(2 * n);
        for(auto& e : edges) {
            scc.ae(e.f ^ 1, e.s);
            scc.ae(e.s ^ 1, e.f);
       scc.gen();
        reverse(all(scc.comps)); // reverse topo order
        for (int i = 0; i < 2 * n; i += 2)
            if (scc.comp[i] == scc.comp[i ^ 1]) return {};
        vt < int > tmp(2 * n);
        for (auto i : scc.comps) {
            if (!tmp[i]) tmp[i] = 1, tmp[scc.comp[i ^ 1]] = -1;
        vt < bool > ans (n);
       FOR (i, n) ans [i] = tmp[scc.comp[2 * i]] == 1;
        return ans;
```

EulerWalk.h

**Description:** Eulerian undirected/directed path/cycle algorithm. For edges, push the edge u came from instead of current node. First, the graph (after removing directivity) must be connected. For undirected graphs, a tour exists when all nodes have even degree. For directed graphs, a tour exists when all nodes have equal in and out degree. For trails, the condition is the same as if you added an edge from  $t \rightarrow s$ . Time:  $\mathcal{O}(V+E)$ 

```
int n, m;
vt<vt<pair<int, int>>> adi;
vt<int> ret, used;
void dfs(int u) {
    while (adj[u].size()) {
        auto [v, ei] = adj[u].back();
        adi[u].pop back();
        if (used[ei]++) continue;
        dfs(v);
    ret.push_back(u);
```

# 7.5 Heuristics

MaximumClique.h

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

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

f7c0bc, 49 lines

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

# MaximumIndependentSet LCA VirtualTree HLD Point

```
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-

# Trees

#### LCA.h

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

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

"../data-structures/RMQ.h" bf464a, 20 lines struct LCA { int t = 0: vi time, path, ret; RMQ<int> rmq; LCA(vt<vi>& adj) : time(size(adj)), rmq((dfs(0, -1, adj), ret void dfs(int u, int p, vt<vi> &adj) { time[u] = t++;for (int v : adj[u]) if (v != p) { path.push\_back(u), ret.push\_back(time[u]); dfs(v, u, adi); int operator()(int u, int v) { if (u == v) return u; tie(u, v) = minmax(time[u], time[v]);

#### VirtualTree.h

return path[rmq.query(u, v)];

Description: Computes virtual tree. pos is inorder dfs time, and returns pairs of (par, child).

```
Time: \mathcal{O}\left(|S|\log|S|\right)
"LCA.h"
                                                         4ff13b, 14 lines
// pos is dfs time
// pairs of {ancestor, child}
vt<pl> virtual_tree(vt<ll>& nodes) {
    auto cmp = [&] (ll u, ll v) { return pos[u] < pos[v]; };</pre>
    sort(all(nodes), cmp);
    int sz = size(nodes);
    FOR (i, sz - 1) nodes.pb(lca(nodes[i], nodes[i + 1]));
    sort (all (nodes), cmp);
    nodes.erase(unique(all(nodes)), nodes.end());
    vt<pl> res;
    FOR (i, (int) size(nodes) - 1)
    res.pb({lca(nodes[i], nodes[i + 1]), nodes[i + 1]});
    return res;
```

#### HLD.h

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

```
to the segtree default. Root must be 0.
Time: \mathcal{O}\left((\log N)^2\right)
"../data-structures/LazySegmentTree.h"
                                                       812f05, 69 lines
template<bool in_edges> struct HLD {
   int n;
    vt<vt<int>> adi;
   vt<int> par, root, depth, sz, pos;
    int time;
    SegTree tree;
    void ae(int u, int v) {
       adj[u].pb(v);
        adj[v].pb(u);
   void dfs_sz(int u) {
        sz[u] = 1;
        for (int& v : adj[u]) {
            par[v] = u;
            depth[v] = depth[u] + 1;
            adj[v].erase(find(all(adj[v]), u));
            dfs sz(v);
            sz[u] += sz[v];
            if (sz[v] > sz[adj[u][0]]) swap(v, adj[u][0]);
   void dfs_hld(int u) {
        pos[u] = time++;
        for (int& v : adj[u]) {
            root[v] = (v == adj[u][0] ? root[u] : v);
            dfs_hld(v);
    void init(int _n) {
       n = _n;
        adj.resize(n);
        par = root = depth = sz = pos = vt<int>(n);
   void gen(int r = 0) {
       par[r] = depth[r] = time = 0;
       dfs sz(r);
        root[r] = r;
        dfs hld(r);
        tree.init(n);
   int lca(int u, int v) {
        while (root[u] != root[v]) {
            if (depth[root[u]] > depth[root[v]]) swap(u, v);
            v = par[root[v]];
        return depth[u] < depth[v] ? u : v;</pre>
   template <class Op>
   void process(int u, int v, Op op) {
        for (;; v = par[root[v]]) {
            if (pos[u] > pos[v]) swap(u, v);
```

if (root[u] == root[v]) break;

op(pos[root[v]], pos[v] + 1);

op(pos[u] + in\_edges, pos[v] + 1);

process(u, v, [&] (int 1, int r) {

void upd(int u, int v, ll upd) {

tree.upd(1, r, upd);

```
});
11 query(int u, int v) {
    11 \text{ res} = 0;
    process(u, v, [&] (int l, int r) {
       res = res + tree.querv(1, r);
    });
    return res;
```

# Math

# 7.7.1 Number of Spanning Trees

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

14

# 7.7.2 Erdős–Gallai theorem

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

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

# Geometry (8)

# 8.1 Geometric primitives

# Point.h

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

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

15

b70280, 11 lines

```
P rotate(double a) const {
        return P(x * cos(a) - y * sin(a), x * sin(a) + y * cos(a)
             a));
    friend ostream& operator<<(ostream& os, P p) {</pre>
        return os << "(" << p.x << "," << p.v << ")";
};
```

# lineDistance.h

#### Description:

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



3b34c4, 4 lines

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

# SegmentDistance.h

Description:

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

Usage: Point < double > a, b(2,2), p(1,1);



579797, 5 lines

bool on\_segment = seg\_dist(a,b,p) < 1e-10;

double seg dist(P& s, P& e, P& p) { if (s == e) return (p - s).dist(); auto d = (e - s).dist2(), t = min(d, max(.0, (p - s).dot(e s))); return ((p - s) \* d - (e - s) \* t).dist() / d;

# SegmentIntersection.h

#### Description:

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



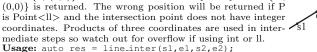
```
Usage: vector <P > inter = seq_inter(s1,e1,s2,e2);
if (size(inter) == 1)
cout << "segments intersect at " << inter[0] << endl;</pre>
                                                         bd6e14, 13 lines
"Point.h", "OnSegment.h"
```

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

## lineIntersection.h

#### Description:

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



```
if (res.first == 1)
cout << "intersection point at " << res.second << endl;</pre>
"Point.h"
                                                       b0d826, 8 lines
template<class P>
pair<int, P> line_inter(P s1, P e1, P s2, P e2) {
    auto d = (e1 - s1).cross(e2 - s2);
    if (d == 0) // if parallel
        return \{-(s1.cross(e1, s2) == 0), P(0, 0)\};
    auto p = s2.cross(e1, e2), q = s2.cross(e2, s1);
    return \{1, (s1 * p + e1 * q) / d\};
```

#### sideOf.h

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

```
Usage: bool left = side_of(p1,p2,q) ==1;
"Point.h"
                                                      9e71fb, 9 lines
template<class P>
int side_of(P s, P e, P p) { return sqn(s.cross(e, p)); }
template<class P>
int side of (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 (seg\_dist(s,e,p) < epsilon) instead when using Point < double >.

```
0c1f75, 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;
```

# linearTransformation.h Description:

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



"Point.h" 8e73be, 6 lines typedef Point < double > P; P linear\_transformation(const P& p0, const P& p1, const P& q0, const P& q1, const P& r) { P dp = p1-p0, dq = q1-q0, num(dp.cross(dq), dp.dot(dq)); return q0 + P((r-p0).cross(num), (r-p0).dot(num))/dp.dist2();

# Angle.h

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

```
Usage: vector<Angle> v = \{w[0], w[0].t360() ...\}; // sorted
int j = 0; rep(i,0,n) { while (v[j] < v[i].t180()) ++j; }
// sweeps j such that (j-i) represents the number of positively
oriented triangles with vertices at 0 and i
struct Angle {
  int x, v;
  int t:
  Angle(int x, int y, int t = 0) : x(x), y(y), t(t) {}
  Angle operator-(Angle b) const { return {x - b.x, y - b.y, t}
  int half() const {
    assert(x || y);
    return y < 0 \mid | (y == 0 \&\& x < 0);
  Angle t90() const { return \{-y, x, t + (half() \&\& x >= 0)\}; \}
  Angle t180() const { return \{-x, -y, t + half()\}; }
  Angle t360() const { return \{x, y, t + 1\}; \}
bool operator<(Angle a, Angle b) {</pre>
  // add a. dist2() and b. dist2() to also compare distances
  return make_tuple(a.t, a.half(), a.y * (11) b.x) <</pre>
         make tuple(b.t, b.half(), a.x \star (ll) b.v);
 / Given two points, this calculates the smallest angle between
 // them, i.e., the angle that covers the defined line segment.
pair < Angle, Angle > segment Angles (Angle a, Angle b) {
  if (b < a) swap(a, b);
  return (b < a.t180() ?
          make_pair(a, b) : make_pair(b, a.t360()));
Angle operator+(Angle a, Angle b) { // point a + vector b
  Angle r(a.x + b.x, a.y + b.y, a.t);
  if (a.t180() < r) r.t--;
  return r.t180() < a ? r.t360() : r;
Angle angleDiff(Angle a, Angle b) { // angle b- angle a
  int tu = b.t - a.t; a.t = b.t;
  return {a.x * b.x + a.y * b.y, a.x * b.y - a.y * b.x, tu - (b
        < a);
```

## 8.2 Circles

"Point.h"

#### CircleIntersection.h

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

```
using P = Point <db>;
bool circle inter(P a, P b, db r1, db r2, pair<P, P> *out) {
  if (a == b) { assert(r1 != r2); return false; }
  P \text{ vec} = b - a;
  db d2 = vec.dist2(), sum = r1 + r2, dif = r1 - r2,
          p = (d2 + r1 * r1 - r2 * r2) / (d2 * 2), h2 = r1 * r1
                - p * p * d2;
  if (sum * sum < d2 || dif * dif > d2) return false;
  P \text{ mid} = a + \text{vec} * p, \text{ per} = \text{vec.perp}() * \text{sqrt}(\text{fmax}(0, h2) / d2)
  *out = {mid + per, mid - per};
  return true:
```

# CircleTangents.h

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

# CircleLine.h

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

#### CirclePolygonIntersection.h

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

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

```
"../../content/geometry/Point.h"
                                                      e876aa, 19 lines
typedef Point < double > P;
#define arg(p, q) atan2(p.cross(q), p.dot(q))
double circlePoly(P c, double r, vector<P> ps) {
  auto tri = [&](P p, P q) {
   auto r2 = r * r / 2;
   P d = q - p;
    auto a = d.dot(p) / d.dist2(), b = (p.dist2() - r * r) / d.
        dist2();
    auto det = a * a - b;
    if (det <= 0) return arg(p, q) * r2;</pre>
    auto s = max(0., -a - sqrt(det)), t = min(1., -a + sqrt(det))
        ));
    if (t < 0 \mid | 1 \le s) return arg(p, q) * r2;
   P u = p + d * s, v = q + d * (t - 1);
   return arg(p, u) * r2 + u.cross(v) / 2 + arg(v, q) * r2;
  };
  auto sum = 0.0;
  FOR (i, size(ps))
    sum += tri(ps[i] - c, ps[(i + 1) % size(ps)] - c);
  return sum;
```

# circumcircle.h

#### Description:

"Point.h"

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



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

# MinimumEnclosingCircle.h

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

```
"circumcircle.h"
                                                     256373, 17 lines
pair<P, double> mec(vector<P> ps) {
 shuffle(all(ps), mt19937(time(0)));
 P \circ = ps[0];
 double r = 0, EPS = 1 + 1e-8;
 FOR (i, size(ps)) if ((o - ps[i]).dist() > r * EPS) {
   o = ps[i], r = 0;
   FOR (j, i) if ((o - ps[j]).dist() > r * EPS) {
     o = (ps[i] + ps[j]) / 2;
      r = (o - ps[i]).dist();
     FOR (k, j) if ((o - ps[k]).dist() > r * EPS) {
       o = cc_center(ps[i], ps[j], ps[k]);
       r = (o - ps[i]).dist();
   }
 return {o, r};
```

# 8.3 Polygons

# InsidePolygon.h

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

```
Usage: vector<P> v = {P{4,4}, P{1,2}, P{2,1}};
bool in = in_polygon(v, P{3, 3}, false);
Time: \mathcal{O}(n)
```

"Point.h", "OnSegment.h", "SegmentDistance.h"

# 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" c9f086, 6 lines template<class T>
```

```
T polygon_area(vector<Point<T>>& v) {
   T a = v.back().cross(v[0]);
   FOR (i, size(v) - 1) a += v[i].cross(v[i + 1]);
   return a;
}
```

# PolygonCenter.h

**Description:** Returns the center of mass for a polygon.

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

# 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));



#### PolygonUnion.h

b915a1, 11 lines

**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"
                                                     b287c2, 33 lines
typedef Point < db > P;
double rat(P a, P b) { return sqn(b.x) ? a.x/b.x : a.y/b.y; }
double polyUnion(vector<vector<P>>& poly) {
 double ret = 0:
 FOR (i, size(poly)) FOR (v, size(poly[i])) {
    P A = poly[i][v], B = poly[i][(v + 1) % sz(poly[i])];
    vector<pair<double, int>> segs = {{0, 0}, {1, 0}};
    FOR (j, size(poly)) if (i != j) {
      rep(u, 0, sz(polv[i])) {
        P C = poly[j][u], D = poly[j][(u + 1) % sz(poly[j])];
        int sc = side_of(A, B, C), sd = side_of(A, B, D);
        if (sc != sd) {
          db sa = C.cross(D, A), sb = C.cross(D, B);
          if (\min(sc, sd) < 0)
            segs.emplace_back(sa / (sa - sb), sgn(sc - sd));
        } else if (!sc && !sd && j < i && sgn((B - A).dot(D-C))</pre>
          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;
 FOR (j, 1, size(segs)) {
   if (!cnt) sum += segs[j].first - segs[j - 1].first;
   cnt += segs[j].second;
 ret += A.cross(B) * sum;
return ret / 2;
```

## ConvexHull.h

#### Description:

Returns a vector of the points of the convex hull in 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"
                                                       5e9915, 12 lines
vt<P> convex_hull(vt<P> pts) {
    if (size(pts) <= 1) return pts;</pre>
    sort(all(pts));
    vector<P> h(size(pts)+1);
    int s = 0, t = 0;
    for (int it = 2; it--; s = --t, reverse(all(pts)))
        for (P p : pts) {
            while (t \ge s + 2 \&\& h[t - 2].cross(h[t - 1], p) \le
            h[t++] = p;
    return \{h.begin(), h.begin() + t - (t == 2 && h[0] == h[1])
        };
```

#### HullDiameter.h

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

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

```
"Point.h"
                                                      8279d4, 12 lines
typedef Point<ll> P;
array<P, 2> hull_diameter(vector<P> s) {
  int n = size(s), j = n < 2 ? 0 : 1;
  pair<11, array<P, 2 >> res({0, {s[0], s[0]}});
  FOR (i, j)
    for (;; j = (j + 1) % n) {
      res = \max(\text{res}, \{(s[i] - s[j]).dist2(), \{s[i], s[j]\}\});
      if ((s[(j+1) % n] - s[j]).cross(s[i+1] - s[i]) >= 0)
       break;
  return res.second;
```

#### PointInsideHull.h

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

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

```
"Point.h", "sideOf.h", "OnSegment.h"
                                                        c366a3, 13 lines
using P = Point<11>;
bool in_hull(const vector<P>& 1, P p, bool strict = true) {
    int a = 1, b = size(1) - 1, r = !strict;
    if (size(1) < 3) return r && on_segment(1[0], 1.back(), p);</pre>
    if (side_of(1[0], 1[a], 1[b]) > 0) swap(a, b);
```

```
if (side_of(1[0], 1[a], p) >= r || side_of(1[0], 1[b], p)
     \langle = -r \rangle
    return false;
while (abs(a - b) > 1) {
    int c = (a + b) / 2;
    (side_of(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"
                                                     bf84fc, 39 lines
#define cmp(i,j) sgn(dir.perp().cross(poly[(i) % n] - poly[(j)
    % n]))
#define extr(i) cmp(i + 1, i) >= 0 && cmp(i, i - 1 + n) < 0
template <class P> int extrVertex(vector<P>& poly, P dir) {
 int n = size(poly), lo = 0, hi = n;
 if (extr(0)) return 0;
 while (lo + 1 < hi) {
   int m = (lo + hi) / 2;
   if (extr(m)) return m;
   int 1s = cmp(1o + 1, 1o), ms = cmp(m + 1, m);
    (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;
 FOR (i, 2) {
   int lo = endB, hi = endA, n = size(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;
```

# 8.4 Misc. Point Set Problems

## ClosestPair.h

Description: Finds the closest pair of points. Time:  $\mathcal{O}(n \log n)$ 

```
be22ff, 16 lines
pair<P, P> closest(vector<P> v) {
```

```
assert(size(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;
```

#### kdTree.h

# **Description:** KD-tree (2d)

ad9b75, 53 lines

```
using P = array<int, 2>;
struct Node {
    \#define sq(x) (x) * (x)
   P lo, hi;
    struct Node *lc, *rc;
    11 dist2(const P &a, const P &b) const {
        return 111 * _sq(a[0] - b[0]) + 111 * _sq(a[1] - b[1]);
   ll dist2(P &p) {
    #define _loc(i) (p[i] < lo[i] ? lo[i] : (p[i] > hi[i] ? hi[
        i] : p[i]))
    return dist2(p, {_loc(0), _loc(1)});
       // ll res = 0;
        // FOR (i, 2) {
             ll \ tmp = (p[i] < lo[i] ? lo[i] - p[i] : 0) + (hi)
             [i] < p[i] ? p[i] - hi[i] : 0);
               res \neq = tmp * tmp;
        // return res;
    template<class ptr>
   Node (ptr 1, ptr r, int d) : lc(0), rc(0) {
       lo = {inf, inf}, hi = {-inf, -inf};
       for (ptr p = 1; p < r; p++) {
            FOR (i, 2) lo[i] = min(lo[i], (*p)[i]), hi[i] = max
                 (hi[i], (*p)[i]);
       if (r - 1 == 1) return;
       ptr m = 1 + (r - 1) / 2;
       nth_element(l, m, r, [&] (auto a, auto b) { return a[d]
             < b[d]; });
       lc = new Node(1, m, d ^ 1);
       rc = new Node(m, r, d^1);
   void search(P p, ll &best) {
       if (lc) { // rc will also exist
            11 dl = 1c -> dist2(p), dr = rc -> dist2(p);
            if (dl > dr) swap(lc, rc), dr = dl;
            lc->search(p, best);
            if (dr < best) rc->search(p, best);
       } else best = min(best, dist2(p, lo));
    // fill pg with k infinities for nearest k points
```

void search(P p, priority\_queue<11> &pq) {

# FastDelaunay.h

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

```
t[0][1],\,t[0][2],\,\bar{t}[1][0],\,\dots\}, all counter-clockwise.
Time: \mathcal{O}(n \log n)
typedef Point<11> P;
typedef struct Quad* Q;
typedef __int128_t 111; // (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; }
  O& r() { return rot->rot; }
  Q prev() { return rot->o->rot; }
  O 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]);
    0 c = side ? connect(b, a) : 0;
    return {side < 0 ? c->r() : a, side < 0 ? c : b->r() };
\#define H(e) e \rightarrow F(), e \rightarrow p
\#define\ valid(e)\ (e->F().cross(H(base)) > 0)
  Q A, B, ra, rb;
  int half = sz(s) / 2;
  tie(ra, A) = rec({all(s) - half});
  tie(B, rb) = rec({sz(s) - half + all(s)});
  while ((B->p.cross(H(A)) < 0 \&& (A = A->next())) | |
```

```
(A->p.cross(H(B)) > 0 && (B = B->r()->o)));
 Q base = connect(B->r(), A);
 if (A->p == ra->p) ra = base->r();
 if (B->p == rb->p) rb = base;
#define DEL(e, init, dir) Q e = init->dir; if (valid(e)) \
   while (circ(e->dir->F(), H(base), e->F())) { }
     Q t = e->dir; \
     splice(e, e->prev()); \
     splice(e->r(), e->r()->prev()); \
     e->o = H; H = e; e = t; \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 {};</pre>
 Q e = rec(pts).first;
 vector<Q> q = {e};
 int qi = 0;
 while (e->o->F().cross(e->F(), e->p) < 0) e = e->o;
#define ADD { Q c = e; do { c->mark = 1; pts.push_back(c->p); \
 q.push\_back(c->r()); c = c->next(); } while (c != e); }
 ADD; pts.clear();
 while (qi < sz(q)) if (!(e = q[qi++]) \rightarrow mark) ADD;
 return pts;
```

# 8.5 3D

# 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 = 0) and t3 (0 = 0) from z axis ((0 = 0)). 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 points in the x direction and d\*radius is the total distance between the points.

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

# Strings (9)

#### KMP.h

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

```
int g = p[i-1];
while (g && s[i] != s[g]) g = p[g-1];
p[i] = g + (s[i] == s[g]);
}
return p;
}

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

#### Zfunc.h

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

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

#### Manacher.h

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

```
Time: O(N)

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

#### 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;
}
```

# SuffixArray.h

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# SuffixTree Hashing AhoCorasick

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

Time:  $O(n \log n)$ 

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

# 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=1[v];
    if (q==-1 || c==toi(a[q])) q++; else {
     l[m+1]=i; p[m+1]=m; l[m]=l[v]; r[m]=q;
     p[m]=p[v]; t[m][c]=m+1; t[m][toi(a[q])]=v;
     l[v]=q; p[v]=m; t[p[m]][toi(a[l[m]])]=m;
     v=s[p[m]]; q=l[m];
     while (q < r[m]) \{ v = t[v][toi(a[q])]; q + = r[v] - l[v]; \}
     if (q==r[m]) s[m]=v; else s[m]=m+2;
      q=r[v]-(q-r[m]); m+=2; goto suff;
  SuffixTree(string a) : a(a) {
    fill(r,r+N,sz(a));
   memset(s, 0, sizeof s);
   memset(t, -1, sizeof t);
    fill(t[1],t[1]+ALPHA,0);
    s[0] = 1; 1[0] = 1[1] = -1; r[0] = r[1] = p[0] = p[1] = 0;
    rep(i,0,sz(a)) ukkadd(i, toi(a[i]));
```

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

# Hashing.h

```
Description: Self-explanatory methods for string hashing.
```

```
// Arithmetic mod 2^64-1. 2x slower than mod 2^64 and more
// code, but works on evil test data (e.g. Thue-Morse, where
// ABBA... and BAAB... of length 2^10 hash the same mod 2^64).
// "typedef ull H;" instead if you think test data is random,
// or work mod 10^9+7 if the Birthday paradox is not a problem.
typedef uint64_t ull;
struct H {
 ull x; H(ull x=0) : x(x) {}
 H operator+(H o) { return x + o.x + (x + o.x < x); }
  H operator-(H o) { return *this + ~o.x; }
 H 	ext{ operator} * (H 	ext{ o}) { auto } m = (\underline{uint128}\_t) x * o.x;
    return H((ull)m) + (ull)(m >> 64); }
  ull get() const { return x + !\sim x; }
 bool operator==(H o) const { return get() == o.get(); }
 bool operator<(H o) const { return get() < o.get(); }</pre>
static const H C = (11)1e11+3; // (order \sim 3e9; random also ok)
struct HashInterval {
 vector<H> ha, pw;
 HashInterval(string& str) : ha(sz(str)+1), pw(ha) {
    pw[0] = 1;
    rep(i, 0, sz(str))
     ha[i+1] = ha[i] * C + str[i],
      pw[i+1] = pw[i] * C;
 H hashInterval(int a, int b) { // hash [a, b)
    return ha[b] - ha[a] * pw[b - a];
};
vector<H> getHashes(string& str, int length) {
 if (sz(str) < length) return {};</pre>
 H h = 0, pw = 1;
 rep(i,0,length)
   h = h * C + str[i], pw = pw * C;
  vector<H> ret = {h};
 rep(i,length,sz(str)) {
   ret.push_back(h = h * C + str[i] - pw * str[i-length]);
 return ret;
H hashString(string& s){H h{}; for(char c:s) h=h*C+c;return h;}
```

## AhoCorasick.h

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

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

```
struct AhoCorasick {
 enum {alpha = 26, first = 'A'}; // change this!
 struct Node {
    // (nmatches is optional)
   int back, next[alpha], start = -1, end = -1, nmatches = 0;
   Node(int v) { memset(next, v, sizeof(next)); }
 vector<Node> N;
 vi backp;
 void insert(string& s, int j) {
   assert(!s.empty());
   int n = 0:
   for (char c : s) {
     int& m = N[n].next[c - first];
     if (m == -1) { n = m = sz(N); N.emplace_back(-1); }
     else n = m;
   if (N[n].end == -1) N[n].start = j;
   backp.push_back(N[n].end);
   N[n].end = j;
   N[n].nmatches++;
 AhoCorasick(vector<string>& pat) : N(1, -1) {
   rep(i,0,sz(pat)) insert(pat[i], i);
   N[0].back = sz(N);
   N.emplace_back(0);
   queue<int> q:
    for (q.push(0); !q.empty(); q.pop()) {
     int n = q.front(), prev = N[n].back;
     rep(i,0,alpha) {
       int &ed = N[n].next[i], y = N[prev].next[i];
       if (ed == -1) ed = y;
       else {
         N[ed].back = y;
          (N[ed].end == -1 ? N[ed].end : backp[N[ed].start])
           = N[y].end;
         N[ed].nmatches += N[y].nmatches;
         q.push(ed);
 vi find(string word) {
   int n = 0;
   vi res; // ll count = 0;
   for (char c : word) {
     n = N[n].next[c - first];
     res.push_back(N[n].end);
     // count += N[n]. nmatches;
   return res;
 vector<vi> findAll(vector<string>& pat, string word) {
   vi r = find(word);
   vector<vi> res(sz(word));
   rep(i, 0, sz(word)) {
```

```
int ind = r[i];
   while (ind != -1) {
      res[i - sz(pat[ind]) + 1].push_back(ind);
      ind = backp[ind];
    }
   return res;
}
```

# Various (10)

# 10.1 Intervals

# IntervalContainer.h

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

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

#### IntervalCover.h

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

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

9e9d8d, 19 lines

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

```
return R;
}
ConstantIntervals.h
Description: Split a monotone function on [from, to) into a
```

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

 $\begin{array}{lll} \textbf{Usage:} & \text{constantIntervals} (0, \text{ sz}(\text{v}), \text{ [\&] (int x) \{return v[x];\}, } \\ \text{[\&] (int lo, int hi, T val)} \\ \textbf{Time:} & \mathcal{O}\left(k\log\frac{n}{k}\right) & \text{753a4c, 19 lines} \end{array}$ 

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

# 10.2 Misc. algorithms

# TernarySearch.h

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

```
Usage: int ind = ternSearch(0,n-1,[&](int i){return a[i];});

Time: \mathcal{O}(\log(b-a)) 9155b4, 11 lines
```

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

## LIS.h

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

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

```
vi ans(L);
while (L--) ans[L] = cur, cur = prev[cur];
return ans;
}
```

# FastKnapsack.h

**Description:** Given N non-negative integer weights w and a non-negative target t, computes the maximum  $S \le t$  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)) {
      u = v;
      rep(x,0,m) v[x+w[i]] = max(v[x+w[i]], u[x]);
      for (x = 2*m; --x > m;) rep(j, max(0,u[x]), v[x])
      v[x-w[j]] = max(v[x-w[j]], j);
   }
   for (a = t; v[a+m-t] < 0; a--);
   return a;</pre>
```

# 10.3 Dynamic programming

# KnuthDP.h

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

# DivideAndConquerDP.h

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

Time:  $\mathcal{O}\left(\left(N+(hi-lo)\right)\log N\right)$  d38d2b, 18 lines

```
struct DP { // Modify at will:
   int lo(int ind) { return 0; }
   int hi(int ind) { return ind; }
   ll f(int ind, int k) { return dp[ind][k]; }
   void store(int ind, int k, ll v) { res[ind] = pii(k, v); }

void rec(int L, int R, int LO, int HI) {
   if (L >= R) return;
   int mid = (L + R) >> 1;
   pair<ll, int> best(LLONG_MAX, LO);
   rep(k, max(LO,lo(mid)), min(HI,hi(mid)))
      best = min(best, make_pair(f(mid, k), k));
   store(mid, best.second, best.first);
   rec(L, mid, LO, best.second+1);
   rec(mid+1, R, best.second, HI);
   }
   void solve(int L, int R) { rec(L, R, INT_MIN, INT_MAX); }
};
```

# 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.</pre>

# 10.5.2 Pragmas

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

#### FastMod.h

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

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

# FastInput.h

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

Usage: ./a.out < input.txt

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

7b3c70, 17 lines

```
inline char gc() { // like getchar()
  static char buf[1 << 16];
  static size_t bc, be;
  if (bc >= be) {
```

```
buf[0] = 0, bc = 0;
  be = fread(buf, 1, sizeof(buf), stdin);
}
return buf[bc++]; // returns 0 on EOF
}
int readInt() {
  int a, c;
  while ((a = gc()) < 40);
  if (a == '-') return -readInt();
  while ((c = gc()) >= 48) a = a * 10 + c - 480;
  return a - 48;
}
```

#### BumpAllocator.h

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

```
// 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*) {}</pre>
```

## SmallPtr.h

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

"BumpAllocator.h"

2dd6c9, 10 lines

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

# BumpAllocatorSTL.h

Description: BumpAllocator for STL containers.

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

```
char buf[450 << 20] alignas(16);
size_t buf_ind = sizeof buf;

template<class T> struct small {
    typedef T value_type;
    small() {}
    template<class U> small(const U&) {}
    T* allocate(size_t n) {
        buf_ind -= n * sizeof(T);
        buf_ind &= 0 - alignof(T);
        return (T*) (buf + buf_ind);
    }
    void deallocate(T*, size_t) {}
};
```

# SIMD.h

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

```
#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?u:x), movemask_epi8 (hibits of butes)
// 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 < n; ++i) if (a[i] < b[i]) r += a[i] *b[i]; // <- equiv
  return r;
```

# Techniques (A)

# techniques.txt

Combinatorics

159 lines

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

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

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

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