



University of Copenhagen

string & a long

Elias Rasmussen Lolck, Thor Vejen Eriksen, William Bille Meyling

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Setup

bashrc.sh

```
-----4b42a4
setxkbmap -option caps:escape
# fast:
xset r rate 200 120
# normal:
xset r rate 500 35
# debug compile (C++):
dc() {
    bsnm=$(basename "$1" .cpp)
    # EUC uses -std=gnu++20
    command="g++ ${bsnm}.cpp -o $bsnm -Wshadow -Wall -g -
        fsanitize=address,undefined -D_GLIBCXX_DEBUG -std=gnu
        ++20 -Wfatal-errors"
    echo $command
    $command
}
set -o vi
-----
```

hash.sh

```
-----5246ca
# hashes a file, ignoring whitespaces and comments
# use for verifying that code is copied correctly
cpp -dD -P -fpreprocessed | tr -d '[:space:]' | md5sum |
cut -c-6
-----
```

init.lua

```
-----1d7b4f
local options = {
    cmdheight = 1,
    ignorecase = true,
    mouse = "a",
    expandtab = true,
    shiftwidth = 4,
    tabstop = 4,
    cursorline = true,
    number = true,
    relativenumber = true,
    numberwidth = 1,
    signcolumn = "yes",
    wrap = false,
    scrolloff = 6,
    sidescrolloff = 6,
    foldmethod="indent",
    foldlevel=99,
    colorcolumn='80',
}

local keymap = vim.api.nvim_set_keymap
local ops = { noremap = true, silent = true }

keymap("v", "<A-Down>", ":m '>+1<CR>gv=gv", ops)
keymap("v", "<A-Up>", ":m '<-2<CR>gv=gv", ops)
keymap("v", "p", "\\_dP", ops)

for k, v in pairs(options) do
    vim.opt[k] = v
end

local function hashCurrentBuffer()
    local buffer_content = table.concat(vim.api.
        nvim_buf_get_lines(0, 0, -1, false), "\\n")
    local command = "echo '".buffer_content..' ' | cpp -dD
        -P -fpreprocessed | tr -d '[:space:]' | md5sum | cut -c
        -6"
    local hash = vim.fn.system(command)
    hash = hash:gsub("%s+", "")
end
-----
```

```
print("Buffer Hash: " .. hash)
end
vim.api.nvim_create_user_command('Hash', hashCurrentBuffer,
    {})
```

KACTL template

```
-----bd2055
// in addition to template.h, kactl uses:
#define rep(i,a,b) for(int i = a; i < (b); ++i)
#define sz(x) (int)(x).size()
typedef pair<int,int> pii;
typedef vector<int> vi;
-----
```

template

```
-----9aea93
#include <bits/stdc++.h>
using namespace std;
typedef long long ll;
#define all(x) (x).begin(), (x).end()

int main() {
    ios::sync_with_stdio(0); cin.tie(0);
}
-----
```

vimrc

```
-----88d5be
se ch=1 ic mouse=a sw=4 ts=4 nu rnu nuw=4 nowrap so=6 siso
    =8 fdm=indent fdl=99 tm=100
ca Hash w !cpp -dD -P -fpreprocessed \l tr -d '[:space:]'
    \l md5sum \l cut -c-6
vnoremap <silent> <A-Down> :m '>+1<CR>gv=gv
vnoremap <silent> <A-Up> :m '<-2<CR>gv=gv
vnoremap <silent> p "_dP
-----
```

Data_structures

Disjoint Set Union

Description: Classic DSU using path compression and union by rank. unite returns true iff u and v were disjoint.
Usage: Dsu d(n); d.unite(a, b); d.find(a);
Complexity: find(), unite() are amortized $\mathcal{O}(\alpha(n))$, where $\alpha(n)$ is the inverse Ackermann function. Basically $\mathcal{O}(1)$.

```
-----5168ab
struct Dsu {
    vector<int> p, rank;
    Dsu(int n) {
        p.resize(n); rank.resize(n, 0);
        iota(p.begin(), p.end(), 0);
    }
    int find(int x) {
        return p[x] == x ? x : p[x] = find(p[x]);
    }
    bool unite(int u, int v) {
        if ((u = find(u)) == (v = find(v))) return false;
        if (rank[u] < rank[v]) swap(u, v);
        p[v] = u;
        rank[u] += rank[v] == rank[v];
        return true;
    }
};
-----
```

Li-Chao tree

Description: Contianer of lines, online insertion/querying. Retrieve the line f with minimum $f(x)$ for a given x .
Usage: LCT lct(n); lct.insert(line, 0, n); lct.query(x, 0, n);
Complexity: $\mathcal{O}(\log n)$ per insertion/query

```
-----ba9fe6
struct Line { ll a, b; ll f(ll x) { return a * x + b; } };
constexpr const Line LINF { 0, 1LL << 60 };
struct LCT {
    vector<Line> v; // coord-compression: modify v[x] -> v[
        conert(x)]
    LCT(int size) { v.resize(size + 1, LINF); }
    void insert(Line line, int l, int r) {
        if (l > r) return;
        int mid = (l + r) >> 1;
        if (line.f(mid) < v[mid].f(mid)) swap(line, v[mid]);
        if (line.f(l) < v[mid].f(l)) insert(line, l, mid - 1);
        else insert(line, mid + 1, r);
    }
    Line query(int x, int l, int r) {
        if (l > r) return LINF;
        int mid = (l + r) >> 1;
        if (x == mid) return v[mid]; // faster on avg. - not
            necessary
        if (x < mid) return best_of(v[mid], query(x, l, mid -
            1), x);
        return best_of(v[mid], query(x, mid + 1, r), x);
    }
    Line best_of(Line a, Line b, ll x) { return a.f(x) < b.f(
        x) ? a : b; }
};
-----
```

Rollback Union Find

Description: Yoinked from kactl. Disjoint-set data structure with undo. If undo is not needed, skip st, time() and rollback().
Usage: int t = uf.time(); ...; uf.rollback(t);
Complexity: $\mathcal{O}(\log n)$.

```
-----de4ad0
struct RollbackUF {
    vi e; vector<pii> st;
    RollbackUF(int n) : e(n, -1) {}
    int size(int x) { return -e[find(x)]; }
    int find(int x) { return e[x] < 0 ? x : find(e[x]); }
    int time() { return sz(st); }
    void rollback(int t) {
        for (int i = time(); i --> t;)
            e[st[i].first] = st[i].second;
        st.resize(t);
    }
    bool join(int a, int b) {
        a = find(a), b = find(b);
        if (a == b) return false;
        if (e[a] > e[b]) swap(a, b);
        st.push_back({a, e[a]});
        st.push_back({b, e[b]});
        e[a] += e[b]; e[b] = a;
        return true;
    }
};
-----
```

Fenwick tree

Description: Computes prefix sums and single element updates. Uses 0-indexing.
Usage: Fen f(n); f.update(ind, val); f.query(ind);
f.lower_bound(sum);
Complexity: $\mathcal{O}(\log n)$ per update/query

```
-----1743e1
struct Fen {
    vector<ll> v;
};
-----
```

```

Fen(int s) : v(s, 0) { }
void update(int ind, ll val) {
    for (; ind < (int) v.size(); ind |= ind + 1) v[ind] += val;
}
ll query(int ind) { // [0, ind), ind < 0 returns 0
    ll res = 0;
    for (; ind > 0; ind &= ind - 1) res += v[ind - 1]; // operation can be modified
    return res;
}
int lower_bound(ll sum) { // returns first i with query(i + 1) >= sum, n if not found
    int ind = 0;
    for (int p = 1 << 25; p; p >= 1) // 1 << 25 can be lowered to ceil(log2(v.size()))
        if (ind + p <= (int) v.size() && v[ind + p - 1] < sum + sum -= v[(ind += p) - 1];
    return ind;
}
};

```

Fast hash map

Description: 3x faster hash map, 1.5x more memory usage, similar API to `std::unordered_map`. Initial capacity, if provided, must be power of 2.

Usage: `hash_map<key_t, val_t> mp; mp[key] = val; mp.find(key); mp.begin(); mp.end(); mp.erase(key); mp.size();`

Complexity: $\mathcal{O}(1)$ per operation on average.

```

-----bf4708
#include <bits/extc++.h>

struct chash {
    const uint64_t C = ll(4e18 * acos(0)) | 71;
    ll operator () (ll x) const { return __builtin_bswap64(x * C); }
};

template <typename KEY_T, typename VAL_T> using hash_map =
    __gnu_pbds::gp_hash_table<KEY_T, VAL_T, chash>;

```

Implicit 2D segment tree

Description: Classic implicit 2D segment tree taken from my solution to IOI game 2013. It is in rough shape, but it works. Designed to be [inclusive, exclusive). It is old and looks shady, only rely slightly on it, maybe even just make a new one if you need one.

Usage: See usage example at the bottom.

Complexity: $\mathcal{O}(\log^2 n)$ per operation *I think*.

```

-----ae92aa
constexpr const int MX_RC = 1 << 30;

struct Inner {
    long long val;
    int lv, rv;
    Inner* lc,* rc;
    Inner(long long _val, int _l, int _r) :
        val(_val), lv(_l), rv(_r), lc(nullptr), rc(nullptr)
    { }
    ~Inner() {
        delete(lc);
        delete(rc);
    }
    void update(int ind, long long nev, int l = 0, int r = MX_RC) {
        MX_RC) {
            if (!(r - l - 1)) {
                assert(lv == l && rv == r);
                assert(ind == l);
                val = nev;
                return;
            }

```

```

        }
        int mid = (l + r) >> 1;
        if (ind < mid) {
            if (lc) {
                if (lc->lv != l || lc->rv != mid) {
                    Inner* tmp = lc;
                    lc = new Inner(0, l, mid);
                    (tmp->lv < ((l + mid) >> 1) ? lc->lc : lc->rc) = tmp;
                }
                lc->update(ind, nev, l, mid);
            } else lc = new Inner(nev, ind, ind + 1);
        } else {
            if (rc) {
                if (rc->lv != mid || rc->rv != r) {
                    Inner* tmp = rc;
                    rc = new Inner(0, mid, r);
                    (tmp->lv < ((mid + r) >> 1) ? rc->lc : rc->rc) = tmp;
                }
                rc->update(ind, nev, mid, r);
            } else rc = new Inner(nev, ind, ind + 1);
        }
        val = std::gcd(lc ? lc->val : 0, rc ? rc->val : 0);
    }
    long long query(int tl, int tr, int l = 0, int r = MX_RC)
    {
        if (l >= tr || r <= tl) return 0;
        if (!(rv - lv - 1)) {
            if (lv >= tr || rv <= tl) return 0;
            return val;
        }
        assert(l == lv && r == rv);
        if (l >= tl && r <= tr) return val;
        int mid = (l + r) >> 1;
        return std::gcd(lc ? lc->query(tl, tr, l, mid) : 0, rc ? rc->query(tl, tr, mid, r) : 0);
    }
    void fill(Inner* source) {
        val = source->val;
        if (!(lv - rv - 1)) return;
        if (source->lc) {
            lc = new Inner(source->lc->val, source->lc->lv, source->lc->rv);
            source->lc->rv);
            lc->fill(source->lc);
        }
        if (source->rc) {
            rc = new Inner(source->rc->val, source->rc->lv, source->rc->rv);
            source->rc->lv);
            rc->fill(source->rc);
        }
    }
};

struct Outer {
    Inner* inner;
    int lv, rv;
    Outer* lc,* rc;
    Outer(Inner* _inner, int _l, int _r) :
        inner(_inner), lv(_l), rv(_r), lc(nullptr), rc(nullptr)
    { }
    void update(int ind_outer, int ind_inner, long long nev, int l = 0, int r = MX_RC) {
        if (!(r - l - 1)) {
            assert(lv == l && rv == r);
            assert(ind_outer == l);
            assert(inner);
            inner->update(ind_inner, nev);
            return;
        }
        int mid = (l + r) >> 1;
        if (ind_outer < mid) {

```

```

            if (lc) {
                if (lc->lv != l || lc->rv != mid) {
                    Outer* tmp = lc;
                    lc = new Outer(new Inner(0, 0, MX_RC), l, mid);
                    lc->inner->fill(tmp->inner);
                    (tmp->lv < ((l + mid) >> 1) ? lc->lc : lc->rc) = tmp;
                }
                lc->update(ind_outer, ind_inner, nev, l, mid);
            } else {
                lc = new Outer(new Inner(0, 0, MX_RC), ind_outer, ind_outer + 1);
                lc->inner->update(ind_inner, nev);
            }
        } else {
            if (rc) {
                if (rc->lv != mid || rc->rv != r) {
                    Outer* tmp = rc;
                    rc = new Outer(new Inner(0, 0, MX_RC), mid, r);
                    rc->inner->fill(tmp->inner);
                    (tmp->lv < ((mid + r) >> 1) ? rc->lc : rc->rc) = tmp;
                }
                rc->update(ind_outer, ind_inner, nev, mid, r);
            } else {
                rc = new Outer(new Inner(nev, 0, MX_RC), ind_outer, ind_outer + 1);
                rc->inner->update(ind_inner, nev);
            }
        }
        inner->update(ind_inner, std::gcd(
            lc ? lc->inner->query(ind_inner, ind_inner + 1) : 0,
            rc ? rc->inner->query(ind_inner, ind_inner + 1) : 0));
    }
    long long query(int tl_outer, int tr_outer, int tl_inner, int tr_inner, int l = 0, int r = MX_RC) {
        if (l >= tr_outer || r <= tl_outer) return 0;
        if (!(rv - lv - 1)) {
            if (lv >= tr_outer || rv <= tl_outer) return 0;
            return inner->query(tl_inner, tr_inner);
        }
        assert(l == lv && r == rv);
        if (l >= tl_outer && r <= tr_outer)
            return inner->query(tl_inner, tr_inner);
        int mid = (l + r) >> 1;
        return std::gcd(
            lc ? lc->query(tl_outer, tr_outer, tl_inner, tr_inner, l, mid) : 0,
            rc ? rc->query(tl_outer, tr_outer, tl_inner, tr_inner, mid, r) : 0);
    }
};

// this is how it has been used in the solution to IOI game 2013
Outer root(new Inner(0, 0, MX_RC), 0, MX_RC);
void update(int r, int c, long long k) {
    root.update(r, c, k);
}
long long calculate(int r_l, int c_l, int r_r, int c_r) {
    return root.query(r_l, r_r + 1, c_l, c_r + 1);
}

```

Lazy segment tree

Description: Zero-indexed, bounds are [l, r), operations can be modified. $\mathcal{O}(\log n)$ find_first and the like can be implemented by checking bounds, then checking left tree, then right tree, recursively.

Usage: `Lazy_segtree seg(n); seg.update(l, r, val); seg.query(l, r);`

Complexity: $\mathcal{O}(\log n)$ per update/query

```
-----
struct Lazy_segtree {
    typedef ll T; // change type here
    typedef ll LAZY_T; // change type here
    static constexpr T unit = 0; // change unit here
    static constexpr LAZY_T lazy_unit = 0; // change lazy
        unit here
    T f(T l, T r) { return l + r; } // change operation here
    void push(int now, int l, int r) {
        if (w[now] == lazy_unit) return;
        v[now] += w[now] * (r - l); // operation can be
            modified
        if (r - l - 1)
            w[now * 2 + 1].first += w[now],
            w[now * 2 + 2].first += w[now];
        w[now] = lazy_unit;
    }
    int size;
    vector<T> v;
    vector<LAZY_T> w;
    Lazy_segtree(int s = 0) : size(s ? 1 << (32 -
        __builtin_clz(s)) : 0), v(size << 1, unit), w(size <<
        1, lazy_unit) { }
    template<typename U> void update(int l, int r, U val) {
        update(l, r, val, 0, 0, size); }
    T query(int l, int r) { return query(l, r, 0, 0, size); }
    template<typename U> void update(int tl, int tr, U val,
        int now, int l, int r) {
        push(now, l, r);
        if (l >= tr || r <= tl) return;
        if (l >= tl && r <= tr) {
            // this does not *have* to accumulate, push is called
            before this:
            w[now] += val; // operation can be modified
            push(now, l, r);
            return;
        }
        int mid = (l + r) >> 1;
        update(tl, tr, val, now * 2 + 1, l, mid);
        update(tl, tr, val, now * 2 + 2, mid, r);
        v[now] = f(v[now * 2 + 1], v[now * 2 + 2]);
    }
    T query(int tl, int tr, int now, int l, int r) {
        push(now, l, r);
        if (l >= tr || r <= tl) return unit;
        if (l >= tl && r <= tr) return v[now];
        int mid = (l + r) >> 1;
        return f(query(tl, tr, now * 2 + 1, l, mid), query(tl,
            tr, now * 2 + 2, mid, r));
    }
    template<typename U> void build(const vector<U>& a) {
        for (int i = 0; i < (int) a.size(); i++) v[size - 1 + i
            ] = a[i]; // operation can be modified
        for (int i = size - 2; i >= 0; i--) v[i] = f(v[i * 2 +
            1], v[i * 2 + 2]);
    }
};
-----
```

Matrix

Description: Yoinked from kactl. Basic operations on square matrices.
Usage: Matrix<int, 3> A; A.d = {{{{1,2,3}}}, {{4,5,6}}}, {{{7,8,9}}}}; vector<int> vec = {1,2,3}; vec = (A^N) * vec;
Complexity: $\mathcal{O}(n^3)$ per multiplication, $\mathcal{O}(n^3 \log p)$ per exponentiation.

```
c43c7d
-----
template<class T, int N> struct Matrix {
    typedef Matrix M;
    array<array<T, N>, N> d{};
    M operator*(const M& m) const {
        M a;
```

```

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

Ordered Map

Description: extc++.h order statistics tree. find_by_order returns an iterator to the k th element (0-indexed), order_of_key returns the index of the element (0-indexed), i.e. the number of elements less than the argument.
Usage: ordered_set<int> s; s.insert(1); s.insert(2); *s.find_by_order(0) = 3; s.erase(3); s.order_of_key(2);
Complexity: Everything is $\mathcal{O}(\log n)$.

```
-----225ff4
#include <bits/extc++.h>
// if judge does not have extc++.h, use:
// #include <ext/pb_ds/assoc_container.hpp>
// #include <ext/pb_ds/tree_policy.hpp>
using namespace __gnu_pbds;
template<typename T> using ordered_set = tree<T,
    null_type, less<T>, rb_tree_tag,
    tree_order_statistics_node_update>;
template<typename T, typename U> using ordered_map = tree
    <T, U, less<T>, rb_tree_tag,
    tree_order_statistics_node_update>;

// yeet from kactl:
void example() {
    ordered_set<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
}
-----
```

Persistent segment tree

Description: Zero-indexed, bounds are [l, r), operations can be modified. update(...) returns a pointer to a new tree with the applied update, all other trees remain unchanged. $\mathcal{O}(\log n)$ find.first and the like can be implemented by checking bounds, then checking left tree, then right tree, recursively.
Usage: Node* root = build(arr, 0, n); Node* another_root = update(root, ind, val, 0, 0, n); query(some_root, l, r, 0, n).val; Node* empty_root = nullptr; Node* another_version = update(empty_root, ind, val, 0, 0, n);
Complexity: $\mathcal{O}(\log n)$ per update/query, $\mathcal{O}(n)$ per build

```
-----b73a59
struct Node {
    Node* l,* r;
    int val; // i.e. data
    Node(int _v) : l(nullptr), r(nullptr), val(_v) { }
```

```

    Node(Node* _l, Node* _r) : l(_l), r(_r), val(0) {
        // i.e. merge two nodes:
        if (l) val += l->val;
        if (r) val += r->val;
    }
};

// slightly more memory, much faster:
template<typename... ARGS> Node* new_node(ARGS&&... args)
{
    static std::deque<Node> pool;
    pool.emplace_back(std::forward<ARGS>(args)...);
    return &pool.back();
}

// slightly less memory, much slower:
// #define new_node(...) new Node(__VA_ARGS__)

// optional:
Node* build(const std::vector<int>& a, int l, int r) {
    if (!(r - l - 1)) return new_node(a[l]);
    int mid = (l + r) >> 1;
    return new_node(build(a, l, mid), build(a, mid, r));
}

// can be called with node == nullptr
Node* update(Node* node, int ind, int val, int now, int l,
    int r) {
    if (!(r - l - 1)) return new_node(val); // i.e. point
        update
    int mid = (l + r) >> 1;
    Node* lf = node ? node->l : nullptr;
    Node* rg = node ? node->r : nullptr;
    return new_node
        (ind < mid ? update(lf, ind, val, now, l, mid) : lf,
        ind >= mid ? update(rg, ind, val, now, mid, r) : rg);
}

Node query(Node* node, int tl, int tr, int l, int r) {
    if (l >= tr || r <= tl || !node) return Node(0); // i.e.
        empty node
    if (l >= tl && r <= tr) return *node;
    int mid = (l + r) >> 1;
    Node lf = query(node->l, tl, tr, l, mid);
    Node rg = query(node->r, tl, tr, mid, r);
    return Node(&lf, &rg);
}
-----
```

Segment tree

Description: Zero-indexed, bounds are [l, r), operations can be modified. $\mathcal{O}(\log n)$ find.first and the like can be implemented by checking bounds, then checking left tree, then right tree, recursively.
Usage: Segtree seg(n); seg.update(ind, val); seg.query(l, r);
Complexity: $\mathcal{O}(\log n)$ per update/query.

```
-----f40463
struct Segtree {
    typedef ll T; // change type here
    static constexpr T unit = 0; // change unit here
    T f(T l, T r) { return l + r; } // change operation here
    int size;
    vector<T> v;
    Segtree(int s = 0) : size(s ? 1 << (32 - __builtin_clz(s)
        ) : 0), v(size << 1, unit) { }
    void update(int ind, T val) { update(ind, val, 0, 0, size
        ); }
    T query(int l, int r) { return query(l, r, 0, 0, size); }
    void update(int ind, T val, int now, int l, int r) {
        if (!(r - l - 1)) { v[now] = val; return; } //
            operation can be modified
        int mid = (l + r) >> 1;
        if (ind < mid) update(ind, val, now * 2 + 1, l, mid);
        else update(ind, val, now * 2 + 2, mid, r);
    }
};
-----
```

```

    v[now] = f(v[now * 2 + 1], v[now * 2 + 2]);
}
T query(int tl, int tr, int now, int l, int r) {
    if (l >= tr || r <= tl) return unit;
    if (l >= tl && r <= tr) return v[now];
    int mid = (l + r) >> 1;
    return f(query(tl, tr, now * 2 + 1, l, mid), query(tl,
tr, now * 2 + 2, mid, r));
}
template <typename U> void build(const vector <U& a) {
    for (int i = 0; i < (int) a.size(); i++) v[size - 1 + i
] = a[i]; // operation can be modified
    for (int i = size - 2; i >= 0; i--) v[i] = f(v[i * 2 +
1], v[i * 2 + 2]);
}
};
```

Sparse table

Description: Yoinked from kactl. Classic sparse table, implemented with range minimum queries, can be modified.
Usage: Sparse s(vec); s.query(a, b);
Complexity: $O(|V| \log |V| + Q)$.

```

-----5b1135
template<class T> struct Sparse {
    vector<vector<T>> jmp;
    Sparse(const vector<T>& V) : jmp(1, V) {
        for (int pw = 1, k = 1; pw * 2 <= sz(V); pw *= 2, ++k)
            {
                jmp.emplace_back(sz(V) - pw * 2 + 1);
                rep(j,0,sz(jmp[k]))
                    jmp[k][j] = min(jmp[k - 1][j], jmp[k - 1][j + pw]);
            }
    }
    T query(int a, int b) { // interval [a, b)
        assert(a < b); // or return inf if a == b
        int dep = 31 - __builtin_clz(b - a);
        return min(jmp[dep][a], jmp[dep][b - (1 << dep)]);
    }
};
```

Treap

Description: Yoinked from kactl. A short self-balancing tree. It acts as a sequential container with log-time splits/joins, and is easy to augment with additional data.
Complexity: $O(\log n)$ operations.

```

-----9556fc
struct Node {
    Node *l = 0, *r = 0;
    int val, y, c = 1;
    Node(int val) : val(val), y(rand()) {}
    void recalc();
};

int cnt(Node* n) { return n ? n->c : 0; }
void Node::recalc() { c = cnt(l) + cnt(r) + 1; }

template<class F> void each(Node* n, F f) {
    if (!n) return;
    if (cnt(n->l) >= cnt(n->r)) { each(n->l, f); each(n->r, f); }
    else { each(n->r, f); each(n->l, f); }
}

pair<Node*, Node*> split(Node* n, int k) {
    if (!n) return {0, 0};
    if (cnt(n->l) >= k) { // "n->val >= k" for lower_bound(k)
        auto pa = split(n->l, k);
        n->l = pa.second;
        n->recalc();
        return {pa.first, n};
    } else {
        auto pa = split(n->r, k - cnt(n->l) - 1); // and just "k"
        n->r = pa.first;
        n->recalc();
        return {n, pa.second};
    }
}
```

```

    n->r = pa.first;
    n->recalc();
    return {n, pa.second};
}

Node* merge(Node* l, Node* r) {
    if (!l) return r;
    if (!r) return l;
    if (l->y > r->y) {
        l->r = merge(l->r, r);
        l->recalc();
        return l;
    } else {
        r->l = merge(l, r->l);
        r->recalc();
        return r;
    }
}

Node* ins(Node* t, Node* n, int pos) {
    auto pa = split(t, pos);
    return merge(merge(pa.first, n), pa.second);
}

// Example application: move the range [l, r) to index k
void move(Node& t, int l, int r, int k) {
    Node *a, *b, *c;
    tie(a,b) = split(t, l); tie(b,c) = split(b, r - l);
    if (k <= l) t = merge(merge(a, b), c);
    else t = merge(a, merge(b, c));
}
```

Geometry

3D convex hull

Description: Yoinked from kactl. Computes all faces of the 3-dimension hull of a point set. *No four points must be coplanar*, or else random results will be returned. All faces will point outwards.
Complexity: $O(n^2)$.

```

-----6e20a4
#include "Point_3D.h"

typedef Point3D<double> P3;

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

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

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

```

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

Angle

Description: Yoinked from kactl. 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

```

-----0f0602
struct Angle {
    int x, y;
    int t;
    Angle(int x, int y, int t=0) : x(x), y(y), t(t) {}
    Angle operator-(Angle b) const { return {x-b.x, y-b.y, t
        - b.t}; }
    int half() const {
        assert(x || y);
        return y < 0 || (y == 0 && x < 0);
    }
    Angle t90() const { return {-y, x, t + (half() && x >= 0)
        }; }
    Angle t180() const { return {-x, -y, t + half()}; }
    Angle t360() const { return {x, y, t + 1}; }
};

bool operator<(Angle a, Angle b) {
    // add a.dist2() and b.dist2() to also compare distances
    return make_tuple(a.t, a.half(), a.y * (1ll)b.x) <
        make_tuple(b.t, b.half(), a.x * (1ll)b.y);
}

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

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



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

Circle circle intersection

Description: Yoinked from kactl. Computes the pair of points at which two circles intersect. Returns false in case of no intersection.
Complexity: $\mathcal{O}(1)$.

```
-----583f44
#include "Point.h"

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

Circle line intersection

Description: Yoinked from kactl. 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>.

```
-----a84598
#include "Point.h"

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

Circle polygon intersection

Description: Yoinked from kactl. Returns the area of the intersection of a circle with a ccw polygon.
Complexity: $\mathcal{O}(n)$.

```
-----8d446d
#include "Point.h"

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

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

Circle tangents

Description: Yoinked from kactl. 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.

```
-----15aa0d
#include "Point.h"

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

Circumcircle

Description: Yoinked from kactl. 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.

```
-----fca990
#include "Point.h"

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

Closest pair of points

Description: Yoinked from kactl. Finds the closest pair of points.
Complexity: $\mathcal{O}(n \log n)$.

```
-----ac9ec9
#include "Point.h"

typedef Point<ll> P;
pair<P, P> closest(vector<P> v) {
    assert(sz(v) > 1);
    set<P> S;
    sort(all(v), [](P a, P b) { return a.y < b.y; });
    pair<ll, pair<P, P>> ret{LLONG_MAX, {P(), P()}};
    int j = 0;
    for (P p : v) {
        P d{1 + (ll)sqrt(ret.first), 0};
        while (v[j].y <= p.y - d.x) S.erase(v[j++]);
        auto lo = S.lower_bound(p - d), hi = S.upper_bound(p +
            d);
        for (; lo != hi; ++lo)
```

```
        ret = min(ret, {(ll) - p).dist2(), {(ll) - p).dist2()});
        S.insert(p);
    }
    return ret.second;
}
```

Convex hull

Description: Yoinked from kactl. Returns a vector of the points of the convex hull in counter-clockwise order. Points on the edge of the hull between two other points are not considered part of the hull.
Complexity: $\mathcal{O}(n \log n)$.

```
-----ec8c60
#include "Point.h"

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

Delaunay triangulation

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

Complexity: $\mathcal{O}(n^2)$.

```
-----0da7c8
#include "Point.h"
#include "3d_hull.h"

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

Dynamic Convex Hull

Description: Supports building a convex hull one point at a time. Viewing the convex hull along the way.

```
-----431bba
struct point {
    ll x, y;
    point(ll x=0, ll y=0): x(x), y(y) {}
    point operator-(const point &p) const { return point(x-
        p.x, y-p.y); }
    point operator*(const ll k) const { return point(k*x, k
        *y); }
    ll cross(const point &p) const { return x*p.y - p.x*y;
    }
    bool operator<(const point &p) const { return x < p.x
        || x == p.x && y < p.y; }
};
```

```

bool above(set<point> &hull, point p, ll scale = 1) {
    auto it = hull.lower_bound(point((p.x+scale-1)/scale, 0));
    if (it == hull.end()) return true;
    if (p.y <= it->y*scale) return false;
    if (it == hull.begin()) return true;
    auto jt = it--;
    return (p-*it*scale).cross(*jt-*it) < 0;
}

void add(set<point> &hull, point p) {
    if (!above(hull, p)) return;
    auto pit = hull.insert(p).first;
    while (pit != hull.begin()) {
        auto it = prev(pit);
        if (it->y <= p.y || (it != hull.begin() && (*it-*prev(it)).cross(*pit-*it) >= 0))
            hull.erase(it);
        else
            break;
    }
    auto it = next(pit);
    while (it != hull.end()) {
        if (next(it) != hull.end() && (*it-p).cross(*next(it)-*it) >= 0)
            hull.erase(it++);
        else
            break;
    }
}

```

Hull diameter

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

Complexity: $\mathcal{O}(n)$.

```

-----326872
#include "Point.h"

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

```

Inside polygon

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

Usage: vector<P> v = {P{4,4}, P{1,2}, P{2,1}};

bool in = inPolygon(v, P{3, 3}, false);

Complexity: $\mathcal{O}(n)$.

```

-----bbf6f5
#include "Point.h"
#include "On_segment.h"
#include "Segment_distance.h"

template<class P>
bool inPolygon(vector<P> &p, P a, bool strict = true) {
    int cnt = 0, n = sz(p);

```

```

rep(i,0,n) {
    P q = p[(i + 1) % n];
    if (onSegment(p[i], q, a)) return !strict;
    //or: if (segDist(p[i], q, a) <= eps) return !strict;
    cnt ^= ((a.y<p[i].y) - (a.y<q.y)) * a.cross(p[i], q) > 0;
}
return cnt;
}

```

KD-tree

Description: Yoinked from kactl. 2D, can be extended to 3D. See comments for details.

```

-----08fbca
#pragma once

#include "Point.h"

typedef long long T;
typedef Point<T> P;
const T INF = numeric_limits<T>::max();

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

struct Node {
    P pt; // if this is a leaf, the single point in it
    T x0 = INF, x1 = -INF, y0 = INF, y1 = -INF; // bounds
    Node *first = 0, *second = 0;

    T distance(const P& p) { // min squared distance to a point
        T x = (p.x < x0 ? x0 : p.x > x1 ? x1 : p.x);
        T y = (p.y < y0 ? y0 : p.y > y1 ? y1 : p.y);
        return (P(x,y) - p).dist2();
    }

    Node(vector<P>&& vp) : pt(vp[0]) {
        for (P p : vp) {
            x0 = min(x0, p.x); x1 = max(x1, p.x);
            y0 = min(y0, p.y); y1 = max(y1, p.y);
        }
        if (vp.size() > 1) {
            // split on x if width >= height (not ideal...)
            sort(all(vp), x1 - x0 >= y1 - y0 ? on_x : on_y);
            // divide by taking half the array for each child (
            not
            // best performance with many duplicates in the middle)
            int half = sz(vp)/2;
            first = new Node({vp.begin(), vp.begin() + half});
            second = new Node({vp.begin() + half, vp.end()});
        }
    }
};

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

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

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

        // search closest side first, other side if needed

```

```

    auto best = search(f, p);
    if (bsec < best.first)
        best = min(best, search(s, p));
    return best;
}

```

```

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

```

Line hull intersection

Description: Yoinked from kactl. 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 poly:

- $(-1, -1)$ if no collision,
- $(i, -1)$ if touching the corner i ,
- (i, i) if along side $(i, i + 1)$,
- (i, j) if crossing sides $(i, i + 1)$ and $(j, j + 1)$.

In the last case, if a corner i is crossed, this is treated as happening on side $(i, i + 1)$. The points are returned in the same order as the line hits the polygon.

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

```

-----32bb13
#include "Point.h"

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

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

```

```
        case 2: return {res[1], res[1]};
    }
    return res;
}
```

Line line intersection

Description: Yoinked from kactl. 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{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 ll.

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

```
-----ae219b
#include "Point.h"

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

Line projection and reflection

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

```
-----c08b13
#include "Point.h"

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

Linear transformation

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

```
-----a1c4fb
#include "Point.h"

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

Manhattan MST

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

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

```
-----c264e8
#include "Point.h"
```

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

Minimum enclosing circle

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

Complexity: $\mathcal{O}(n)$.

```
-----0b9ff5
#include "circumcircle.h"

pair<P, double> mec(vector<P> ps) {
    shuffle(all(ps), mt19937(time(0)));
    P o = ps[0];
    double r = 0, EPS = 1 + 1e-8;
    rep(i,0,sz(ps)) if ((o - ps[i]).dist() > r * EPS) {
        o = ps[i], r = 0;
        rep(j,0,i) if ((o - ps[j]).dist() > r * EPS) {
            o = (ps[i] + ps[j]) / 2;
            r = (o - ps[i]).dist();
            rep(k,0,j) if ((o - ps[k]).dist() > r * EPS) {
                o = ccCenter(ps[i], ps[j], ps[k]);
                r = (o - ps[i]).dist();
            }
        }
    }
    return {o, r};
}
```

Is on segment

Description: Yoinked from kactl. Returns true iff p lies on the line segment from s to e. Use (segDist(s,e,p)<=epsilon) instead when using Point <double>.

```
-----5fde08
#include "Point.h"

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

2D Point

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

```
-----47ec0a
template <class T> int sgn(T x) { return (x > 0) - (x < 0); }
template<class T>
struct Point {
```

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

3D Point

Description: Yoinked from kactl. Class to handle points in 3D space. T can be e.g. double or long long. (Avoid int.).

```
-----8058ae
template<class T> struct Point3D {
    typedef Point3D P;
    typedef const P& R;
    T x, y, z;
    explicit Point3D(T x=0, T y=0, T z=0) : x(x), y(y), z(z) {}
    bool operator<(R p) const {
        return tie(x, y, z) < tie(p.x, p.y, p.z); }
    bool operator==(R p) const {
        return tie(x, y, z) == tie(p.x, p.y, p.z); }
    P operator+(R p) const { return P(x+p.x, y+p.y, z+p.z); }
    P operator-(R p) const { return P(x-p.x, y-p.y, z-p.z); }
    P operator*(T d) const { return P(x*d, y*d, z*d); }
    P operator/(T d) const { return P(x/d, y/d, z/d); }
    T dot(R p) const { return x*p.x + y*p.y + z*p.z; }
    P cross(R p) const {
        return P(y*p.z - z*p.y, z*p.x - x*p.z, x*p.y - y*p.x); }
    T dist2() const { return x*x + y*y + z*z; }
    double dist() const { return sqrt((double)dist2()); }
    //Azimuthal angle (longitude) to x-axis in interval [-pi, pi]
    double phi() const { return atan2(y, x); }
    //Zenith angle (latitude) to the z-axis in interval [0, pi]
    double theta() const { return atan2(sqrt(x*x+y*y),z); }
    P unit() const { return *this/(T)dist(); } //makes dist()==1
    //returns unit vector normal to *this and p
    P normal(P p) const { return cross(p).unit(); }
    //returns point rotated 'angle' radians ccw around axis
    P rotate(double angle, P axis) const {
        double s = sin(angle), c = cos(angle); P u = axis.unit();
        return u*dot(u)*(1-c) + (*this)*c - cross(u)*s;
```



```
    }
};
```

Is point in convex polygon

Description: Yoinked from kactl. 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.

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

```
-----78b7ca
#include "Point.h"
#include "Side_of.h"
#include "On_segment.h"

typedef Point<ll> P;

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

Polygon area

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

```
-----714de9
#include "Point.h"

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

Polygon center of mass

Description: Yoinked from kactl. Returns the center of mass for a polygon.

Complexity: $\mathcal{O}(n)$.

```
-----6906c1
#include "Point.h"

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

Polygon cut

Description: Yoinked from kactl. 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));

```
#include "Point.h"
#include "Line_intersection.h"

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

Polygon union

Description: Yoinked from kactl. 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.)

Complexity: $\mathcal{O}(n^2)$ where n is the total number of points.

```
-----82e22a
#include "Point.h"
#include "Side_of.h"

typedef Point<double> P;
double rat(P a, P b) { return sgn(b.x) ? a.x/b.x : a.y/b.y;
}

double polyUnion(vector<vector<P>>& poly) {
    double ret = 0;
    rep(i,0,sz(poly)) rep(v,0,sz(poly[i])) {
        P A = poly[i][v], B = poly[i][(v + 1) % sz(poly[i])];
        vector<pair<double, int>> segs = {{0, 0}, {1, 0}};
        rep(j,0,sz(poly)) if (i != j) {
            rep(u,0,sz(poly[j])) {
                P C = poly[j][u], D = poly[j][(u + 1) % sz(poly[j])
            ];
            int sc = sideOf(A, B, C), sd = sideOf(A, B, D);
            if (sc != sd) {
                double sa = C.cross(D, A), sb = C.cross(D, B);
                if (min(sc, sd) < 0)
                    segs.emplace_back(sa / (sa - sb), sgn(sc - sd));
            } else if (!sc && !sd && j < i && sgn((B-A).dot(D-C))
>0){
                segs.emplace_back(rat(C - A, B - A), 1);
                segs.emplace_back(rat(D - A, B - A), -1);
            }
        }
    }
    sort(all(segs));
    for (auto& s : segs) s.first = min(max(s.first, 0.0),
1.0);
    double sum = 0;
    int cnt = segs[0].second;
    rep(j,1,sz(segs)) {
        if (!cnt) sum += segs[j].first - segs[j - 1].first;
        cnt += segs[j].second;
    }
    ret += A.cross(B) * sum;
}

return ret / 2;
}
```

Polyhedron volume

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

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

Points line-segments distance

Description: Yoinked from kactl. Returns the shortest distance between point p and the line segment from point s to e.

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

bool onSegment = segDist(a,b,p) < 1e-10;

```
-----a3104f
#include "Point.h"

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

Line segment line segment intersection

Description: Yoinked from kactl. 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||_ and the intersection point does not have integer coordinates. Products of three coordinates are used in intermediate steps so watch out for overflow if using int or long long.

Usage: vector <P> inter = segInter(s1,e1,s2,e2); if (sz(inter)==1) cout << "segments intersect at " << inter[0] << endl;

```
-----12fa1d
#include "Point.h"
#include "OnSegment.h"

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

Side of

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

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

```
-----f459d8
#include "Point.h"

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

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

Spherical distance

Description: Yoinked from kactl. Returns the shortest distance on the sphere with radius *radius* between the points with azimuthal angles (longitude) f_1 (ϕ_1) and f_2 (ϕ_2) from x axis and zenith angles (latitude) t_1 (θ_1) and t_2 (θ_2) from z axis (0 = north pole). All angles measured in radians. The algorithm starts by converting the spherical coordinates to cartesian coordinates so if that is what you have you can use only the two last rows. $dx \cdot radius$ is then the difference between the two points in the x direction and $d \cdot radius$ is the total distance between the points.

```
-----611f07-----
double sphericalDistance(double f1, double t1,
    double f2, double t2, double radius) {
    double dx = sin(t2)*cos(f2) - sin(t1)*cos(f1);
    double dy = sin(t2)*sin(f2) - sin(t1)*sin(f1);
    double dz = cos(t2) - cos(t1);
    double d = sqrt(dx*dx + dy*dy + dz*dz);
    return radius*2*asin(d/2);
}
```

Line distance

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

```
-----d1d032-----
#include "Point.h"

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

Graphs

Articulation points finding

Description: Yoinked from CP-algorithms. Standard articulation points finding algorithm.
Complexity: $\mathcal{O}(V + E)$.

```
-----23f413-----
int n; // number of nodes
vector<vector<int>>> adj; // adjacency list of graph

vector<bool> visited;
vector<int> tin, low;
int timer;

void dfs(int v, int p = -1) {
    visited[v] = true;
    tin[v] = low[v] = timer++;
    int children=0;
    for (int to : adj[v]) {
        if (to == p) continue;
        if (visited[to]) {
```

```
        low[v] = min(low[v], tin[to]);
    } else {
        dfs(to, v);
        low[v] = min(low[v], low[to]);
        if (low[to] >= tin[v] && p!=-1)
            IS_CUTPOINT(v);
        ++children;
    }
}
if(p == -1 && children > 1)
    IS_CUTPOINT(v);
}

void find_cutpoints() {
    timer = 0;
    visited.assign(n, false);
    tin.assign(n, -1);
    low.assign(n, -1);
    for (int i = 0; i < n; ++i) {
        if (!visited[i])
            dfs(i);
    }
}
```

Bellman-Ford

Description: Yoinked from kactl. 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 $V^2 \max |w_i| < 2^{63}$.
Usage: `bellmanFord(nodes, edges, s)`.
Complexity: $\mathcal{O}(VE)$.

```
-----830a8f-----
const ll inf = LLONG_MAX;
struct Ed { int a, b, w, s() { return a < b ? a : -a; }};
struct Node { ll dist = inf; int prev = -1; };

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

Biconnected components

Description: Yoinked from kactl. Finds all biconnected components in an undirected graph, and runs a callback for the edges in each. In a biconnected component there are at least two distinct paths between any two nodes. Note that a node can be in several components. An edge which is not in a component is a bridge, i.e., not part of any cycle.
Usage: `int eid = 0; ed.resize(n); for each edge (a, b) { ed[a].emplace(b, eid); ed[b].emplace(a, eid++); } bicomps([&] (const vi& edgelist) { ... });`
Complexity: $\mathcal{O}(E + V)$.

```
-----2965e5-----
```

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

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

Binary lifting with LCA

Description: Yoinked from kactl. Finds power of two jumps in a tree - and standard LCA. Assumes the root node points to itself!
Usage: `vector<vi> jmps = treeJump(parents); int l = lca(jmps, depth, a, b);`
Complexity: $\mathcal{O}(N \log N)$ construction. $\mathcal{O}(\log N)$ per query.

```
-----bfc085-----
vector<vi> treeJump(vi& P){
    int on = 1, d = 1;
    while(on < sz(P)) on *= 2, d++;
    vector<vi> jmp(d, P);
    rep(i,1,d) rep(j,0,sz(P))
        jmp[i][j] = jmp[i-1][jmp[i-1][j]];
    return jmp;
}

int jmp(vector<vi>& tbl, int nod, int steps){
    rep(i,0,sz(tbl))
        if(steps&(1<<i)) nod = tbl[i][nod];
    return nod;
}

int lca(vector<vi>& tbl, vi& depth, int a, int b) {
    if (depth[a] < depth[b]) swap(a, b);
    a = jmp(tbl, a, depth[a] - depth[b]);
    if (a == b) return a;
    for (int i = sz(tbl); i--;) {
        int c = tbl[i][a], d = tbl[i][b];
        if (c != d) a = c, b = d;
    }
    return tbl[0][a];
}
```

Bridge finding

Description: Yoinked from CP-algorithms. Standard bridge finding algorithm.
Complexity: $\mathcal{O}(V + E)$.

```
-----
int n; // number of nodes
vector<vector<int>>> adj; // adjacency list of graph
```

```
vector<bool> visited;
vector<int> tin, low;
int timer;
```

```
void dfs(int v, int p = -1) {
    visited[v] = true;
    tin[v] = low[v] = timer++;
    for (int to : adj[v]) {
        if (to == p) continue;
        if (visited[to]) {
            low[v] = min(low[v], tin[to]);
        } else {
            dfs(to, v);
            low[v] = min(low[v], low[to]);
            if (low[to] > tin[v])
                IS_BRIDGE(v, to);
        }
    }
}
```

```
void find_bridges() {
    timer = 0;
    visited.assign(n, false);
    tin.assign(n, -1);
    low.assign(n, -1);
    for (int i = 0; i < n; ++i) {
        if (!visited[i])
            dfs(i);
    }
}
```

DFS Bipartite Matching

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

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

Complexity: $\mathcal{O}(VE)$.

```
-----
bool find(int j, vector<vi>& g, vi& btoa, vi& vis) {
    if (btoa[j] == -1) return 1;
    vis[j] = 1; int di = btoa[j];
    for (int e : g[di])
        if (!vis[e] && find(e, g, btoa, vis)) {
            btoa[e] = di;
            return 1;
        }
    return 0;
}

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

Dinic's Algorithm

Description: Yoinked from kactl. Finds the maximum flow from s to t in a directed graph. To obtain the actual flow values, look at all edges

with capacity > 0 (zero capacity edges are residual edges).

Usage: Dinic dinic(n); dinic.addEdge(a, b, c); dinic.maxFlow(s, t);

Complexity: $\mathcal{O}(VE \log U)$ where $U = \max | \text{capacity} |$. $\mathcal{O}(\min(\sqrt{E}, V^{2/3})E)$ if $U = 1$; so $\mathcal{O}(\sqrt{VE})$ for bipartite matching.

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

MST in directed graphs

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

Usage: pair<ll, vi> res = DMST(n, edges, root);

Complexity: $\mathcal{O}(E \log V)$.

```
-----
#include "../Data_structures/dsu_rollback.h"
```

```
struct Edge { int a, b; ll w; };
struct Node { /// lazy skew heap node
    Edge key;
    Node *l, *r;
    ll delta;
    void prop() {
        key.w += delta;
        if (l) l->delta += delta;
        if (r) r->delta += delta;
        delta = 0;
    }
};
```

```
    }
    Edge top() { prop(); return key; }
};

Node *merge(Node *a, Node *b) {
    if (!a || !b) return a ? b;
    a->prop(), b->prop();
    if (a->key.w > b->key.w) swap(a, b);
    swap(a->l, (a->r = merge(b, a->r)));
    return a;
}

void pop(Node*& a) { a->prop(); a = merge(a->l, a->r); }

pair<ll, vi> dmst(int n, int r, vector<Edge>& g) {
    RollbackUF uf(n);
    vector<Node*> heap(n);
    for (Edge e : g) heap[e.b] = merge(heap[e.b], new Node{e});
    ll res = 0;
    vi seen(n, -1), path(n), par(n);
    seen[r] = r;
    vector<Edge> Q(n), in(n, {-1, -1}), comp;
    deque<tuple<int, int, vector<Edge>>> cys;
    rep(s, 0, n) {
        int u = s, qi = 0, w;
        while (seen[u] < 0) {
            if (!heap[u]) return {-1, {}};
            Edge e = heap[u]->top();
            heap[u]->delta -= e.w, pop(heap[u]);
            Q[qi] = e, path[qi++] = u, seen[u] = s;
            res += e.w, u = uf.find(e.a);
            if (seen[u] == s) { /// found cycle, contract
                Node* cyc = 0;
                int end = qi, time = uf.time();
                do cyc = merge(cyc, heap[w = path[--qi]]);
                while (uf.join(u, w));
                u = uf.find(u), heap[u] = cyc, seen[u] = -1;
                cys.push_front({u, time, {&Q[qi], &Q[end]}});
            }
        }
        rep(i, 0, qi) in[uf.find(Q[i].b)] = Q[i];
    }

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

(D + 1)-edge coloring

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

Usage: vi res = edgeColoring(N, eds);

Complexity: $\mathcal{O}(NM)$.

```
-----
e210e2
vi edgeColoring(int N, vector<pii> eds) {
    vi cc(N + 1), ret(sz(eds)), fan(N), free(N), loc;
    for (pii e : eds) ++cc[e.first], ++cc[e.second];
    int u, v, ncols = *max_element(all(cc)) + 1;
    vector<vi> adj(N, vi(ncols, -1));
    for (pii e : eds) {
        tie(u, v) = e;
        fan[0] = v;
        loc.assign(ncols, 0);
        int at = u, end = u, d, c = free[u], ind = 0, i = 0;
```

```
while (d = free[v], !loc[d] && (v = adj[u][d]) != -1)
    loc[d] = ++ind, cc[ind] = d, fan[ind] = v;
cc[loc[d]] = c;
for (int cd = d; at != -1; cd ^= c ^ d, at = adj[at][cd])
    swap(adj[at][cd], adj[end = at][cd ^ c ^ d]);
while (adj[fan[i]][d] != -1) {
    int left = fan[i], right = fan[++i], e = cc[i];
    adj[u][e] = left;
    adj[left][e] = u;
    adj[right][e] = -1;
    free[right] = e;
}
adj[u][d] = fan[i];
adj[fan[i]][d] = u;
for (int y : {fan[0], u, end})
    for (int& z = free[y] = 0; adj[y][z] != -1; z++);
}
rep(i,0,sz(eds))
for (tie(u, v) = eds[i]; adj[u][ret[i]] != v;) ++ret[i];
return ret;
}
```

Edmonds-Karp

Description: Yoinked from kactl. Flow algorithm with guaranteed complexity $\mathcal{O}(VE^2)$. To get edge flow values, compare capacities before and after, and take the positive values only.
Usage: edmondsKarp(graph, source, sink);
Complexity: $\mathcal{O}(EV^2)$.

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

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

        rep(i,0,ptr) {
            int x = q[i];
            for (auto e : graph[x]) {
                if (par[e.first] == -1 && e.second > 0) {
                    par[e.first] = x;
                    q[ptr++] = e.first;
                    if (e.first == sink) goto out;
                }
            }
        }
        return flow;
    out:
    T inc = numeric_limits<T>::max();
    for (int y = sink; y != source; y = par[y])
        inc = min(inc, graph[par[y]][y]);

    flow += inc;
    for (int y = sink; y != source; y = par[y]) {
        int p = par[y];
        if ((graph[p][y] -= inc) <= 0) graph[p].erase(y);
        graph[y][p] += inc;
    }
}
```

Floyd-Warshall

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

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

General matching

Description: Yoinked from kactl. Matching for general graphs. Finds a maximum subset of edges such that each vertex is incident to at most one edge. Fails with probability $\frac{N}{\text{mod}}$.
Usage: generalMatching(N, ed)
Complexity: $\mathcal{O}(N^3)$.

```
#include "../Maths/Matrix_inverse_mod.h"

vector<pii> generalMatching(int N, vector<pii>& ed) {
    vector<vector<ll>> mat(N, vector<ll>(N)), A;
    for (pii pa : ed) {
        int a = pa.first, b = pa.second, r = rand() % mod;
        mat[a][b] = r, mat[b][a] = (mod - r) % mod;
    }

    int r = matInv(A = mat), M = 2*N - r, fi, fj;
    assert(r % 2 == 0);

    if (M != N) do {
        mat.resize(M, vector<ll>(M));
        rep(i,0,M) {
            mat[i].resize(M);
            rep(j,N,M) {
                int r = rand() % mod;
                mat[i][j] = r, mat[j][i] = (mod - r) % mod;
            }
        }
    } while (matInv(A = mat) != M);

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

```
}
return ret;
}
```

Global minimum cut

Description: Yoinked from kactl. Finds a global minimum cut in an undirected graph, as represented by an adjacency matrix.
Usage: pair<int, vi> res = globalMinCut(mat);
Complexity: $\mathcal{O}(V^3)$.

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

Heavy-light decomposition

Description: Yoinked from kactl. 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. NOTE: below implementation uses kactl lazy segtree, this detail must be modified!
Usage: HLD <false> hld(adj); hld.query_path(u, v); ...
Complexity: $\mathcal{O}(\log n)$ segtree operations per operation.

```
#include "...kactl_segtree..."

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



```

    pos[v] = tim++;
    for (int u : adj[v]) {
        rt[u] = (u == adj[v][0] ? rt[v] : u);
        dfsHld(u);
    }
}
template <class B> void process(int u, int v, B op) {
    for (; rt[u] != rt[v]; v = par[rt[v]]) {
        if (depth[rt[u]] > depth[rt[v]]) swap(u, v);
        op(pos[rt[v]], pos[v] + 1);
    }
    if (depth[u] > depth[v]) swap(u, v);
    op(pos[u] + VALS_EDGES, pos[v] + 1);
}
void modifyPath(int u, int v, int val) {
    process(u, v, [&](int l, int r) { tree->add(l, r, val); });
}
int queryPath(int u, int v) { // Modify depending on
    problem
    int res = -1e9;
    process(u, v, [&](int l, int r) {
        res = max(res, tree->query(l, r));
    });
    return res;
}
int querySubtree(int v) { // modifySubtree is similar
    return tree->query(pos[v] + VALS_EDGES, pos[v] + siz[v]
    ]);
}
};

```

Hopcroft-Karp Bipartite Matching

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

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

Complexity: $\mathcal{O}(\sqrt{VE})$.

```

-----f612e4
bool dfs(int a, int L, vector<vi>& g, vi& btoa, vi& A, vi&
    B) {
    if (A[a] != L) return 0;
    A[a] = -1;
    for (int b : g[a]) if (B[b] == L + 1) {
        B[b] = 0;
        if (btoa[b] == -1 || dfs(btoa[b], L + 1, g, btoa, A, B)
        )
            return btoa[b] = a, 1;
    }
    return 0;
}

int hopcroftKarp(vector<vi>& g, vi& btoa) {
    int res = 0;
    vi A(g.size()), B(btoa.size()), cur, next;
    for (;;) {
        fill(all(A), 0);
        fill(all(B), 0);
        /// Find the starting nodes for BFS (i.e. layer 0).
        cur.clear();
        for (int a : btoa) if (a != -1) A[a] = -1;
        rep(a, 0, sz(g)) if (A[a] == 0) cur.push_back(a);
        /// Find all layers using bfs.
        for (int lay = 1; lay++) {
            bool islast = 0;
            next.clear();
            for (int a : cur) for (int b : g[a]) {
                if (btoa[b] == -1) {

```

```

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

```

Link-cut tree

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

Usage: See comments in code.

Complexity: Amortized $\mathcal{O}(\log n)$ per (any) operation.

```

-----5909e2
struct Node { // Splay tree. Root's pp contains tree's
    parent.
    Node *p = 0, *pp = 0, *c[2];
    bool flip = 0;
    Node() { c[0] = c[1] = 0; fix(); }
    void fix() {
        if (c[0]) c[0]->p = this;
        if (c[1]) c[1]->p = this;
        /// (+ update sum of subtree elements etc. if wanted)
    }
    void pushFlip() {
        if (!flip) return;
        flip = 0; swap(c[0], c[1]);
        if (c[0]) c[0]->flip ^= 1;
        if (c[1]) c[1]->flip ^= 1;
    }
    int up() { return p ? p->c[1] == this : -1; }
    void rot(int i, int b) {
        int h = i ^ b;
        Node *x = c[i], *y = b == 2 ? x : x->c[h], *z = b ? y :
        x;
        if ((y->p == p)) p->c[up()] = y;
        c[i] = z->c[i ^ 1];
        if (b < 2) {
            x->c[h] = y->c[h ^ 1];
            z->c[h ^ 1] = b ? x : this;
        }
        y->c[i ^ 1] = b ? this : x;
        fix(); x->fix(); y->fix();
        if (p) p->fix();
        swap(pp, y->pp);
    }
    void splay() { /// Splay this up to the root. Always
        finishes without flip set.
        for (pushFlip(); p; ) {
            if (p->p) p->p->pushFlip();
            p->pushFlip(); pushFlip();
            int c1 = up(), c2 = p->up();
            if (c2 == -1) p->rot(c1, 2);
            else p->p->rot(c2, c1 != c2);
        }
    }
    Node* first() { /// Return the min element of the subtree
        rooted at this, splayed to the top.

```

```

        pushFlip();
        return c[0] ? c[0]->first() : (splay(), this);
    }
};

struct LinkCut {
    vector<Node> node;
    LinkCut(int N) : node(N) {}

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

```

Maximum clique callbacks

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

Usage: `cliques(eds, callback, ...);`

Complexity: $\mathcal{O}(3^{n/3})$ - much faster for sparse graphs.

```

-----b045b1
typedef bitset<128> B;
template<class F>
void cliques(vector<B>& eds, F f, B P = ~B(), B X={}, B R
    ={}) {
    if (!P.any()) { if (!X.any()) f(R); return; }
    auto q = (P | X)._Find_first();
    auto cand = P & ~eds[q];
    rep(i, 0, sz(eds)) if (cand[i]) {
        R[i] = 1;

```



```
    cliques(eds, f, P & eds[i], X & eds[i], R);
    R[i] = P[i] = 0; X[i] = 1;
}
}
```

Maximum clique

Description: Yoinked from kactl. Finds a maximum clique of a graph given as a symmetric bitset matrix. Can be used to find a maximum independent set by finding a clique of the complement graph.
Complexity: About 1 second for $n = 155$, worst case random graphs ($p = .90$). Runs faster for sparse graphs.

```
-----_f7c0bc
typedef vector<bitset<200>> vb;
struct MaxClique {
    double limit=0.025, pk=0;
    struct Vertex { int i, d=0; };
    typedef vector<Vertex> vv;
    vb e;
    vv V;
    vector<vi> C;
    vi qmax, q, S, old;
    void init(vv& r) {
        for (auto& v : r) v.d = 0;
        for (auto& v : r) for (auto j : r) v.d += e[v.i][j.i];
        sort(all(r), [](auto a, auto b) { return a.d > b.d; });
        int mxD = r[0].d;
        rep(i,0,sz(r)) r[i].d = min(i, mxD) + 1;
    }
    void expand(vv& R, int lev = 1) {
        S[lev] += S[lev - 1] - old[lev];
        old[lev] = S[lev - 1];
        while (sz(R)) {
            if (sz(q) + R.back().d <= sz(qmax)) return;
            q.push_back(R.back().i);
            vv T;
            for(auto v:R) if (e[R.back().i][v.i]) T.push_back({v.i});
            if (sz(T)) {
                if (S[lev]++ / ++pk < limit) init(T);
                int j = 0, mxk = 1, mnk = max(sz(qmax) - sz(q) + 1, 1);
                C[1].clear(), C[2].clear();
                for (auto v : T) {
                    int k = 1;
                    auto f = [&](int i) { return e[v.i][i]; };
                    while (any_of(all(C[k]), f)) k++;
                    if (k > mxk) mxk = k, C[mxk + 1].clear();
                    if (k < mnk) T[j++] .i = v.i;
                    C[k].push_back(v.i);
                }
                if (j > 0) T[j - 1].d = 0;
                rep(k,mnk,mxk + 1) for (int i : C[k])
                    T[j].i = i, T[j++].d = k;
                expand(T, lev + 1);
            } else if (sz(q) > sz(qmax)) qmax = q;
            q.pop_back(), R.pop_back();
        }
    }
    vi maxClique() { init(V), expand(V); return qmax; }
    MaxClique(vb conn) : e(conn), C(sz(e)+1), S(sz(C)), old(S)
    ) {
        rep(i,0,sz(e)) V.push_back({i});
    }
};
```

Minimum cost maximum flow

Description: Yoinked from kactl. $cap[i][j] \neq cap[j][i]$ is allowed; double edges are not. 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.

Complexity: $\mathcal{O}(E^2)$ o.O.

```
-----fe85cc
#include <bits/extc++.h>

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

struct MCMF {
    int N;
    vector<vi> ed, red;
    vector<VL> cap, flow, cost;
    vi seen;
    VL dist, pi;
    vector<pii> par;

    MCMF(int N) :
        N(N), ed(N), red(N), cap(N, VL(N)), flow(cap), cost(cap),
        seen(N), dist(N), pi(N), par(N) {}

    void addEdge(int from, int to, ll cap, ll cost) {
        this->cap[from][to] = cap;
        this->cost[from][to] = cost;
        ed[from].push_back(to);
        red[to].push_back(from);
    }

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

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

        auto relax = [&](int i, ll cap, ll cost, int dir) {
            ll val = di - pi[i] + cost;
            if (cap && val < dist[i]) {
                dist[i] = val;
                par[i] = {s, dir};
                if (its[i] == q.end()) its[i] = q.push({-dist[i], i});
            } else q.modify(its[i], {-dist[i], i});
        };

        while (!q.empty()) {
            s = q.top().second; q.pop();
            seen[s] = 1; di = dist[s] + pi[s];
            for (int i : ed[s]) if (!seen[i])
                relax(i, cap[s][i] - flow[s][i], cost[s][i], 1);
            for (int i : red[s]) if (!seen[i])
                relax(i, flow[i][s], -cost[i][s], 0);
        }
        rep(i,0,N) pi[i] = min(pi[i] + dist[i], INF);
    }

    pair<ll, ll> maxflow(int s, int t) {
        ll totflow = 0, totcost = 0;
        while (path(s), seen[t]) {
            ll f1 = INF;
            for (int p,r,x = t; tie(p,r) = par[x], x != s; x = p)
                f1 = min(f1, r ? cap[p][x] - flow[p][x] : flow[x][p]);
            totflow += f1;
            for (int p,r,x = t; tie(p,r) = par[x], x != s; x = p)
                if (r) flow[p][x] += f1;
                else flow[x][p] -= f1;
        }
        rep(i,0,N) rep(j,0,N) totcost += cost[i][j] * flow[i][j];
    }
};
```

```
    return {totflow, totcost};
}

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

Minimum vertex cover

Description: Yoinked from kactl. Finds a minimum vertex cover in a bipartite graph. The size is the same as the size of a maximum matching, and the complement is a maximum independent set.
Complexity: Idk, look code.

```
-----b09e48
#include "DFS_matching.h"

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

Strongly connected components

Description: Yoinked from kactl. Finds strongly connected components in a directed graph. If vertices u, v belong to the same component, we can reach u from v and vice versa.
Usage: `scc(graph, [&](vi& v) { ... })` visits all components in reverse topological order. `comp[i]` holds the component index of a node (a component only has edges to components with lower index). `ncomps` will contain the number of components.
Complexity: $\mathcal{O}(E + V)$.

```
-----76b5c9
vi val, comp, z, cont;
int Time, ncomps;
template<class G, class F> int dfs(int j, G& g, F& f) {
    int low = val[j] = ++Time, x; z.push_back(j);
    for (auto e : g[j]) if (comp[e] < 0)
        low = min(low, val[e] ? dfs(e,g,f));

    if (low == val[j]) {
        do {
            x = z.back(); z.pop_back();
            comp[x] = ncomps;
            cont.push_back(x);
        } while (x != j);
        f(cont); cont.clear();
    }
}
```

```
    ncomps++;
}
return val[j] = low;
}
template<class G, class F> void scc(G& g, F f) {
    int n = sz(g);
    val.assign(n, 0); comp.assign(n, -1);
    Time = ncomps = 0;
    rep(i,0,n) if (comp[i] < 0) dfs(i, g, f);
}
```

Topological sort

Description: Yoinked from kactl. Given is an oriented graph. Output is an ordering of vertices, such that there are edges only from left to right. If there are cycles, the returned list will have size smaller than n – nodes reachable from cycles will not be returned.
Usage: `vi res = topoSort(gr);`
Complexity: $\mathcal{O}(V + E)$.

```
-----66a137
vi topoSort(const vector<vi>& gr) {
    vi indeg(sz(gr)), ret;
    for (auto& li : gr) for (int x : li) indeg[x]++;
    queue<int> q; // use priority_queue for lexic. largest ans.
    rep(i,0,sz(gr)) if (indeg[i] == 0) q.push(i);
    while (!q.empty()) {
        int i = q.front(); // top() for priority queue
        ret.push_back(i);
        q.pop();
        for (int x : gr[i])
            if (--indeg[x] == 0) q.push(x);
    }
    return ret;
}
```

Weighted bipartite matching

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

```
-----1e0fe9
pair<int, vi> hungarian(const vector<vi> &a) {
    if (a.empty()) return {0, {}};
    int n = sz(a) + 1, m = sz(a[0]) + 1;
    vi u(n), v(m), p(m), ans(n - 1);
    rep(i,1,n) {
        p[0] = i;
        int j0 = 0; // add "dummy" worker 0
        vi dist(m, INT_MAX), pre(m, -1);
        vector<bool> done(m + 1);
        do { // dijkstra
            done[j0] = true;
            int i0 = p[j0], j1, delta = INT_MAX;
            rep(j,1,m) if (!done[j]) {
                auto cur = a[i0 - 1][j - 1] - u[i0] - v[j];
                if (cur < dist[j]) dist[j] = cur, pre[j] = j0;
                if (dist[j] < delta) delta = dist[j], j1 = j;
            }
            rep(j,0,m) {
                if (done[j]) u[p[j]] += delta, v[j] -= delta;
                else dist[j] -= delta;
            }
            j0 = j1;
        } while (p[j0]);
        while (j0) { // update alternating path
            int j1 = pre[j0];
```

```
            p[j0] = p[j1], j0 = j1;
        }
    }
    rep(j,1,m) if (p[j]) ans[p[j] - 1] = j - 1;
    return {-v[0], ans}; // min cost
}
```

Two SAT

Description: Yoinked from kactl. Solves 2-SAT.
Usage: `TwoSat ts(n)` where n is the number of variables. `ts.either(i,j)` means that either i or j must be true. `ts.setValue(i)` means that i must be true. `ts.atMostOne(1)` means that at most one of the variables in l can be true. `ts.solve()` returns true iff it is solvable. `ts.values` will contain one possible solution. Negated variables are represented by bit-inversions ($\sim x$).
Complexity: $\mathcal{O}(N + E)$ where N is the number of variables and E is the number of clauses.

```
-----5f9706
struct TwoSat {
    int N;
    vector<vi> gr;
    vi values; // 0 = false, 1 = true
    TwoSat(int n = 0) : N(n), gr(2*n) {}
    int addVar() { // (optional)
        gr.emplace_back();
        gr.emplace_back();
        return N++;
    }
    void either(int f, int j) {
        f = max(2*f, -1-2*f);
        j = max(2*j, -1-2*j);
        gr[f].push_back(j^1);
        gr[j].push_back(f^1);
    }
    void setValue(int x) { either(x, x); }
    void atMostOne(const vi& li) { // (optional)
        if (sz(li) <= 1) return;
        int cur = ~li[0];
        rep(i,2,sz(li)) {
            int next = addVar();
            either(cur, ~li[i]);
            either(cur, next);
            either(~li[i], next);
            cur = ~next;
        }
        either(cur, ~li[1]);
    }
    vi val, comp, z; int time = 0;
    int dfs(int i) {
        int low = val[i] = ++time, x; z.push_back(i);
        for(int e : gr[i]) if (!comp[e])
            low = min(low, val[e] ? : dfs(e));
        if (low == val[i]) do {
            x = z.back(); z.pop_back();
            comp[x] = low;
            if (values[x]>>1) == -1)
                values[x]>>1 = x&1;
        } while (x != i);
        return val[i] = low;
    }
    bool solve() {
        values.assign(N, -1);
        val.assign(2*N, 0); comp = val;
        rep(i,0,2*N) if (!comp[i]) dfs(i);
        rep(i,0,N) if (comp[2*i] == comp[2*i+1]) return 0;
        return 1;
    }
};
```

Maths

Chinese remainder theorem

Description: Yoinked from kactl. `crt(a, m, b, n)` computes x such that $x \equiv a \pmod m$, $x \equiv b \pmod n$. If $|a| < m$ and $|b| < n$, x will obey $0 \leq x < \text{lcm}(m, n)$. Assumes $mn < 2^{62}$.
Complexity: $\mathcal{O}(\log n)$.

```
-----9a1b35
#include "Euclid.h"

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

Continued fractions

Description: Yoinked from kactl. Given N and a real number $x \geq 0$, finds the closest rational approximation p/q with $p, q \leq N$. It will obey $|p/q - x| \leq 1/qN$. For consecutive convergents, $p_{k+1}q_k - q_{k+1}p_k = (-1)^k$. (p_k/q_k alternates between $> x$ and $< x$.) If x is rational, y eventually becomes ∞ ; if x is the root of a degree 2 polynomial then a 's eventually become cyclic.
Complexity: $\mathcal{O}(\log n)$.

```
-----dd6c5e
typedef double d; // for N ~ 1e7; long double for N ~ 1e9
pair<ll, ll> approximate(d x, ll N) {
    ll LP = 0, LQ = 1, P = 1, Q = 0, inf = LLONG_MAX; d y = x;
    for (;;) {
        ll lim = min(P ? (N-LP) / P : inf, Q ? (N-LQ) / Q : inf),
            a = (ll)floor(y), b = min(a, lim),
            NP = b*P + LP, NQ = b*Q + LQ;
        if (a > b) {
            // If b > a/2, we have a semi-convergent that gives
            // us a
            // better approximation; if b = a/2, we *may* have
            // one.
            // Return {P, Q} here for a more canonical
            // approximation.
            return (abs(x - (d)NP / (d)NQ) < abs(x - (d)P / (d)Q)
                ) ?
                make_pair(NP, NQ) : make_pair(P, Q);
        }
        if (abs(y = 1/(y - (d)a)) > 3*N) {
            return {NP, NQ};
        }
        LP = P; P = NP;
        LQ = Q; Q = NQ;
    }
}
```

Determinant

Description: Yoinked from kactl. Calculates determinant of a matrix. Destroys the matrix.
Complexity: $\mathcal{O}(N^3)$.

```
-----bd5cec
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;
}
```

Divisor Count

Description: Counts number of divisors

```
-----7ff470
ll divisor_cnt(ll n) {
    ll cnt = 1;
    map<ll, ll> factors = factorize(n);
    for (auto p : factors) cnt *= p.second+1;
    return cnt;
}
```

Sieve of Eratosthenes

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

Complexity: $lim = 100'000'000 \approx 0.8$ s. Runs 30% faster if only odd indices are stored.

```
-----7c144c
const int MAX_PR = 5'000'000;
bitset<MAX_PR> isprime;
vi eratosthenesSieve(int lim) {
    isprime.set(); isprime[0] = isprime[1] = 0;
    for (int i = 4; i < lim; i += 2) isprime[i] = 0;
    for (int i = 3; i*i < lim; i += 2) if (isprime[i])
        for (int j = i*i; j < lim; j += i*2) isprime[j] = 0;
    vi pr;
    rep(i,2,lim) if (isprime[i]) pr.push_back(i);
    return pr;
}
```

Euclid

Description: Yoinked from kactl. Finds two integers x and y , such that $ax + by = \gcd(a, b)$. If a and b are coprime, then x is the inverse of $a \pmod b$.

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

```
-----33ba8f
ll euclid(ll a, ll b, ll &x, ll &y) {
    if (!b) return x = 1, y = 0, a;
    ll d = euclid(b, a % b, y, x);
    return y -= a/b * x, d;
}
```

Fast fourier transform

Description: Yoinked from kactl. $\text{fft}(\mathbf{a})$ computes $\hat{f}(k) = \sum_x a[x] \exp(2\pi i \cdot kx/N)$ for all k . N must be a power of 2. Useful for convolution: $\text{conv}(\mathbf{a}, \mathbf{b}) = \mathbf{c}$, 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.

Complexity: $\mathcal{O}(n \log n)$ with $N = |A| + |B|$. (~ 1 s for $N = 2^{22}$)

```
-----3dd197
typedef complex<double> C;
typedef vector<double> vd;
void fft(vector<C>& a) {
    int n = sz(a), L = 31 - __builtin_clz(n);
    static vector<complex<long double>> R(2, 1);
    static vector<C> rt(2, 1); // (^ 10% faster if double)
    for (static int k = 2; k < n; k *= 2) {
        R.resize(n); rt.resize(n);
```

```
auto x = polar(1.0L, acos(-1.0L) / k);
rep(i,k,2*k) rt[i] = R[i] = i&1 ? R[i/2] * x : R[i/2];
}
vi rev(n);
rep(i,0,n) rev[i] = (rev[i / 2] | (i & 1) << L) / 2;
rep(i,0,n) if (i < rev[i]) swap(a[i], a[rev[i]]);
for (int k = 1; k < n; k *= 2)
    for (int i = 0; i < n; i += 2 * k) rep(j,0,k) {
        // C z = rt[j+k] * a[i+j+k]; // (25% faster if hand-
        rolled) // include-line
        auto x = (double *)&rt[j+k], y = (double *)&a[i+j+k];
        // exclude-line
        C z(x[0]*y[0] - x[1]*y[1], x[0]*y[1] + x[1]*y[0]);
        // exclude-line
        a[i + j + k] = a[i + j] - z;
        a[i + j] += z;
    }
}
vd conv(const vd& a, const vd& b) {
    if (a.empty() || b.empty()) return {};
    vd res(sz(a) + sz(b) - 1);
    int L = 32 - __builtin_clz(sz(res)), n = 1 << L;
    vector<C> in(n), out(n);
    copy(all(a), begin(in));
    rep(i,0,sz(b)) in[i].imag(b[i]);
    fft(in);
    for (C& x : in) x *= x;
    rep(i,0,n) out[i] = in[-i & (n - 1)] - conj(in[i]);
    fft(out);
    rep(i,0,sz(res)) res[i] = imag(out[i]) / (4 * n);
    return res;
}
```

Fast fourier transform under arbitrary MOD

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

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

```
-----499494
#include "FFT.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));
    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)) {
        ll av = ll(real(outl[i])+.5), cv = ll(imag(outs[i])+.5);
        ll bv = ll(imag(outl[i])+.5) + ll(real(outs[i])+.5);
        res[i] = ((av % M * cut + bv) % M * cut + cv) % M;
    }
    return res;
}
```

Fast sieve of Eratosthenes

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

Complexity: LIM= 1e9 ≈ 1.5 s. Utalizes cache locality.

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

Gauss-Jordan elimination

Description: Yoinked from CP-algorithms. The description is taken from CP-algorithms as well: Following is an implementation of Gauss-Jordan. Choosing the pivot row is done with heuristic: choosing maximum value in the current column. The input to the function gauss is the system matrix a . The last column of this matrix is vector b . The function returns the number of solutions of the system (0,1, or ∞). If at least one solution exists, then it is returned in the vector ans . Implementation notes:

- The function uses two pointers - the current column col and the current row row .
- For each variable x_i , the value $where(i)$ is the line where this column is not zero. This vector is needed because some variables can be independent.
- In this implementation, the current i th line is not divided by a_{ii} as described above, so in the end the matrix is not identity matrix (though apparently dividing the i th line can help reducing errors).
- After finding a solution, it is inserted back into the matrix - to check whether the system has at least one solution or not. If the test solution is successful, then the function returns 1 or inf, depending on whether there is at least one independent variable.

kactl also has code for solving linear systems somewhere in the document, if needed.

Complexity: $\mathcal{O}(\min(n, m) \cdot nm)$ – I.e. cubic.

```
-----4847fe
const double EPS = 1e-9;
const int INF = 2; // it doesn't actually have to be
                    infinity or a big number

int gauss (vector < vector<double> > a, vector<double> &
ans) {
    int n = (int) a.size();
    int m = (int) a[0].size() - 1;

    vector<int> where (m, -1);
    for (int col=0, row=0; col<m && row<n; ++col) {
        int sel = row;
        for (int i=row; i<n; ++i)
            if (abs (a[i][col]) > abs (a[sel][col]))
```

```
        sel = i;
    if (abs (a[sel][col]) < EPS)
        continue;
    for (int i=col; i<=m; ++i)
        swap (a[sel][i], a[row][i]);
    where[col] = row;

    for (int i=0; i<n; ++i)
        if (i != row) {
            double c = a[i][col] / a[row][col];
            for (int j=col; j<=m; ++j)
                a[i][j] -= a[row][j] * c;
        }
    ++row;
}

ans.assign (m, 0);
for (int i=0; i<m; ++i)
    if (where[i] != -1)
        ans[i] = a[where[i]][m] / a[where[i]][i];
for (int i=0; i<n; ++i) {
    double sum = 0;
    for (int j=0; j<m; ++j)
        sum += ans[j] * a[i][j];
    if (abs (sum - a[i][m]) > EPS)
        return 0;
}

for (int i=0; i<m; ++i)
    if (where[i] == -1)
        return INF;
return 1;
}
```

Integer determinant

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

Complexity: $O(n^3)$.

```
-----3313dc
const ll mod = 12345;
ll det(vector<vector<ll>>& a) {
    int n = sz(a); ll ans = 1;
    rep(i,0,n) {
        rep(j,i+1,n) {
            while (a[j][i] != 0) { // gcd step
                ll t = a[i][i] / a[j][i];
                if (t) rep(k,i,n)
                    a[i][k] = (a[i][k] - a[j][k] * t) % mod;
                swap(a[i], a[j]);
                ans *= -1;
            }
        }
        ans = ans * a[i][i] % mod;
        if (!ans) return 0;
    }
    return (ans + mod) % mod;
}
```

Integration

Description: Yoinked from kactl. 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.

Complexity: $O(n)$ evaluations of f .

```
-----4756fc
template<class F> double quad(double a, double b, F f,
    const int n = 1000) {
    double h = (b - a) / 2 / n, v = f(a) + f(b);
    rep(i,1,n*2)
```

```
        v += f(a + i*h) * (i&1 ? 4 : 2);
    return v * h / 3;
}
```

Linear Diophantine Equation

Description: See below

-----002852

A Linear Diophantine Equation (in two variables) is an equation of the general form:

$$ax + by = c$$

where a, b, c are given integers, and x, y are unknown integers.

The degenerate case

A degenerate case that need to be taken care of is when $a = b = 0$. It is easy to see that we either have no solutions or infinitely many solutions, depending on whether $c = 0$ or not. In the rest of this article, we will ignore this case.

Analytic solution

When $a \neq 0$ and $b \neq 0$, the equation $ax+by=c$ can be equivalently treated as either of the following:

$$\begin{gathered} ax \equiv c \pmod b, \\ by \equiv c \pmod a. \end{gathered}$$

Without loss of generality, assume that $b \neq 0$ and consider the first equation. When a and b are co-prime, the solution to it is given as

$$x \equiv ca^{-1} \pmod b,$$

where a^{-1} is the [modular inverse](module-inverse.md) of a modulo b .

When a and b are not co-prime, values of ax modulo b for all integer x are divisible by $g=\gcd(a, b)$, so the solution only exists when c is divisible by g . In this case, one of solutions can be found by reducing the equation by g :

$$(a/g) x \equiv (c/g) \pmod{b/g}.$$

By the definition of g , the numbers a/g and b/g are co-prime, so the solution is given explicitly as

$$\begin{cases} x \equiv (c/g)(a/g)^{-1} \pmod{b/g}, \\ y = \frac{c-ax}{b}. \end{cases}$$

Algorithmic solution

To find one solution of the Diophantine equation with 2 unknowns, you can use the [Extended Euclidean algorithm](extended-euclid-algorithm.md). First, assume that a and b are non-negative. When we apply Extended Euclidean algorithm for a and b , we can find their greatest common divisor g and 2 numbers x_g and y_g such that:

$$ax_g + by_g = g$$

If c is divisible by $g = \gcd(a, b)$, then the given Diophantine equation has a solution, otherwise it does not have any solution. The proof is straight-forward: a linear combination of two numbers is divisible by

their common divisor.

Now supposed that c is divisible by g , then we have:

$$ax_g \cdot \frac{c}{g} + by_g \cdot \frac{c}{g} = c$$

Therefore one of the solutions of the Diophantine equation is:

$$x_0 = x_g \cdot \frac{c}{g},$$

$$y_0 = y_g \cdot \frac{c}{g}.$$

The above idea still works when a or b or both of them are negative. We only need to change the sign of x_0 and y_0 when necessary.

Finally, we can implement this idea as follows (note that this code does not consider the case $a = b = 0$):

```
/*
int gcd(int a, int b, int& x, int& y) {
    if (b == 0) {
        x = 1;
        y = 0;
        return a;
    }
    int x1, y1;
    int d = gcd(b, a % b, x1, y1);
    x = y1;
    y = x1 - y1 * (a / b);
    return d;
}

bool find_any_solution(int a, int b, int c, int& x0, int& y0, int& g) {
    g = gcd(abs(a), abs(b), x0, y0);
    if (c % g) {
        return false;
    }

    x0 *= c / g;
    y0 *= c / g;
    if (a < 0) x0 = -x0;
    if (b < 0) y0 = -y0;
    return true;
}
'''
/*
## Getting all solutions

From one solution  $(x_0, y_0)$ , we can obtain all the solutions of the given equation.

Let  $g = \gcd(a, b)$  and let  $x_0, y_0$  be integers which satisfy the following:

 $ax_0 + by_0 = c$ 

Now, we should see that adding  $b / g$  to  $x_0$ , and, at the same time subtracting  $a / g$  from  $y_0$  will not break the equality:

 $a \cdot \left(x_0 + \frac{b}{g}\right) + b \cdot \left(y_0 - \frac{a}{g}\right) = a \cdot x_0 + b \cdot y_0 + a \cdot \frac{b}{g} - b \cdot \frac{a}{g} = c$ 

Obviously, this process can be repeated again, so all the numbers of the form:


$$x = x_0 + k \cdot \frac{b}{g}$$


$$y = y_0 - k \cdot \frac{a}{g}$$


are solutions of the given Diophantine equation.
```


Moreover, this is the set of all possible solutions of the given Diophantine equation.

Finding the number of solutions and the solutions in a given interval

From previous section, it should be clear that if we don't impose any restrictions on the solutions, there would be infinite number of them. So in this section, we add some restrictions on the interval of x and y , and we will try to count and enumerate all the solutions.

Let there be two intervals: $[min_x; max_x]$ and $[min_y; max_y]$ and let's say we only want to find the solutions in these two intervals.

Note that if a or b is 0 , then the problem only has one solution. We don't consider this case here.

First, we can find a solution which have minimum value of x , such that $x \geq min_x$. To do this, we first find any solution of the Diophantine equation. Then, we shift this solution to get $x \geq min_x$ (using what we know about the set of all solutions in previous section). This can be done in $O(1)$.

Denote this minimum value of x by l_{x1} .

Similarly, we can find the maximum value of x which satisfy $x \leq max_x$. Denote this maximum value of x by r_{x1} .

Similarly, we can find the minimum value of y $(y \geq min_y)$ and maximum values of y $(y \leq max_y)$. Denote the corresponding values of x by l_{x2} and r_{x2} .

The final solution is all solutions with x in intersection of $[l_{x1}, r_{x1}]$ and $[l_{x2}, r_{x2}]$. Let denote this intersection by $[l_x, r_x]$.

Following is the code implementing this idea. Notice that we divide a and b at the beginning by g . Since the equation $ax + by = c$ is equivalent to the equation $\frac{a}{g}x + \frac{b}{g}y = \frac{c}{g}$, we can use this one instead and have $\gcd(\frac{a}{g}, \frac{b}{g}) = 1$, which simplifies the formulas.

```
void shift_solution(int & x, int & y, int a, int b, int cnt) {
    x += cnt * b;
    y -= cnt * a;
}

int find_all_solutions(int a, int b, int c, int minx, int maxx, int miny, int maxy) {
    int x, y, g;
    if (!find_any_solution(a, b, c, x, y, g))
        return 0;
    a /= g;
    b /= g;

    int sign_a = a > 0 ? +1 : -1;
    int sign_b = b > 0 ? +1 : -1;

    shift_solution(x, y, a, b, (minx - x) / b);
    if (x < minx)
        shift_solution(x, y, a, b, sign_b);
    if (x > maxx)
        return 0;
    int lx1 = x;

    shift_solution(x, y, a, b, (maxx - x) / b);
    if (x > maxx)
        shift_solution(x, y, a, b, -sign_b);
    int rx1 = x;
```

```
    shift_solution(x, y, a, b, -(miny - y) / a);
    if (y < miny)
        shift_solution(x, y, a, b, -sign_a);
    if (y > maxy)
        return 0;
    int lx2 = x;

    shift_solution(x, y, a, b, -(maxy - y) / a);
    if (y > maxy)
        shift_solution(x, y, a, b, sign_a);
    int rx2 = x;

    if (lx2 > rx2)
        swap(lx2, rx2);
    int lx = max(lx1, lx2);
    int rx = min(rx1, rx2);

    if (lx > rx)
        return 0;
    return (rx - lx) / abs(b) + 1;
}
/*
Once we have  $l_x$  and  $r_x$ , it is also simple to enumerate through all the solutions. Just need to iterate through  $x = l_x + k \cdot \frac{b}{g}$  for all  $k \geq 0$  until  $x = r_x$ , and find the corresponding  $y$  values using the equation  $ax + by = c$ .

## Find the solution with minimum value of  $x + y$  { data-toc-label='Find the solution with minimum value of <script type="math/tex">x + y</script>' }

Here,  $x$  and  $y$  also need to be given some restriction, otherwise, the answer may become negative infinity.

The idea is similar to previous section: We find any solution of the Diophantine equation, and then shift the solution to satisfy some conditions.

Finally, use the knowledge of the set of all solutions to find the minimum:

 $x' = x + k \cdot \frac{b}{g}$ ,
 $y' = y - k \cdot \frac{a}{g}$ .

Note that  $x + y$  change as follows:

 $x' + y' = x + y + k \cdot \left(\frac{b}{g} - \frac{a}{g}\right) = x + y + k \cdot \frac{b-a}{g}$ 

If  $a < b$ , we need to select smallest possible value of  $k$ . If  $a > b$ , we need to select the largest possible value of  $k$ . If  $a = b$ , all solution will have the same sum  $x + y$ .
```

Linear Recurrences

Description: Having a linear recurrence of the form $f(n) = a_1 \cdot f(n-1) + a_2 \cdot f(n-2) \dots$ can be solved in log time with matrix exponentation.

```
#define Matrix vector<vector<ll>>
const ll m = 1000000007;
Matrix operator*(const Matrix& a, const Matrix& b) {
    Matrix c = Matrix(len(a), vector<ll>(len(b[0])));
    for (int i = 0; i < len(a); i++) {
        for (int j = 0; j < len(b[0]); j++) {
            for (int k = 0; k < len(b); k++) {
                c[i][j] += a[i][k]*b[k][j]%m;
                c[i][j] %= m;
            }
        }
    }
}
```

```
    return c;
}
// DOES THIS WORK? Why dp needed?
Matrix fast_exp(const Matrix& a, ll b, map<ll, Matrix>& dp) {
    if (dp.count(b)) return dp[b];
    if (b == 1) return a;
    if (b%2) return dp[b] = fast_exp(a, b/2, dp)*fast_exp(a, b/2, dp)*a;
    return dp[b] = fast_exp(a, b/2, dp)*fast_exp(a, b/2, dp);
}
Matrix operator^(const Matrix& a, ll b) {
    map<ll, Matrix> dp;
    return fast_exp(a, b, dp);
}
void linear_recurrence() {
    /*
        dp[j] += dp[i] * X[i][j] <-- genral case
    */
}
```

Matrix inverse

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

Complexity: $O(n^3)$.

```
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;
        }

        /// forget A at this point, just eliminate tmp backward
        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;
    }
```

Matrix inverse mod prime

Description: Yoinked from kactl. Returns rank; result is stored in A unless singular ($\text{rank} < n$). For prime powers, repeatedly set $A^{-1} = A^{-1}(2I - AA^{-1}) \pmod{p^k}$ where A^{-1} starts as the inverse of $A \bmod p$, and k is increased in each step.

Complexity: $\mathcal{O}(n^3)$.

```
-----b4a981
#include "Mod_pow.h"

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

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

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

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

Miller-Rabin primality test

Description: Yoinked from kactl. Deterministic Miller-Rabin primality test. Guaranteed to work for numbers up to $7 \cdot 10^{18}$.

Complexity: 7 times the complexity of $a^b \bmod c$.

```
-----922927
#include "Mod_mul_LL.h"

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;
}
```

Modular inverses

Description: Yoinked from kactl. Pre-computation of modular inverses. Assumes $\text{LIM} \leq \text{mod}$ and that mod is a prime.

```
-----b4a981
// const ll mod = 1000000007, LIM = 200000; ///include-line
ll* inv = new ll[LIM] - 1; inv[1] = 1;
rep(i,2,LIM) inv[i] = mod - (mod / i) * inv[mod % i] % mod;
```

Modulo multiplication for 64-bit integers

Description: Yoinked from kactl. Calculate $a \cdot b \bmod c$ (or $a^b \bmod c$) for $0 \leq a, b \leq c \leq 7.2 \cdot 10^{18}$. This runs 2x faster than the naive $(_int128.t)a * b \% M$.

Complexity: $\mathcal{O}(1)$ for modmul, $\mathcal{O}(\log b)$ for modpow.

```
-----bbb88f
typedef unsigned long long ull;
ull modmul(ull a, ull b, ull M) {
    ll ret = a * b - M * ull(1.L / M * a * b);
    return ret + M * (ret < 0) - M * (ret >= (ll)M);
}

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

Mod pow

Description: Yoinked from kactl. What u think mans. (this interface is used by a few other things, hence included in the document)

Complexity: $\mathcal{O}(\log e)$.

```
-----b83e45
const ll mod = 1000000007; // faster if const

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

Modular arithmetic

Description: Yoinked from kactl. Simple operators for modular arithmetic. You need to set mod to some number first and then you can use the structure.

```
-----bf099e
#include "Euclid.h"

const ll mod = 17; // change to something else
struct Mod {
    ll x;
    Mod(ll xx) : x(xx) {}
    Mod operator+(Mod b) { return Mod((x + b.x) % mod); }
    Mod operator-(Mod b) { return Mod((x - b.x + mod) % mod); }
    Mod operator*(Mod b) { return Mod((x * b.x) % mod); }
    Mod operator/(Mod b) { return *this * invert(b); }
    Mod invert(Mod a) {
        ll x, y, g = euclid(a.x, mod, x, y);
        assert(g == 1); return Mod((x + mod) % mod);
    }
    Mod operator^(ll e) {
        if (!e) return Mod(1);
        Mod r = *this ^ (e / 2); r = r * r;
        return e&1 ? *this * r : r;
    }
};
```

Number theoretic transform

Description: Yoinked from kactl. $\text{ntt}(a)$ computes $\hat{f}(k) = \sum_x a[x]g^{xk}$ for all k , where $g = \text{root}^{(\text{mod}-1)/N}$. 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. $\text{conv}(a, b) = c$, where $c[x] = \sum a[i]b[x-i]$. For manual convolution: NTT the inputs, multiply pointwise, divide by n , reverse($\text{start}+1, \text{end}$), NTT back. Inputs must be in $[0, \text{mod})$.

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

```
-----9c7e10
#include "Mod_pow.h"

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

Polynomial root finding

Description: Yoinked from kactl. Finds the real roots to a polynomial. Usage: `polyRoots({{2,-3,1}}, -1e9, 1e9);` // solve $x^2-3x+2 = 0$

Complexity: $\mathcal{O}(n^2 \log(\frac{1}{\epsilon}))$.

```
-----7c7d9e
#include "Polynomial.h"

vector<double> polyRoots(Poly p, double xmin, double xmax) {
    if (sz(p.a) == 2) { return {-p.a[0]/p.a[1]}; }
    vector<double> ret;
    Poly der = p;
    der.diff();
    auto dr = polyRoots(der, xmin, xmax);
    dr.push_back(xmin-1);
    dr.push_back(xmax+1);
    sort(all(dr));
    rep(i,0,sz(dr)-1) {
        double l = dr[i], h = dr[i+1];
```

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

Polynomial thing

Description: Yoinked from kactl. Some poly things I guess.

```
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();
    }
};
```

SOS DP

Description: Some solution from some problem Elias solved. For each of n elements x : The number of elements y such that $x \mid y = x$. The number of elements y such that $x \& y = x$. The number of elements y such that $x \& y \neq 0$. NOTE: if TLE issues, try loop unrolling or C style arrays. Complexity: $\mathcal{O}(V \log V + n)$ where V is the maximum value.

```
constexpr const int lgmxV = 20;
constexpr const int mxV = 1 << lgmxV;

int main(){
    ios_base::sync_with_stdio(0); cin.tie(0); cout.tie(0);
    int n; cin >> n;
    vector<int> v(n);
    for(auto &x : v)cin >> x;
    vector<vector<int>> sos1(mxV, vector<int> (lgmxV + 1, 0));
    vector<vector<int>> sos2(mxV, vector<int> (lgmxV + 1, 0));
    for(int i = 0; i < n; ++i){
        sos1[v[i]][0]++;
        sos2[v[i] ^ (mxV - 1)][0]++;
    }
    for(int i = 0; i < mxV; ++i){
        for(int j = 0; j < lgmxV; ++j){
            sos1[i][j + 1] = sos1[i][j];
            sos2[i][j + 1] = sos2[i][j];
            if(i & (1 << j)) { sos1[i][j + 1] += sos1[i - (1 << j)][j]; };
            if(i & (1 << j)) { sos2[i][j + 1] += sos2[i - (1 << j)][j]; };
        }
    }
}
```

```
for(int i = 0; i < n; ++i){
    cout << sos1[v[i]][lgmxV] << ' ' << sos2[v[i] ^ (mxV - 1)][lgmxV] << ' ' << n - sos1[v[i] ^ (mxV - 1)][lgmxV] << '\n';
}
}
```

Simplex

Description: Yoinked from kactl. Solves a general linear maximization problem: maximize $c^T x$ subject to $Ax \leq b, x \geq 0$. Returns -inf if there is no solution, inf if there are arbitrarily good solutions, or the maximum value of $c^T x$ otherwise. The input vector is set to an optimal x (or in the unbounded case, an arbitrary solution fulfilling the constraints). Numerical stability is not guaranteed. For better performance, define variables such that $x = 0$ is viable. Usage: vvd A = 1,-1, -1,1, -1,-2; vd b = 1,1,-4, c = -1,-1, x; T val = LPSolver(A, b, c).solve(x); Complexity: $\mathcal{O}(NM \cdot \#pivots)$, 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 || 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;
                if (r == -1 || MP(D[i][n+1] / D[i][s], B[i]) < MP(D[r][n+1] / D[r][s], B[r])) r = i;
            }
            if (r == -1) return false;
        }
    }
};
```

```
    pivot(r, s);
}

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

Solve linear equations

Description: Yoinked from kactl. 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. Complexity: $\mathcal{O}(n^2m)$.

```
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)
}
```

Solve linear equations extended

Description: Yoinked from kactl. To get all uniquely determined values of x back from SolveLinear, make the following changes:

```
-----cf7d6d
#include "Solve_linear.h"

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:; }

```

Phi function

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

```
-----cf7d6d
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;
}

```

Strings

Aho-Corasick automaton

Description: Yoinked from kactl. 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. **Complexity:** $26 \cdot \mathcal{O}(N)$ to construct, where N = sum of length of patterns. find(x) is $\mathcal{O}(N)$, where N = length of x . findAll is $\mathcal{O}(NM)$.

```
f35677
-----
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) { m = m = sz(N); N.emplace_back(-1); }
            else m = 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;
}
};

```

String hashing

Description: Yoinked from kactl. Self-explanatory methods for string hashing.

```
-----2d2a67
// Arithmetic mod 2^64-1. 2x slower than mod 2^64 and more
// code, but works on evil test data (e.g. Thue-Morse,
// where
// ABBA... and BAAB... of length 2^10 hash the same mod 2^
// 64).
// "typedef ull H;" instead if you think test data is
// random,
// or work mod 10^9+7 if the Birthday paradox is not a
// problem.
typedef uint64_t ull;
struct H {
    ull x; H(ull x=0) : x(x) {}
    H operator+(H o) { return x + o.x + (x + o.x < x); }
    H operator-(H o) { return *this + ~o.x; }
    H operator*(H o) { auto m = (__uint128_t)x * o.x;
        return H((ull)m) + (ull)(m >> 64); }
    ull get() const { return x + !~x; }
}

```

```

    bool operator==(H o) const { return get() == o.get(); }
    bool operator<(H o) const { return get() < o.get(); }
};
static const H C = (1ll)1e11+3; // (order ~ 3e9; random also
    ok)

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

vector<H> getHashes(string& str, int length) {
    if (sz(str) < length) return {};
    H h = 0, pw = 1;
    rep(i,0,length)
        h = h * C + str[i], pw = pw * C;
    vector<H> ret = {h};
    rep(i,length,sz(str)) {
        ret.push_back(h = h * C + str[i] - pw * str[i-length]);
    }
    return ret;
}

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

```

Knuth-Morris-Pratt algorithm

Description: Yoinked from kactl. Finds all occurrences of a pattern in a string. p[x] computes the length of the longest prefix of s that ends at x, other than s[0...x] itself (abacaba -> 0010123). **Complexity:** $\mathcal{O}(n)$.

```
-----d4375c
vi pi(const string& s) {
    vi p(sz(s));
    rep(i,1,sz(s)) {
        int g = p[i-1];
        while (g && s[i] != s[g]) g = p[g-1];
        p[i] = g + (s[i] == s[g]);
    }
    return p;
}

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

```

Manacher

Description: Yoinked from kactl. 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). **Complexity:** $\mathcal{O}(n)$.

```
-----e7ad79
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;
}

```

Min rotation

Description: Yoinked from kactl. Finds the lexicographically smallest rotation of a string.

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

Complexity: $O(n)$.

```

-----d07a42
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;
}

```

Suffix array

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

Complexity: $O(n \log n)$ per update/query

```

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

```

Suffix automaton

Description: Standard suffix automaton. Does what you'd expect.

Usage: See example main function below. This was thrown in last minute from a working cses solution.

Complexity: $O(\log n)$ per update/query

```

-----34d234e
struct SA {
    struct State {
        int length;

```

```

        int link;
        int next[26];
        int cnt;
        bool is_clone;
        int first_pos;
        State(int _length, int _link) :
            length(_length),
            link(_link),
            cnt(0),
            is_clone(false),
            first_pos(-1)
        {
            memset(next, -1, sizeof(next));
        }
};
std::vector<State> states;
int size;
int last;
bool did_init_count;
int str_len;
bool did_init_css;
SA() :
    states(1, State(0, -1)),
    size(1),
    last(0),
    did_init_count(false),
    str_len(0),
    did_init_css(false)
{ }
void push(char c) {
    str_len++;
    did_init_count = false;
    did_init_css = false;
    int cur = size;
    states.resize(++size, State(states[last].length + 1,
        -1));
    states[cur].first_pos = states[cur].length - 1;
    int p = last;
    while (p != -1 && states[p].next[c - 'a'] == -1) {
        states[p].next[c - 'a'] = cur;
        p = states[p].link;
    }
    if (p == -1) {
        states[cur].link = 0;
    } else {
        int q = states[p].next[c - 'a'];
        if (states[p].length + 1 == states[q].length) {
            states[cur].link = q;
        } else {
            int clone = size;
            states.resize(++size, State(states[p].length + 1,
                states[q].link));
            states[clone].is_clone = true;
            memcpy(states[clone].next, states[q].next, sizeof(
                State::next));
            states[clone].first_pos = states[q].first_pos;
            while (p != -1 && states[p].next[c - 'a'] == q) {
                states[p].next[c - 'a'] = clone;
                p = states[p].link;
            }
            states[q].link = states[cur].link = clone;
        }
    }
    last = cur;
}
bool exists(const std::string& pattern) {
    int node = 0;
    int index = 0;
    while (index < (int) pattern.length() && states[node].
        next[pattern[index] - 'a'] != -1) {
        node = states[node].next[pattern[index] - 'a'];
        index++;
    }
}

```

```

    }
    return index == (int) pattern.size();
}
int count(const std::string& pattern) {
    if (!did_init_count) {
        did_init_count = true;
        for (int i = 1; i < size; i++) {
            states[i].cnt = !states[i].is_clone;
        }
        std::vector<std::vector<int>> of_length(str_len +
            1);
        for (int i = 0; i < size; i++) {
            of_length[states[i].length].push_back(i);
        }
        for (int l = str_len; l >= 0; l--) {
            for (int node : of_length[l]) {
                if (states[node].link != -1) {
                    states[states[node].link].cnt += states[node].
                        cnt;
                }
            }
        }
    }
    int node = 0;
    int index = 0;
    while (index < (int) pattern.length() && states[node].
        next[pattern[index] - 'a'] != -1) {
        node = states[node].next[pattern[index] - 'a'];
        index++;
    }
    return index == (int) pattern.size() ? states[node].cnt
        : 0;
}
int first_occ(const std::string& pattern) {
    int node = 0;
    int index = 0;
    while (index < (int) pattern.length() && states[node].
        next[pattern[index] - 'a'] != -1) {
        node = states[node].next[pattern[index] - 'a'];
        index++;
    }
    return index == (int) pattern.size() ? states[node].
        first_pos - (int) pattern.size() + 1 : -1;
}
size_t count_substrings() {
    static std::vector<size_t> dp;
    if (!did_init_css) {
        did_init_css = true;
        dp = std::vector<size_t>(size, 0);
        auto dfs = [&](auto&& self, int node) -> size_t {
            if (node == -1) {
                return 0;
            }
            if (dp[node]) {
                return dp[node];
            }
            dp[node] = 1;
            for (int i = 0; i < 26; i++) {
                dp[node] += self(self, states[node].next[i]);
            }
            return dp[node];
        };
        dfs(dfs, 0);
    }
    return dp[0] - 1;
}
}
// usage example: Repeating Substring submission on cses.fi
int main() {
    std::ios::sync_with_stdio(0); std::cin.tie(0);
    std::string s; std::cin >> s;
}

```

```
int n; std::cin >> n;
SA sa;
for (char c : s) {
    sa.push(c);
}
sa.count("");
int len = -1;
int ind = -1;
for (int i = 1; i < sa.size; i++) {
    if (sa.states[i].cnt > 1) {
        if (len < sa.states[i].length) {
            len = sa.states[i].length;
            ind = sa.states[i].first_pos - len + 1;
        }
    }
}
if (len == -1) {
    std::cout << "-1\n";
    return 0;
}
for (int i = 0; i < len; i++) {
    std::cout << s[i + ind];
}
std::cout << "\n";
}
```

Suffix tree

Description: Yoinked from kactl. 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).

Complexity: $26 \cdot \mathcal{O}(n)$.

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

    void ukkadd(int i, int c) { suff:
        if (r[v] <= q) {
            if (t[v][c] == -1) { t[v][c] = m; l[m] = i;
                p[m++] = v; v = s[v]; q = r[v]; goto suff; }
            v = t[v][c]; q = l[v];
        }
        if (q == -1 || c == toi(a[q])) q++; else {
            l[m+1] = i; p[m+1] = m; l[m] = 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; l[0] = l[1] = -1; r[0] = r[1] = p[0] = p[1] = 0;
        rep(i, 0, sz(a)) ukkadd(i, toi(a[i]));
    }
}
```

```
// example: find longest common substring (uses ALPHA = 28)
pii best;
int lcs(int node, int i1, int i2, int olen) {
    if (l[node] <= i1 && i1 < r[node]) return 1;
    if (l[node] <= i2 && i2 < r[node]) return 2;
    int mask = 0, len = node ? olen + (r[node] - l[node]) : 0;
    rep(c, 0, ALPHA) if (t[node][c] != -1)
        mask |= lcs(t[node][c], i1, i2, len);
    if (mask == 3)
        best = max(best, {len, r[node] - len});
    return mask;
}

static pii LCS(string s, string t) {
    SuffixTree st(s + (char)('z' + 1) + t + (char)('z' + 2));
    st.lcs(0, sz(s), sz(s) + 1 + sz(t), 0);
    return st.best;
}
};
```

Z function

Description: Yoinked from kactl. z[x] computes the length of the longest common prefix of s[1:] and s, except z[0] = 0. (abacaba -> 0010301)

Complexity: $\mathcal{O}(n)$.

```
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;
}
```

Various

Bump allocator

Description: Yoinked from kactl. 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.

Complexity: $\mathcal{O}(n)$.

```
// 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*) {}
```

Fast integer input

Description: Yoinked from kactl. USE THIS IF TRYING TO CONSTANT TIME OPTIMIZE SOLUTION READING IN LOTS OF INTEGERS!!! Read an integer from stdin. Usage requires your program to pipe in input from file.

Usage: ./a.out < input.txt

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

```
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 - 48;
    return a - 48;
}
```

Fast knapsack

Description: Yoinked from kactl. Given N non-negative integer weights w and a non-negative target t , computes the maximum $S \leq t$ such that S is the sum of some subset of the weights.

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

```
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;
}
```

Fast mod reduction

Description: Yoinked from kactl. 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 $[0, 2b)$. (proven correct)

```
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;
    }
};
```

Interval container

Description: Yoinked from kactl. 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).

Complexity: $\mathcal{O}(\log n)$ per update/query

```
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);
}

```

Interval cover

Description: Yoinked from kactl. 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).

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

```

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

```

Knuth DP optimization

Description: Yinked from kactl. When doing DP on intervals: $a[i][j] = \min_{i < k < j} (a[i][k] + a[k][j]) + f(i, j)$, where the (minimal) optimal k increases with both i and j , one can solve intervals in increasing

order of length, and search $k = p[i][j]$ for $a[i][j]$ only between $p[i][j-1]$ and $p[i+1][j]$. This is known as Knuth DP. Sufficient criteria for this are if $f(b, c) \leq f(a, d)$ and $f(a, c) + f(b, d) \leq f(a, d) + f(b, c)$ for all $a \leq b \leq c \leq d$.

Complexity: $\mathcal{O}(N^2)$.

```

// generic implementation from cp-algorithms:
int solve() {
    int N;
    ... // read N and input
    int dp[N][N], opt[N][N];
    auto C = [&](int i, int j) {
        ... // Implement cost function C.
    };
    for (int i = 0; i < N; i++) {
        opt[i][i] = i;
        ... // Initialize dp[i][i] according to the problem
    }
    for (int i = N-2; i >= 0; i--) {
        for (int j = i+1; j < N; j++) {
            int mn = INT_MAX;
            int cost = C(i, j);
            for (int k = opt[i][j-1]; k <= min(j-1, opt[i+1][j]); k++) {
                if (mn >= dp[i][k] + dp[k+1][j] + cost) {
                    opt[i][j] = k;
                    mn = dp[i][k] + dp[k+1][j] + cost;
                }
            }
            dp[i][j] = mn;
        }
    }
    cout << dp[0][N-1] << endl;
}

```

Longest increasing subsequence

Description: Yoinked from kactl. Computes the longest increasing subsequence of a sequence.

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

```

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

```

```

        *it = {S[i], i};
        prev[i] = it == res.begin() ? 0 : (it-1)->second;
    }
    int L = sz(res), cur = res.back().second;
    vi ans(L);
    while (L--) ans[L] = cur, cur = prev[cur];
    return ans;
}

```

Small ptr

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

```
#include "Bump_allocator.h"

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

Xor Basis

Description: basis of vectors in Z_2^d

```

int basis[d]; // basis[i] keeps the mask of the vector
              whose f value is i
int sz; // Current size of the basis
void insertVector(int mask) {
    for (int i = 0; i < d; i++) {
        if ((mask & 1 << i) == 0) continue; // continue if i !=
            f(mask)

        if (!basis[i]) { // If there is no basis vector with
            the i'th bit set, then insert this vector into the
            basis
            basis[i] = mask;
            ++sz;

            return;
        }

        mask ^= basis[i]; // Otherwise subtract the basis
            vector from this vector
    }
}

```