

North South University

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```
1 Contest
                                                                 #define all(v) v.begin(), v.end()
                                                                 #define input(v) for(auto&x:v)cin>>x;
2 Data structures
                                                                 #define print(v) for(auto&x:v)cout<<x<<' ';cout<<endl;</pre>
                                                                 #define dbg(a) cout<<#a<<" = "<<a<<endl;
3 Mathematics
                                                             6
                                                                 void solve() {}
4 Numerical
                                                             7
                                                                 int32_t main()
                                                                   ios_base::sync_with_stdio(0);
5 Number theory
                                                                  cin.tie(0); cout.tie(0);
                                                                   int t = 1, tc = 1;
                                                                  // cin >> t;
6 Combinatorial
                                                            12
                                                                   while (t--) {
                                                                    // cout \ll "Case" \ll tc++ \ll ":",
                                                                    solve();
7 Graph
                                                            13
8 Geometry
                                                                   return 0;
9 Strings
                                                                 RandomInputGenerator.cpp
                                                                 // Random input generator
10 Various
                                                                 auto seed = chrono::high_resolution_clock::now().
                                                                     time_since_epoch().count();
Contest (1)
                                                                 std::mt19937 mt(seed);
                                                                 int myrand(int mod) {
                                                                     return mt()%mod;
sublimeSetup.txt
                                                                 // Generates a random number within 100
// Tools -> Build System -> New Build System (write script &
                                                                 // int random_num = myrand(100) + 1;
// Tools -> Build System (select recently created script)
                                                                 DoubleInequality.cpp
"cmd" : ["q++ -std=c++20 $file_name -o $file_base_name &&
                                                                 // EPS | Double Inequality
    timeout 10s ./$file_base_name < input.txt > output.txt 2>
                                                                 const double eps = 1e-9;
    debug.txt && rm $file base name"],
                                                                 bool isEqual(double a, double b) {return abs(a-b) <= eps;} // a</pre>
"selector" : "source.cpp",
"shell": true,
"working_dir" : "$file_path"
                                                                 bool isSmaller(double a, double b) {return a+eps < b;} // a < b
                                                                 bool isGreater (double a, double b) {return a > b+eps;} // a > b
// Press 'Alt + Shift + 4' to split window in 4 parts.
                                                                 BitStuff.cpp
// save 'inputf.in', 'outputf.in', 'debugf.in'
                                                                 // Bit stuffs
// Press 'Ctrl + B' to run code.
                                                                 #define isOdd(n) n&1
/// Precompile HeaderFile
                                                                 #define isEven(n) !(n&1)
// just go to file explorer and serach 'stdc++.h'
// go to that folder and open folder in terminal
                                                                 #define div2pk(n, k) n>>k // n / pow(2, k)
                                                                 #define mul2pk(n, k) n<<k // n * pow(2, k)
// sudo g++ -std=c++20 stdc++.h
// stdc++.h.gch is created precompile done
                                                                 #define isDiv2pK(n, k) !(n & ((1<<k)-1)) // n \% pow(2, k) == 0
// preferences -> settings add "save_on_window_deactivation":
                                                                 #define isPowOf2(n) n && !(n&(n-1))
    true
                                                                 #define set(n, pos) (n \mid= (1<<pos))
// windows
                                                                 #define unset(n, pos) (n &= \sim(1<<pos))
"cmd": [ "q++.exe", "-std=c++14", "${file}", "-o", "${
                                                                 #define flip(n, pos) (n ^= (1<<pos))
    file_base_name}.exe", "&&", "${file_base_name}.exe",
                                                                 #define getBit(n, pos) (n & (1<<pos))</pre>
"<", "input.txt", ">", "output.txt", "2>", "debug.txt", "&&", "
    del", "${file_base_name}.exe"],
                                                                 // #pragma GCC target("popent")
                                                                 #define posL1(n) 32-__builtin_ctz(n)
template.cpp
                                                                 #define posR1(n) __builtin_ffs(n)
#include <bits/stdc++.h>
using namespace std;
                                                                 #define clearL1(n) n ^= (1 << (32-__builtin_ctz(n)-1))
#define endl '\n'
                                                                 #define clearR1(n) n &= (n-1)
#define 11 long long
```

#define countSetBits(n) __builtin_popcount(n)

#define len(v) (int) v.size()

```
// bitset
bitset<64> a, b(value), c("1011");
b.test(i); b[i]; // test i'th bit
// works on i'th bit or all
b.set(i); b.set();
b.reset(i); b.reset();
b.flip(i);b.flip();
b.count(); // counts set bits
b.to_string();
b.to_ulong();b.to_ullong();
CustomPQ.cpp
/// Custom Priority Queue
#define pii pair<int, int>
struct comp{
    bool operator() (pii& a, pii& b) {
        return a.second < b.second;</pre>
};
priority_queue<pii, vector<pii>, comp> pq;
Data structures (2)
CustomHashing.h
Description: Custom Hashing by Bashem vi.
Time: Usual
<ext/pb_ds/assoc_container.hpp>
                                                     6bc9dd, 18 lines
// For gp_hash_table
using namespace __gnu_pbds;
struct custom_hash {
    static uint64_t splitmix64(uint64_t x) {
        x += 0x9e3779b97f4a7c15;
        x = (x ^ (x >> 30)) * 0xbf58476d1ce4e5b9;
        x = (x ^ (x >> 27)) * 0x94d049bb133111eb;
        return x ^ (x >> 31);
    size_t operator()(uint64_t x) const {
        static const uint64_t FIXED_RANDOM = chrono::
             steady_clock::now().time_since_epoch().count();
        return splitmix64(x + FIXED_RANDOM);
};
unordered_map<int, int, custom_hash> mp; // Example Use
gp_hash_table<int, int, custom_hash> mp; // Faster
IndexedSet.h
Description: PBDS Indexed Set — Order Statistics Tree
Time: \mathcal{O}(log N)
<ext/pb_ds/assoc_container.hpp>, <ext/pb_ds/tree_policy.hpp>
                                                     0b0c96, 22 lines
// pbds set (with comparator)
using namespace __gnu_pbds;
struct Node { ll id, score, penalty; };
namespace std { template<> struct less<Node> {
    bool operator() (const Node& a, const Node& b) const {
        if (a.score > b.score) return true;
        if (a.score == b.score) {
            if (a.penalty == b.penalty) return a.id < b.id;</pre>
```

8 lines

38 lines

```
return a.penalty < b.penalty;</pre>
        return false;
};}
typedef tree<Node, null_type, less<Node>,
rb_tree_tag, tree_order_statistics_node_update> pbds;
// pbds s; s.insert(x);
// int value = *s.find_by_order(index);
// int index = s.order_of_key(value);
IndexedMultiset.h
Description: PBDS Indexed Multiset
Time: \mathcal{O}(log N)
<ext/pb_ds/assoc_container.hpp>, <ext/pb_ds/tree_policy.hpp>
                                                       edaf94, 20 lines
// pbds multiset // more like a indexed multiset
using namespace __gnu_pbds;
template < class T>
class multiset{
  using MS = tree<T, null_type, less_equal<T>,
  rb_tree_tag, tree_order_statistics_node_update>;
public:
  _multiset() {s.clear();}
  void erase(T xx){s.erase(s.upper_bound(xx));}
  typename MS::iterator lower_bound(T xx) {return s.upper_bound(
  typename MS::iterator upper bound(T xx) {return s.lower bound(
  // same
  size t size() {return s.size();}
  void insert(T xx){s.insert(xx);}
  T find by order(int xx) {return s.find by order(xx);}
  int order_of_key(T xx){return s.order_of_key(xx);}
  void erase(typename MS::iterator xx) {s.erase(xx);}
SparseTable.h
Description: Sparse Table. Range Minimum Queries on an array. Returns
\min(V[a], V[a+1], ... V[b-1]) in constant time.
Usage: SparseTable<int> table(v);
table.query(a, b); // [a, b) // index 0 based]
Time: \mathcal{O}(|V|\log|V|+Q)
                                                       2a5fd2, 17 lines
template<class T>
struct SparseTable {
    vector<vector<T>> jmp;
    SparseTable(const vector<T>& V) {
        int n = V.size();
        int log = 32 - __builtin_clz(n); // Maximum depth
        jmp.assign(log, V);
        for (int k = 1, pw = 1; pw * 2 <= n; ++k, pw *= 2)
            for (int i = 0; i + pw * 2 <= n; ++i)</pre>
                 jmp[k][i] = min(jmp[k - 1][i], jmp[k - 1][i +
                     ; ([wq
   T query(int a, int b) {
        assert(a < b);
        int dep = 31 - __builtin_clz(b - a); // log2(b - a)
        return min(jmp[dep][a], jmp[dep][b - (1 << dep)]);</pre>
};
```

MoQueries.h

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

```
Time: \mathcal{O}\left(N\sqrt{Q}\right)
<ext/pb_ds/assoc_container.hpp>, <ext/pb_ds/tree_policy.hpp>
                                                      75ee06, 54 lines
// mos
// pbds set // more like a indexed set
using namespace __gnu_pbds;
typedef tree<int, null_type, less<int>,
rb_tree_tag, tree_order_statistics_node_update> pbds;
void getMoAnswer(vector<int>& v, vector<array<int, 5>>& queries
     , vector<int>& ans) {
    pbds oset; // ordered set
    auto add = [&](int x) -> void { oset.insert(v[x]); };
    auto remove = [&](int x) -> void { oset.erase(v[x]); };
    auto get = [&](int k) -> int { return *oset.find_by_order(k
         -1); };
    sort (all (queries));
    int left = 0, right = -1;
    for (auto& [b, r, l, idx, k] : queries) {
        while(right < r) add(++right); while(right > r) remove
             (right--);
        while(left < 1) remove(left++); while(left > 1) add(--
             left);
        ans[idx] = get(k);
//v = main \ array, \ //N = v. size()
queries.push_back({l/sqrtN, r, l, idx, k}); // for each query
// sort quiries according to -> starting block, and then r wise
      sort
// gives k'th smallest number's index in [l, r) rangereturn res
// tree
vi moTree(vector<array<int, 2>> Q, vector<vi>& ed, int root=0) {
  int N = sz(ed), pos[2] = {}, blk = 350; // \sim N/sqrt(Q)
  vi s(sz(0)), res = s, I(N), L(N), R(N), in(N), par(N);
  add(0, 0), in[0] = 1;
  auto dfs = [&](int x, int p, int dep, auto& f) -> void {
    par[x] = p;
    L[x] = N;
    if (dep) I[x] = N++;
    for (int y : ed[x]) if (y != p) f(y, x, !dep, f);
    if (!dep) I[x] = N++;
    R[x] = N;
  dfs(root, -1, 0, dfs);
#define K(x) pii(I[x[0]] / blk, I[x[1]] ^ -(I[x[0]] / blk & 1))
  iota(all(s), 0);
  sort(all(s), [\&](int s, int t){ return K(Q[s]) < K(Q[t]); });
  for (int qi : s) rep(end, 0, 2) {
    int &a = pos[end], b = Q[qi][end], i = 0;
#define step(c) { if (in[c]) { del(a, end); in[a] = 0; } \
                  else { add(c, end); in[c] = 1; } a = c; }
    while (!(L[b] <= L[a] && R[a] <= R[b]))</pre>
     I[i++] = b, b = par[b];
    while (a != b) step(par[a]);
    while (i--) step(I[i]);
    if (end) res[qi] = calc();
  return res;
```

SegmentTree.h

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

```
Time: \mathcal{O}(\log N)
template <typename T>
struct SegmentTree {
    static constexpr T identity = INT_MIN;
    T f(T a, T b) { return max(a, b); }
    int n: vector<T> s:
    SegmentTree(int n = 0, T def = identity) : n(n), s(2*n, def
    SegmentTree(const vector<T>& v) : n(n), s(2*n, identity),
        { for (int i = 0; i < n; i++) { set(i, v[i]); } }
    void set(int pos, T val) {
        for (s[pos += n] = val; pos >>= 1;)
            s[pos] = f(s[pos << 1], s[pos << 1 | 1]);
    void update(int pos, T val) {
        for (s[pos += n] += val; pos >>= 1;)
            s[pos] = f(s[pos << 1], s[pos << 1 | 1]);
    T query(int b, int e) {
        T ra = identity, rb = identity;
        for (b += n, e += n; b < e; b >>= 1, e >>= 1) {
            if (b & 1) ra = f(ra, s[b++]);
            if (e & 1) rb = f(s[--e], rb);
        return f(ra, rb);
// SegmentTree < int > segTree(v);
// segTree.query(left-1, right); /l, r)
// segTree.set(index-1, increasingValue);
template <class T>
struct SegmentTree{
private:
    int n;
    vector<T> tree;
    void buildTree(const vector<T>& v, int node, int b, int e){
        if (b==e) {tree[node] = v[b]; return; }
        int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>
        buildTree(v, ln, b, mid);
        buildTree(v, rn, mid+1, e);
        tree[node] = merge(tree[ln], tree[rn]);
    T query(int node, int b, int e, int l, int r){
        if(l > e or r < b) return identity;</pre>
        if(1<=b and r>=e) return tree[node];
        int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>
        T c1 = query(ln, b, mid, l, r);
        T c2 = query(rn, mid+1, e, l, r);
        return merge(c1,c2);
    void set(int node, int b, int e, int ind, T val){
        if(ind > e or ind < b) return;</pre>
        if(ind<=b and ind>=e){
            tree[node] = val;
            return;
        int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>
        if (ind <= mid) set(ln, b, mid, ind, val);</pre>
        else set(rn, mid+1, e, ind, val);
```

tree[node] = merge(tree[ln], tree[rn]);

public:

LazySegmentTree MergeSortTree

```
vector < int > v(n); cin >> v;
    SegmentTree < int > segTree(v); // All 0 based index
    seqTree.query(left-1, right-1);
    segTree.set(index-1, value);
    segTree.update(index-1, increasingValue);
LazySegmentTree.h
Description: Lazy Segment Tree
Time: \mathcal{O}(\log N).
                                                      55c4d6, 115 lines
template <class T>
struct LazySegtree{
private:
  int n:
  vector<T> tree;
  vector<T> addTree, setTree;
  void buildTree(const vector<T>& v, int node, int b, int e){
    if (b==e) {tree[node] = v[b]; return; }
    int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>
    buildTree(v, ln, b, mid);
    buildTree(v, rn, mid+1, e);
    tree[node] = merge(tree[ln], tree[rn]);
  void propagate(int node, int b, int e){
    int ln = node<<1, rn = ln+1;</pre>
    if (setTree[node]!=set_identity) {
      addTree[node] = add_identity;
      tree[node] = setTree[node] * (e-b+1);
      if(b!=e){
        setTree[ln]=setTree[node];
        setTree[rn]=setTree[node];
      setTree[node]=set_identity;
```

void update(int node, int b, int e, int ind, T val){

tree[node] = merge(tree[node], val);

int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>

if (ind <= mid) update(ln, b, mid, ind, val);</pre>

T query(int 1, int r) {return query(1, 0, n-1, 1, r);}

void update(int ind, T val) {update(1, 0, n-1, ind, val);}

void set(int ind, T val){set(1, 0, n-1, ind, val);}

T merge(const T& a, const T& b) { return a + b; }

else update(rn, mid+1, e, ind, val);

tree[node] = merge(tree[ln], tree[rn]);

if(ind > e or ind < b) return;</pre>

SegmentTree(const vector<T>& input) {

buildTree(input, 1, 0, n-1);

if(ind<=b and ind>=e){

return:

n = input.size();

tree.resize(sz);

T identity = 0;

int sz = n<<2; // 4n

```
if (addTree[node] == add_identity) return;
    tree[node]+=addTree[node]*(e-b+1);
    if(b!=e){
      if (setTree[ln] == set_identity) {
        addTree[ln]+=addTree[node];
      else{
        setTree[ln]+=addTree[node];
        addTree[ln]=0;
      if(setTree[rn] == set_identity) {
        addTree[rn]+=addTree[node];
      else{
        setTree[rn]+=addTree[node];
        addTree[rn]=0;
    addTree[node] = add_identity;
T query(int node, int b, int e, int 1, int r){
  propagate (node, b, e);
  if(l > e or r < b) return identity;</pre>
  if(1<=b and r>=e) return tree[node];
  int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>
  T c1 = query(ln, b, mid, l, r);
  T c2 = query(rn, mid+1, e, l, r);
  return merge(c1,c2);
void range_set(int node, int b, int e, int l, int r, T val){
  propagate(node, b, e);
  if(l > e or r < b) return;</pre>
  if(1 \le b \text{ and } r \ge e)
    setTree[node]=val;
    propagate (node, b, e);
    return;
  int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>
  range set(ln, b, mid, l, r, val);
  range_set(rn, mid+1, e, l, r, val);
  tree[node]=merge(tree[ln], tree[rn]);
void range_update(int node, int b, int e, int l, int r, T val
  propagate (node, b, e);
  if(l > e or r < b) return;</pre>
  if(1<=b and r>=e){
    addTree[node]+=val;
    propagate (node, b, e);
    return;
  int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>
  range update(ln, b, mid, l, r, val);
  range_update(rn, mid+1, e, 1, r, val);
  tree[node] = merge(tree[ln], tree[rn]);
  return;
T query(int 1, int r) {return query(1, 0, n-1, 1, r);}
```

```
void range_set(int 1, int r, T value) { range_set(1, 0, n-1, 1
       , r, value);}
  void range_update(int 1, int r, T value){range_update(1, 0, n
       -1, 1, r, value);}
  LazySegtree(const vector<T>& input) {
    n = input.size();
    int sz = n<<2; // 4n
    tree.resize(sz);
    addTree.resize(sz, add_identity);
    setTree.resize(sz, set_identity);
    buildTree(input, 1, 0, n-1);
  T add_identity = 0;
 T \text{ set\_identity} = 0;
 T identity = 0;
 T merge(const T& a, const T& b) { return a + b; }
/* LazySeqtree<int> seqTree(v);
segTree.query(left-1, right-1);
seqTree.range\_set(left-1, right-1, value);
segTree.range\_update(left-1, right-1, value); */
MergeSortTree.h
Description: Merge Sort Tree
Time: n \log n + (\log n)2
                                                      e8d70e, 63 lines
template <class T>
struct SegmentTree{
private:
    vector<vector<T>> tree;
    // Build Tree n log n
    void buildTree(const vector<T>& v, int node, int b, int e){
        if (b==e) {tree[node] = {v[b]}; return;}
        int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>
        buildTree(v, ln, b, mid);
        buildTree(v, rn, mid+1, e);
        tree[node] = merge(tree[ln],tree[rn]);
    // Merge Nodes (just sort two nodes or vectors)
    vector<int> merge(vector<int> &a, vector<int> &b) {
        vector<int> c;
        int i = 0, j = 0;
        while (i < a.size() and j < b.size()) {</pre>
            if (a[i] < b[j]) c.push_back(a[i++]);</pre>
            else c.push_back(b[j++]);
        while (i < a.size()) c.push_back(a[i++]);</pre>
        while (j < b.size()) c.push_back(b[j++]);</pre>
        return c;
    // do binary search on the sorted node array(ofc if in
         range)
    int get(vector<int> &v, int k) {
        auto it = upper_bound(v.begin(), v.end(), k) - v.begin
        // return it; //number of elements strictly less than
              k in the range
        return v.size() - it; //number of elements strictly
             greater than k in the range
        // return \ v. size() - it - 1; \ // number of elements
             strictly greater than or equal to k in the range
```

Merge Sort Treew Point Update Wavelet Tree

```
int query(int node, int tL, int tR, int qL, int qR, int k)
        { (log n)^2
        if (tL >= qL && tR <= qR) {</pre>
            return get(tree[node], k);
        if (tR < qL || tL > qR) {
            return 0;
        int mid = (tL + tR) / 2;
        int QL = query(2 * node, tL, mid, qL, qR, k);
        int QR = query(2 * node + 1, mid + 1, tR, qL, qR, k);
        return OL + OR;
public:
    int query(int 1, int r, int k) {return query(1, 0, n-1, 1, r
    SegmentTree(const vector<T>& input) {
        n = input.size();
        int sz = n<<2; // 4n
        // tree.assign(vector<T>());
        tree.resize(sz);
        buildTree(input, 1, 0, n-1);
// vector < int > v(n); cin >> v;
// SegmentTree<int> segTree(v); // All 0 based index
// segTree.query(left - 1, right - 1, k);
MergeSortTreewPointUpdate.h
Description: Merge Sort Tree (w point update)
Time: n \log n + (\log n)2
<ext/pb_ds/assoc_container.hpp>, <ext/pb_ds/tree_policy.hpp>
                                                      8fd6e2, 74 lines
using namespace __gnu_pbds;
template <class T>
class multiset{
    using MS = tree<T, null_type, less_equal<T>,
    rb_tree_tag, tree_order_statistics_node_update>;
    MS s:
public:
    _multiset() {s.clear();}
    void erase(T xx){s.erase(s.upper_bound(xx));}
    typename MS::iterator lower_bound(T xx) {return s.
         upper_bound(xx);}
    typename MS::iterator upper_bound(T xx) {return s.
        lower_bound(xx);}
    size t size() { return s.size(); }
    void insert(T xx){s.insert(xx);}
    T find_by_order(int xx) {return s.find_by_order(xx);}
    int order_of_key(T xx){return s.order_of_key(xx);}
    void erase(typename MS::iterator xx){s.erase(xx);}
using T = long long;
int N;
vector<T> vec:
vector<_multiset<ll>> segtree;
void buildTree (int node, int b, int e) { //n \log n
    for (int i = b; i <= e; i++) {</pre>
        segtree[node].insert(vec[i]);
    if (b==e) return;
    int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>
    buildTree(ln, b, mid);
```

```
buildTree(rn, mid+1, e);
T query(int node, int b, int e, int 1, int r, T val) { //(log n)
    if(1 > e or r < b) return 0;</pre>
    if(l<=b and r>=e) return segtree[node].order_of_key(val);
    int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>
    T c1 = query(ln, b, mid, l, r, val);
    T c2 = query(rn, mid+1, e, l, r, val);
    return c1 + c2;
void setValue(int node, int b, int e, int ind, T val){
    segtree[node].erase(vec[ind]);
    segtree[node].insert(val);
    if (b==e) return;
    int mid = (b+e)>>1, ln = node<<1, rn = ln+1;</pre>
    if (ind <= mid) setValue(ln, b, mid, ind, val);</pre>
    else setValue(rn, mid+1, e, ind, val);
void buildTree(vector<T>& input) {
    N = input.size(); vec = input;
    int sz = N<<2; // 4n
    segtree.resize(sz);
    buildTree(1, 0, N-1);
T query(int 1, int r, T val) {return query(1, 0, N-1, 1, r, val)
void setValue(int ind, T val){
    setValue(1, 0, N-1, ind, val);
    vec[ind] = val;
     vector < int > v(n); input(v);
    buildTree(v); // All 0 based index
    query(left-1, right-1, value);
    set(index-1, value);
Wavelet Tree.h
Description: wavelet tree
\textbf{Time:} \ \ \text{Preprocess} \ \ \mathcal{O}\left(nlog(maxval)\right), \ \text{other query} \ \ \mathcal{O}\left(maxval\right)_{148656, \ 85 \ \text{lines}}
// Wavelet Tree
const int MAXN = (int) 3e5 + 9;
const int MAXV = (int)1e9 + 9; // maximum value of any element
// array values can be negative too, use appropriate minimum
     and maximum value
struct wavelet_tree {
    int lo, hi;
    wavelet_tree *1, *r;
    int *b, *c, bsz, csz; // c holds the prefix sum of elements
    wavelet_tree() {
        10 = 1; hi = 1;
        bsz = csz = 0;
        1 = r = NULL;
    void init(int *from, int *to, int x, int y) {
        lo = x, hi = y;
        if (from >= to) return;
```

```
int mid = (lo + hi) >> 1;
        auto f = [mid] (int x) { return x <= mid; };</pre>
        b = (int *)malloc((to - from + 2) * sizeof(int));
        bsz = 0; b[bsz++] = 0;
        c = (int *)malloc((to - from + 2) * sizeof(int));
        csz = 0; c[csz++] = 0;
        for (auto it = from; it != to; it++) {
            b[bsz] = (b[bsz - 1] + f(*it)); bsz++;
            c[csz] = (c[csz - 1] + (*it)); csz++;
        if (hi == lo) return;
        auto pivot = stable_partition(from, to, f);
        1 = new wavelet_tree();
        1->init(from, pivot, lo, mid);
        r = new wavelet_tree();
        r->init(pivot, to, mid + 1, hi);
    // kth smallest element in [l, r]
    int kth(int 1, int r, int k) {
        if (1 > r) return 0;
        if (lo == hi) return lo;
        int inLeft = b[r] - b[1 - 1], 1b = b[1 - 1], rb = b[r];
        if (k <= inLeft) return this->l->kth(lb + 1, rb, k);
        return this->r->kth(l - lb, r - rb, k - inLeft);
    // count of numbers in [l, r] Less than or equal to k
    int LTE(int 1, int r, int k) {
        if (1 > r || k < 10)
            return 0;
        if (hi <= k)
            return r - 1 + 1;
        int 1b = b[1 - 1], rb = b[r];
        return this->1->LTE(lb + 1, rb, k) + this->r->LTE(l -
            lb, r - rb, k);
    // count of numbers in [l, r] equal to k
    int count(int 1, int r, int k) {
        if (1 > r || k < lo || k > hi) return 0;
        if (lo == hi) return r - l + 1;
        int 1b = b[1 - 1], rb = b[r];
        int mid = (lo + hi) >> 1;
        if (k <= mid) return this->1->count(lb + 1, rb, k);
        return this->r->count(1 - 1b, r - rb, k);
    // sum of numbers in [l, r] less than or equal to k
    int sum(int 1, int r, int k) {
        if (1 > r or k < 10) return 0;
        if (hi <= k) return c[r] - c[l - 1];</pre>
        int 1b = b[1 - 1], rb = b[r];
        return this->1->sum(lb + 1, rb, k) + this->r->sum(l -
            lb, r - rb, k);
    ~wavelet tree() { delete 1; delete r; }
int a[MAXN]; // declare
wavelet tree t:
// 1 based \Rightarrow index, l, r
// int n; cin >> n; // size of array
// for (int i=1; i \le n; i++)cin>>a[i]; // array input
// O (n log ( max_ele(array) )), array a changes after init
// t.init(a + 1, a + n + 1, -MAXV, MAXV);
// [l, r] range, below O( max_ele(array)
// t.kth(l, r, k); // kth smallest element
// t.LTE(l, r, k); // count values <= k
// t.count(l, r, k); // count values == k
// t.sum(l, r, k); // sum of numbers <= k
```

UnionFindRollback.h

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

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

```
de4ad0, 21 lines
struct RollbackUF {
  vi e; vector<pii> st;
  RollbackUF(int n) : e(n, -1) {}
  int size(int x) { return -e[find(x)]; }
  int find(int x) { return e[x] < 0 ? x : find(e[x]); }</pre>
  int time() { return sz(st); }
  void rollback(int t) {
   for (int i = time(); i --> t;)
     e[st[i].first] = st[i].second;
   st.resize(t);
  bool join(int a, int b) {
   a = find(a), b = find(b);
   if (a == b) return false;
   if (e[a] > e[b]) swap(a, b);
   st.push_back({a, e[a]});
   st.push back({b, e[b]});
   e[a] += e[b]; e[b] = a;
    return true;
```

SubMatrix.h

};

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

```
Usage: SubMatrix<int> m (matrix);
```

m.sum(0, 0, 2, 2); // top left 4 elements

```
Time: \mathcal{O}\left(N^2+Q\right) c59ada, 13 lines
```

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

Matrix.h

Description: Basic operations on square matrices.

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

c43c7d, 26 lines

```
template < class T, int N> struct Matrix {
    typedef Matrix M;
    array < array < T, N>, N> d{};
    M operator*(const M& m) const {
        M a;
        rep(i,0,N) rep(j,0,N)
            rep(k,0,N) a.d[i][j] += d[i][k]*m.d[k][j];
    return a;
}
vector < T> operator*(const vector < T>& vec) const {
    vector < T> ret(N);
    rep(i,0,N) rep(j,0,N) ret[i] += d[i][j] * vec[j];
    return ret;
}
```

```
M operator^(ll p) const {
   assert(p >= 0);
   M a, b(*this);
   rep(i,0,N) a.d[i][i] = 1;
   while (p) {
      if (p&l) a = a*b;
      b = b*b;
      p >>= 1;
   }
   return a;
}
```

LineContainer.h

Description: Container where you can add lines of the form kx+m, and query maximum values at points x. Useful for dynamic programming ("convex hull trick").

```
Time: \mathcal{O}(\log N)
                                                       8ec1c7, 30 lines
struct Line {
 mutable 11 k, m, p;
 bool operator<(const Line& o) const { return k < o.k; }</pre>
 bool operator<(11 x) const { return p < x; }</pre>
struct LineContainer : multiset<Line, less<>>> {
 // (for doubles, use inf = 1/.0, div(a,b) = a/b)
 static const ll inf = LLONG_MAX;
 ll div(ll a, ll b) { // floored division
    return a / b - ((a ^ b) < 0 && a % b); }
 bool isect(iterator x, iterator y) {
    if (y == end()) return x \rightarrow p = inf, 0;
    if (x->k == y->k) x->p = x->m > y->m ? inf : -inf;
    else x->p = div(y->m - x->m, x->k - y->k);
    return x->p >= y->p;
 void add(ll k, ll m) {
    auto z = insert(\{k, m, 0\}), y = z++, x = y;
    while (isect(v, z)) z = erase(z);
    if (x != begin() \&\& isect(--x, y)) isect(x, y = erase(y));
    while ((y = x) != begin() \&\& (--x)->p >= y->p)
      isect(x, erase(y));
 11 query(11 x) {
    assert(!empty());
    auto 1 = *lower_bound(x);
    return 1.k * x + 1.m;
};
```

Treap.h

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

```
struct Node {
   Node *1 = 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(1) + cnt(r) + 1; }

template<class F> void each(Node* n, F f) {
   if (n) { each(n->1, f); f(n->val); each(n->r, f); }
}
```

```
pair<Node*, Node*> split(Node* n, int k) {
 if (!n) return {};
 if (cnt(n->1) >= k) { // "n-> val >= k" for lower_bound(k)}
    auto pa = split(n->1, k);
   n->1 = pa.second;
   n->recalc();
   return {pa.first, n};
    auto pa = split(n->r, k - cnt(n->1) - 1); // and just "k"
   n->r = pa.first;
   n->recalc();
   return {n, pa.second};
Node* merge(Node* 1, Node* r) {
 if (!1) return r;
 if (!r) return 1;
 if (1->y > r->y) {
   1->r = merge(1->r, r);
   1->recalc();
   return 1;
 } else {
    r->1 = merge(1, r->1);
    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 1, int r, int k) {
 Node *a, *b, *c;
 tie(a,b) = split(t, 1); tie(b,c) = split(b, r - 1);
 if (k \le 1) t = merge(ins(a, b, k), c);
 else t = merge(a, ins(c, b, k - r));
```

FenwickTree.h

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

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

```
struct FT {
  vector<ll> s;
 FT(int n) : s(n) {}
  void update(int pos, 11 dif) { // a[pos] \neq = dif
    for (; pos < sz(s); pos |= pos + 1) s[pos] += dif;</pre>
 11 query (int pos) { // sum of values in [0, pos)
    11 \text{ res} = 0;
    for (; pos > 0; pos &= pos - 1) res += s[pos-1];
    return res;
 int lower_bound(ll sum) \{// min \ pos \ st \ sum \ of \ [0, \ pos] >= sum
    // Returns n if no sum is >= sum, or -1 if empty sum is.
    if (sum \leq 0) return -1;
    int pos = 0;
    for (int pw = 1 << 25; pw; pw >>= 1) {
      if (pos + pw \leq sz(s) && s[pos + pw-1] \leq sum)
        pos += pw, sum -= s[pos-1];
    return pos;
```

FenwickTree2d.h

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

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

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

Mathematics (3)

3.1 Equations

};

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

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

$$ax + by = e$$

$$cx + dy = f$$

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

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

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

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

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

3.2 Recurrences

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

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

Non-distinct roots r become polynomial factors, e.g. $a_n = (d_1n + d_2)r^n$.

3.3 Trigonometry

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

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

$$\sin v + \sin w = 2\sin\frac{v+w}{2}\cos\frac{v-w}{2}$$

$$\cos v + \cos w = 2\cos\frac{v+w}{2}\cos\frac{v-w}{2}$$

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

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

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

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

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

3.4 Geometry

3.4.1 Triangles

Side lengths: a, b, c

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

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

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

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

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

Length of bisector (divides angles in two):

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

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

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

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

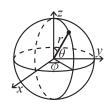
3.4.2 Quadrilaterals

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

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

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

3.4.3 Spherical coordinates



$$\begin{aligned} x &= r \sin \theta \cos \phi & r &= \sqrt{x^2 + y^2 + z^2} \\ y &= r \sin \theta \sin \phi & \theta &= \arccos(z/\sqrt{x^2 + y^2 + z^2}) \\ z &= r \cos \theta & \phi &= \operatorname{atan2}(y, x) \end{aligned}$$

3.5 Derivatives/Integrals

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

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

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

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

Integration by parts:

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

3.6 Sums

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

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

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

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

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

3.7 Series

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

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

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

$$\sin x = x - \frac{x^3}{3!} + \frac{x^5}{5!} - \frac{x^7}{7!} + \dots, (-\infty < x < \infty)$$
$$\cos x = 1 - \frac{x^2}{2!} + \frac{x^4}{4!} - \frac{x^6}{6!} + \dots, (-\infty < x < \infty)$$

3.8 Probability theory

Let X be a discrete random variable with probability $p_X(x)$ of assuming the value x. It will then have an expected value (mean) $\mu = \mathbb{E}(X) = \sum_x x p_X(x)$ and variance $\sigma^2 = V(X) = \mathbb{E}(X^2) - (\mathbb{E}(X))^2 = \sum_x (x - \mathbb{E}(X))^2 p_X(x)$ where σ is the standard deviation. If X is instead continuous it will have a probability density function $f_X(x)$ and the sums above will instead be integrals with $p_X(x)$ replaced by $f_X(x)$.

Expectation is linear:

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

For independent X and Y,

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

3.8.1 Discrete distributions Binomial distribution

The number of successes in n independent yes/no experiments, each which yields success with probability p is $Bin(n, p), n = 1, 2, ..., 0 \le p \le 1$.

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

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

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

First success distribution

The number of trials needed to get the first success in independent yes/no experiments, each which yields success with probability p is Fs(p), $0 \le p \le 1$.

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

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

Poisson distribution

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

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

3.8.2 Continuous distributions Uniform distribution

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

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

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

Exponential distribution

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

$$f(x) = \begin{cases} \lambda e^{-\lambda x} & x \ge 0\\ 0 & x < 0 \end{cases}$$
$$\mu = \frac{1}{\lambda}, \sigma^2 = \frac{1}{\lambda^2}$$

Normal distribution

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

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

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

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

3.9 Markov chains

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

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

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

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

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

Numerical (4)

4.1 Polynomials and recurrences

```
Polynomial.h
```

c9b7b0, 17 lines

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

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

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

PolyRoots.h

return ret;

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

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

PolyInterpolate.h

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

08bf48, 13 lines

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

BerlekampMassey.h

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

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

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

LinearRecurrence.h

Description: Generates the k'th term of an n-order linear recurrence $S[i] = \sum_j S[i-j-1]tr[j]$, given $S[0... \ge n-1]$ and tr[0...n-1]. Faster than matrix multiplication. Useful together with Berlekamp–Massey. Usage: linearRec($\{0, 1\}, \{1, 1\}, k\}$) // k'th Fibonacci number Time: $\mathcal{O}(n^2 \log k)$

```
typedef vector<11> Poly;
ll linearRec(Poly S, Poly tr, ll k) {
  int n = sz(tr);

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

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

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

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

4.2 Optimization

GoldenSectionSearch.h

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

```
Usage: double func(double x) { return 4+x+.3*x*x; } double xmin = gss(-1000,1000,func); 
Time: \mathcal{O}\left(\log((b-a)/\epsilon)\right)
```

31d45b, 14 lines

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

HillClimbing.h

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

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

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

Integrate.h

f4e444, 26 lines

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

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

```
return v * h / 3;
IntegrateAdaptive.h
Description: Fast integration using an adaptive Simpson's rule.
Usage: double sphereVolume = quad(-1, 1, [](double x) {
return quad(-1, 1, [&] (double y)
return quad(-1, 1, [\&] (double z) {
return \bar{x} \times x + y \times y + z \times z < 1; \}); \}); \}); \}
                                                           92dd79, 15 lines
typedef double d;
#define S(a,b) (f(a) + 4*f((a+b) / 2) + f(b)) * (b-a) / 6
template <class F>
d rec(F& f, da, db, deps, dS) {
  dc = (a + b) / 2;
  d S1 = S(a, c), S2 = S(c, b), T = S1 + S2;
  if (abs(T - S) <= 15 * eps || b - a < 1e-10)</pre>
    return T + (T - S) / 15;
  return rec(f, a, c, eps / 2, S1) + rec(f, c, b, eps / 2, S2);
template < class F>
d \text{ quad}(d \text{ a, } d \text{ b, } F \text{ f, } d \text{ eps} = 1e-8)  {
  return rec(f, a, b, eps, S(a, b));
```

Simplex.h

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

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

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

```
aa8530, 68 lines
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;

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

b[s] = a[s] * inv2;

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

Matrices

Determinant.h

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

bd5cec, 15 lines

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

IntDeterminant.h

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

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

3313dc, 18 lines

```
const 11 mod = 12345;
11 det(vector<vector<11>>& a) {
  int n = sz(a); ll ans = 1;
  rep(i,0,n) {
```

```
rep(j,i+1,n) {
    while (a[j][i] != 0) { // gcd step
      11 t = a[i][i] / a[j][i];
      if (t) rep(k,i,n)
        a[i][k] = (a[i][k] - a[j][k] * t) % mod;
      swap(a[i], a[j]);
      ans \star = -1;
  ans = ans * a[i][i] % mod;
  if (!ans) return 0;
return (ans + mod) % mod;
```

SolveLinear.h

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

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

SolveLinear2.h

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

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

```
fail:; }
```

SolveLinearBinarv.h

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

```
typedef bitset<1000> bs;
int solveLinear(vector<bs>& A, vi& b, bs& x, int m) {
 int n = sz(A), rank = 0, br;
 assert(m \le sz(x));
 vi col(m); iota(all(col), 0);
 rep(i,0,n) {
    for (br=i; br<n; ++br) if (A[br].any()) break;</pre>
    if (br == n) {
      rep(j,i,n) if(b[j]) return -1;
      break:
    int bc = (int)A[br]._Find_next(i-1);
    swap(A[i], A[br]);
    swap(b[i], b[br]);
    swap(col[i], col[bc]);
    rep(j,0,n) if (A[j][i] != A[j][bc]) {
     A[j].flip(i); A[j].flip(bc);
    rep(j,i+1,n) if (A[j][i]) {
     b[j] ^= b[i];
      A[j] ^= A[i];
    rank++;
  x = bs();
 for (int i = rank; i--;) {
    if (!b[i]) continue;
   x[col[i]] = 1;
    rep(j,0,i) b[j] ^= A[j][i];
 return rank; // (multiple solutions if rank < m)
```

MatrixInverse.h

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

Time: $\mathcal{O}(n^3)$ ebfff6, 35 lines

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

464cf3, 16 lines

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

Tridiagonal.h

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

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

This is useful for solving problems on the type

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

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

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

Fails if the solution is not unique.

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

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

8f9fa8, 26 lines

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

4.4 Fourier transforms

FastFourierTransform.h

Description: fft(a) computes $\hat{f}(k) = \sum_{x} a[x] \exp(2\pi i \cdot kx/N)$ for all k. N must be a power of 2. Useful for convolution: conv(a, b) = c, where $c[x] = \sum a[i]b[x-i]$. For convolution of complex numbers or more than two vectors: FFT, multiply pointwise, divide by n, reverse(start+1, end), FFT back. Rounding is safe if $(\sum a_i^2 + \sum b_i^2) \log_2 N < 9 \cdot 10^{14}$ (in practice 10^{16}); higher for random inputs). Otherwise, use NTT/FFTMod. **Time:** $O(N \log N)$ with $N = |A| + |B| (\sim 1s \text{ for } N = 2^{22})$

```
00ced6, 35 lines
typedef complex<double> C:
typedef vector<double> vd;
void fft(vector<C>& a) {
 int n = sz(a), L = 31 - builtin clz(n);
 static vector<complex<long double>> R(2, 1);
 static vector<C> rt(2, 1); // (^ 10% faster if double)
 for (static int k = 2; k < n; k *= 2) {
   R.resize(n); rt.resize(n);
   auto x = polar(1.0L, acos(-1.0L) / k);
   rep(i,k,2*k) rt[i] = R[i] = i&1 ? R[i/2] * x : R[i/2];
 vi rev(n);
 rep(i,0,n) rev[i] = (rev[i / 2] | (i & 1) << L) / 2;
 rep(i,0,n) if (i < rev[i]) swap(a[i], a[rev[i]]);
 for (int k = 1; k < n; k *= 2)
   for (int i = 0; i < n; i += 2 * k) rep(j, 0, k) {
     Cz = rt[j+k] * a[i+j+k]; // (25\% faster if hand-rolled)
     a[i + j + k] = a[i + j] - z;
     a[i + j] += z;
vd conv(const vd& a, const vd& b) {
 if (a.empty() || b.empty()) return {};
 vd res(sz(a) + sz(b) - 1);
 int L = 32 - __builtin_clz(sz(res)), n = 1 << L;</pre>
 vector<C> in(n), out(n);
 copy(all(a), begin(in));
 rep(i,0,sz(b)) in[i].imag(b[i]);
 fft(in);
 for (C& x : in) x *= x;
 rep(i, 0, n) out[i] = in[-i & (n - 1)] - conj(in[i]);
 rep(i, 0, sz(res)) res[i] = imag(out[i]) / (4 * n);
 return res;
```

FastFourierTransformMod.h

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

Time: $\mathcal{O}(N \log N)$, where N = |A| + |B| (twice as slow as NTT or FFT) b82773, 22 lines "FastFourierTransform.h"

```
typedef vector<ll> v1;
template<int M> v1 convMod(const v1 &a, const v1 &b) {
 if (a.empty() || b.empty()) return {};
 vl res(sz(a) + sz(b) - 1);
 int B=32-__builtin_clz(sz(res)), n=1<<B, cut=int(sqrt(M));</pre>
 vector<C> L(n), R(n), outs(n), outl(n);
 rep(i,0,sz(a)) L[i] = C((int)a[i] / cut, (int)a[i] % cut);
 rep(i,0,sz(b)) R[i] = C((int)b[i] / cut, (int)b[i] % cut);
 fft(L), fft(R);
 rep(i,0,n) {
   int j = -i \& (n - 1);
   outl[j] = (L[i] + conj(L[j])) * R[i] / (2.0 * n);
   outs[j] = (L[i] - conj(L[j])) * R[i] / (2.0 * n) / 1i;
 fft (outl), fft (outs);
 rep(i, 0, sz(res)) {
```

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

NumberTheoreticTransform.h

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

```
"../number-theory/ModPow.h"
const 11 mod = (119 << 23) + 1, root = 62; // = 998244353
// For p < 2^30 there is also e.g. 5 << 25, 7 << 26, 479 << 21
// and 483 \ll 21 (same root). The last two are > 10^9.
typedef vector<ll> v1;
void ntt(vl &a) {
  int n = sz(a), L = 31 - __builtin_clz(n);
  static v1 rt(2, 1);
  for (static int k = 2, s = 2; k < n; k *= 2, s++) {
    rt.resize(n);
    11 z[] = {1, modpow(root, mod >> s)};
   rep(i,k,2*k) rt[i] = rt[i / 2] * z[i & 1] % mod;
  vi rev(n);
  rep(i,0,n) \ rev[i] = (rev[i / 2] | (i \& 1) << L) / 2;
  rep(i,0,n) if (i < rev[i]) swap(a[i], a[rev[i]]);
  for (int k = 1; k < n; k *= 2)
    for (int i = 0; i < n; i += 2 * k) rep(j, 0, k) {
     11 z = rt[j + k] * a[i + j + k] % mod, &ai = a[i + j];
      a[i + j + k] = ai - z + (z > ai ? mod : 0);
      ai += (ai + z >= mod ? z - mod : z);
vl conv(const vl &a, const vl &b) {
  if (a.empty() || b.empty()) return {};
  int s = sz(a) + sz(b) - 1, B = 32 - _builtin_clz(s),
     n = 1 << B;
  int inv = modpow(n, mod - 2);
  vl L(a), R(b), out(n);
```

FastSubsetTransform.h

L.resize(n), R.resize(n);

return {out.begin(), out.begin() + s};

ntt(L), ntt(R);

rep(i,0,n)

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

out[-i & (n - 1)] = (l1)L[i] * R[i] % mod * inv % mod;

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

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

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

Number theory (5)

5.1 Modular arithmetic

Modular Arithmetic.h

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

35bfea, 18 lines const 11 mod = 17; // change to something else struct Mod { 11 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, q = euclid(a.x, mod, x, y);assert (g == 1); return Mod((x + mod) % mod); Mod operator^(11 e) { if (!e) return Mod(1); Mod $r = *this ^ (e / 2); r = r * r;$ return e&1 ? *this * r : r; };

ModInverse.h

```
// Method 1
const 11 m = 1000000007, N = 200000;
11 inv[N]; inv[1] = 1;
for (11 i=2; i<N; ++i) inv[i] = m - (m/i) * inv[m%i] % m;

// Method 2
const 11 m = (11)1e9+7;
11 binpow(11 a, 11 b) { // log b
    a %= m; 11 res = 1;
    while (b > 0) {
        if (b & 1) res = res * a % m;
        a = a * a % m; b >>= 1;
    }
    return res;
}
11 mod_inv(11 a) { return binpow(a, m-2); } // 1 / a
vector<11> inv0f = {0, mod_inv(1), mod_inv(2)};
```

ModPow.h

b83e45, 8 lines

```
const 11 mod = 1000000007; // faster if const

11 modpow(11 b, 11 e) {
    11 ans = 1;
    for (; e; b = b * b % mod, e /= 2)
        if (e & 1) ans = ans * b % mod;
    return ans;
}
```

```
ModLog.h
```

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

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

ModSum.h

 $\textbf{Description:} \ \ \text{Sums of mod'ed arithmetic progressions.}$

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

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

5c5bc5, 16 lines

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

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

ll modsum(ull to, ll c, ll k, ll m) {
    c = ((c % m) + m) % m;
    k = ((k % m) + m) % m;
    return to * c + k * sumsq(to) - m * divsum(to, c, k, m);
```

ModMulLL.h

Description: Calculate $a \cdot b \mod c$ (or $a^b \mod c$) for $0 \le a, b \le c \le 7.2 \cdot 10^{18}$. **Time:** $\mathcal{O}(1)$ for modmul, $\mathcal{O}(\log b)$ for modpow bbbd8f, 11 lines

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

ModSqrt.h

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

```
"ModPow.h" 19a793, 24 lines

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

5.2 Primality

FastEratosthenes.h

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

```
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]) {</pre>
    cp.push back(\{i, i * i / 2\});
    for (int j = i * i; j <= S; j += 2 * i) sieve[j] = 1;</pre>
  for (int L = 1; L \le R; L += S) {
    array<bool, S> block{};
    for (auto &[p, idx] : cp)
      for (int i=idx; i < S+L; idx = (i+=p)) block[i-L] = 1;</pre>
    rep(i, 0, min(S, R - L))
      if (!block[i]) pr.push_back((L + i) * 2 + 1);
  for (int i : pr) isPrime[i] = 1;
  return pr;
```

MillerRabin.h

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

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

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

Factor.

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

d8d98d, 18 lines

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

"ModMullL.h", "MillerRabin.h"

euclid CRT phiFunction ContinuedFractions FracBinarySearch

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

5.3 Divisibility

euclid.h

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

```
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;
}
```

CRT.h

Description: Chinese Remainder Theorem.

crt (a, m, b, n) computes x such that $x\equiv a\pmod m$, $x\equiv b\pmod n$. If |a|< m and |b|< n, x will obey $0\le x< \mathrm{lcm}(m,n)$. Assumes $mn<2^{62}$. Time: $\log(n)$

5.3.1 Bézout's identity

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

$$ax + by = d$$

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

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

phiFunction.h

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

```
Fermat's little thm: p prime ⇒ a<sup>p-1</sup> ≡ 1 (mod p) ∀a.
const int LIM = 50000000;
int phi[LIM];

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

5.4 Fractions

ContinuedFractions.h

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

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 the a's eventually become cyclic. Time: $\mathcal{O}(\log N)$

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

FracBinarySearch.h

dir = !dir;

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

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

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

```
swap(lo, hi);
A = B; B = !!adv;
}
return dir ? hi : lo;
```

5.5 Pythagorean Triples

The Pythagorean triples are uniquely generated by

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

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

5.6 Primes

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

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

5.7 Estimates

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

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

5.8 Mobius Function

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

Mobius Inversion:

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

Other useful formulas/forms:

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

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

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

Combinatorial (6)

6.1 Permutations

6.1.1 Factorial

						9		
							3628800	
n	11	12	13	14	15	5 16	17	
n!	4.0e7	4.8e	8 6.2e	9 8.7e	10 1.3e	12 2.1e	13 3.6e14	
n	20	25	30	40	50 1	00 15	0 171	
n!	2e18	2e25	3e32	8e47 3	8e64 9e	$157 \ 6e2$	$62 > DBL_M$	AX

IntPerm multinomial BellmanFord

IntPerm.h

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

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

044568, 6 lines

6.1.2 Cycles

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

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

6.1.3 Derangements

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

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

6.1.4 Burnside's lemma

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

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

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

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

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

6.2 Partitions and subsets

6.2.1 Partition function

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

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

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

6.2.2 Lucas' Theorem

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

6.2.3 Binomials

multinomial.h

6.3 General purpose numbers

6.3.1 Bernoulli numbers

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

Sums of powers:

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

Euler-Maclaurin formula for infinite sums:

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

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

6.3.2 Stirling numbers of the first kind

Number of permutations on n items with k cycles.

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

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

6.3.3 Eulerian numbers

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

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

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

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

6.3.4 Stirling numbers of the second kind

Partitions of n distinct elements into exactly k groups.

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

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

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

6.3.5 Bell numbers

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

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

6.3.6 Labeled unrooted trees

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

6.3.7 Catalan numbers

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

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

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

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

$\underline{\text{Graph}}$ (7)

7.1 Fundamentals

BellmanFord.h

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

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

void bellmanFord(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) {</pre>
```

FloydWarshall TopoSort PushRelabel MinCostMaxFlow

```
Node cur = nodes[ed.a], &dest = nodes[ed.b];
  if (abs(cur.dist) == inf) continue;
  11 d = cur.dist + ed.w;
  if (d < dest.dist) {</pre>
   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;
```

FlovdWarshall.h

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

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

const 11 inf = 1LL << 62;</pre> void floydWarshall(vector<vector<ll>>& m) { int n = sz(m);rep(i, 0, n) m[i][i] = min(m[i][i], OLL);rep(k, 0, n) rep(i, 0, n) rep(j, 0, n)

if (m[i][k] != inf && m[k][j] != inf) { **auto** newDist = max(m[i][k] + m[k][j], -inf);m[i][j] = min(m[i][j], newDist);rep(k, 0, n) **if** (m[k][k] < 0) rep(i, 0, n) rep(j, 0, n)if (m[i][k] != inf && m[k][j] != inf) m[i][j] = -inf;

TopoSort.h

Description: Topological sorting. Given is an oriented graph. Output is an ordering of vertices, such that there are edges only from left to right. If there are cycles, the returned list will have size smaller than n – nodes reachable from cycles will not be returned.

Time: $\mathcal{O}(|V| + |E|)$ 1f04c3, 26 lines vi topoSort(const vector<vi>& gr) {

```
vi indeg(sz(gr)), q;
  for (auto& li : gr) for (int x : li) indeg[x]++;
  rep(i, 0, sz(qr)) if (indeg[i] == 0) q.push_back(i);
  rep(j,0,sz(q)) for (int x : gr[q[j]])
   if (--indeg[x] == 0) g.push back(x);
  return q;
// Method 2 (my method)
// Topological sort in DAG (using DFS)
#define clear_graph(n) for(int i=0;i<=n;i++){g[i].clear(),vis[i]</pre>
     ]=false; } path.clear(); nodes = set<int>();
const int N = 1e5+10;
vector<int> q[N];
bool vis[N];
deque<int> path; // path in assending order
set < int > nodes; // all distinct nodes
void dfs(int vertex) {
  vis[vertex] = true;
  for (int child : g[vertex]) if(!vis[child]) dfs(child);
  path.push_front(vertex);
// run dfs from each node
void TopSort() { for (auto& i : nodes) if (!vis[i]) dfs(i); }
// print(path);
```

7.2 Network flow

PushRelabel.h

531245, 12 lines

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

```
Time: \mathcal{O}\left(V^2\sqrt{E}\right)
struct PushRelabel {
 struct Edge {
   int dest, back;
   11 f, c;
 };
  vector<vector<Edge>> g;
  vector<11> ec:
  vector<Edge*> cur;
  vector<vi> hs; vi H;
  PushRelabel(int n): q(n), ec(n), cur(n), hs(2*n), H(n) {}
  void addEdge(int s, int t, ll cap, ll rcap=0) {
    if (s == t) return;
    g[s].push_back({t, sz(g[t]), 0, cap});
    g[t].push_back({s, sz(g[s])-1, 0, rcap});
  void addFlow(Edge& e, ll f) {
    Edge &back = g[e.dest][e.back];
    if (!ec[e.dest] && f) hs[H[e.dest]].push back(e.dest);
    e.f += f; e.c -= f; ec[e.dest] += f;
    back.f -= f; back.c += f; ec[back.dest] -= f;
  11 calc(int s, int t) {
    int v = sz(q); H[s] = v; ec[t] = 1;
    vi co(2*v); co[0] = v-1;
    rep(i,0,v) cur[i] = q[i].data();
    for (Edge& e : g[s]) addFlow(e, e.c);
    for (int hi = 0;;) {
      while (hs[hi].empty()) if (!hi--) return -ec[s];
      int u = hs[hi].back(); hs[hi].pop_back();
      while (ec[u] > 0) // discharge u
        if (cur[u] == g[u].data() + sz(g[u])) {
          H[u] = 1e9;
          for (Edge& e : g[u]) if (e.c && H[u] > H[e.dest]+1)
            H[u] = H[e.dest]+1, cur[u] = &e;
          if (++co[H[u]], !--co[hi] && hi < v)</pre>
            rep(i, 0, v) if (hi < H[i] && H[i] < v)
              --co[H[i]], H[i] = v + 1;
        } else if (cur[u]->c && H[u] == H[cur[u]->dest]+1)
          addFlow(*cur[u], min(ec[u], cur[u]->c));
        else ++cur[u];
 bool leftOfMinCut(int a) { return H[a] >= sz(g); }
```

MinCostMaxFlow.h

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

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

```
#include <bits/extc++.h>
const 11 INF = numeric_limits<11>::max() / 4;
struct MCMF {
  struct edge {
    int from, to, rev;
```

```
11 cap, cost, flow;
};
int N;
vector<vector<edge>> ed;
vi seen;
vector<ll> dist, pi;
vector<edge*> par;
MCMF(int N) : N(N), ed(N), seen(N), dist(N), pi(N), par(N) {}
void addEdge(int from, int to, ll cap, ll cost) {
  if (from == to) return;
  ed[from].push_back(edge{ from, to, sz(ed[to]), cap, cost, 0 });
  ed[to].push_back(edge{ to,from,sz(ed[from])-1,0,-cost,0 });
void path(int s) {
  fill(all(seen), 0);
  fill(all(dist), INF);
  dist[s] = 0; ll di;
  __gnu_pbds::priority_queue<pair<11, int>> q;
  vector<decltype(q)::point_iterator> its(N);
  q.push({ 0, s });
  while (!q.empty()) {
    s = q.top().second; q.pop();
    seen[s] = 1; di = dist[s] + pi[s];
    for (edge& e : ed[s]) if (!seen[e.to]) {
     11 val = di - pi[e.to] + e.cost;
      if (e.cap - e.flow > 0 && val < dist[e.to]) {</pre>
        dist[e.to] = val;
        par[e.to] = &e;
        if (its[e.to] == q.end())
          its[e.to] = q.push({ -dist[e.to], e.to });
          q.modify(its[e.to], { -dist[e.to], e.to });
  rep(i, 0, N) pi[i] = min(pi[i] + dist[i], INF);
pair<11, 11> maxflow(int s, int t) {
  11 totflow = 0, totcost = 0;
  while (path(s), seen[t]) {
   11 fl = INF;
    for (edge* x = par[t]; x; x = par[x->from])
      fl = min(fl, x->cap - x->flow);
    totflow += fl;
    for (edge* x = par[t]; x; x = par[x->from]) {
      x \rightarrow flow += fl;
      ed[x->to][x->rev].flow -= fl;
  rep(i,0,N) for(edge& e : ed[i]) totcost += e.cost * e.flow;
  return {totflow, totcost/2};
// If some costs can be negative, call this before maxflow:
void setpi(int s) { // (otherwise, leave this out)
  fill(all(pi), INF); pi[s] = 0;
  int it = N, ch = 1; l1 v;
  while (ch-- && it--)
    rep(i,0,N) if (pi[i] != INF)
      for (edge& e : ed[i]) if (e.cap)
        if ((v = pi[i] + e.cost) < pi[e.to])
          pi[e.to] = v, ch = 1;
```

```
assert(it >= 0); // negative cost cycle
};
```

EdmondsKarp.h

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

```
template < class T > T edmonds Karp (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);</pre>
      graph[y][p] += inc;
```

MinCut.h

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

GlobalMinCut.h

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

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

8b0e19, 21 lines

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

```
best = min(best, \{w[t] - mat[t][t], co[t]\});
  co[s].insert(co[s].end(), all(co[t]));
  rep(i,0,n) mat[s][i] += mat[t][i];
  rep(i, 0, n) mat[i][s] = mat[s][i];
  mat[0][t] = INT_MIN;
return best;
```

GomoryHu.h

Description: Given a list of edges representing an undirected flow graph, returns edges of the Gomory-Hu tree. The max flow between any pair of vertices is given by minimum edge weight along the Gomory-Hu tree path. **Time:** $\mathcal{O}(V)$ Flow Computations

```
"PushRelabel.h"
                                                     0418b3, 13 lines
typedef array<11, 3> Edge;
vector<Edge> gomoryHu(int N, vector<Edge> ed) {
 vector<Edge> tree;
 vi par(N);
 rep(i,1,N) {
   PushRelabel D(N); // Dinic also works
   for (Edge t : ed) D.addEdge(t[0], t[1], t[2], t[2]);
   tree.push_back({i, par[i], D.calc(i, par[i])});
   rep(j,i+1,N)
     if (par[j] == par[i] && D.leftOfMinCut(j)) par[j] = i;
 return tree;
```

7.3 Matching

hopcroftKarp.h

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

```
Time: \mathcal{O}\left(\sqrt{V}E\right)
bool dfs(int a, int L, vector<vi>& q, vi& btoa, vi& A, vi& B) {
  if (A[a] != L) return 0;
```

```
A[a] = -1;
 for (int b : q[a]) if (B[b] == L + 1) {
   if (btoa[b] == -1 || dfs(btoa[b], L + 1, q, btoa, A, B))
      return btoa[b] = a, 1;
 return 0;
int hopcroftKarp(vector<vi>& g, vi& btoa) {
 int res = 0;
 vi A(g.size()), B(btoa.size()), cur, next;
 for (;;) {
    fill(all(A), 0);
   fill(all(B), 0);
   cur.clear();
    for (int a : btoa) if (a !=-1) A[a] = -1;
    rep(a, 0, sz(q)) if(A[a] == 0) cur.push_back(a);
    for (int lay = 1;; lay++) {
     bool islast = 0;
     next.clear();
      for (int a : cur) for (int b : g[a]) {
       if (btoa[b] == -1) {
         B[b] = lay;
         islast = 1;
        else if (btoa[b] != a && !B[b]) {
```

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

DFSMatching.h

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

Usage: vi btoa(m, -1); dfsMatching(q, btoa); Time: $\mathcal{O}(VE)$

```
522b98, 22 lines
bool find(int j, vector<vi>& q, vi& btoa, vi& vis) {
 if (btoa[j] == -1) return 1;
 vis[j] = 1; int di = btoa[j];
 for (int e : q[di])
    if (!vis[e] && find(e, g, btoa, vis)) {
     btoa[e] = di;
      return 1:
 return 0;
int dfsMatching(vector<vi>& g, vi& btoa) {
 vi vis;
  rep(i,0,sz(q)) {
    vis.assign(sz(btoa), 0);
    for (int j : q[i])
      if (find(j, g, btoa, vis)) {
        btoa[j] = i;
        break:
  return sz(btoa) - (int)count(all(btoa), -1);
```

MinimumVertexCover.h

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

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

WeightedMatching GeneralMatching BFS DFS

```
WeightedMatching.h
```

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

```
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;</pre>
       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
```

General Matching.h

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

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

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

7.4 DFS algorithms

BFS.h

```
e080d1, 66 lines
// BFS
const int N = 2e5+25;
vector<int> q[N];
vector<bool> vis;
void clear_graph(int n) {
    for (auto& v : g) v.clear();
    vis.assign(n, false);
void bfs(int vertex) {
    queue<int> q;
    q.push (vertex);
    vis[vertex] = true;
    while (!q.empty()) {
        int v = q.front();
        q.pop();
        for (auto& child : g[v]) {
            if (!vis[child]) {
                q.push(child);
                vis[child] = true;
    }
// Grid BFS
             U D R L 1 2 3 4
int dx[] = \{-1, 1, 0, 0, -1, -1, 1\};
int dy[] = \{ 0, 0, 1, -1, 1, -1, -1, 1\};
int n, m;
vector<vector<int>> grid, vis;
bool check(int x, int y) {
    return (x >= 0 \text{ and } x < n) and (y >= 0 \text{ and } y < m);
void bfs(int a, int b) {
    queue<pair<int, int>> q;
    vis[a][b] = 0;
    q.push({a, b});
    while (!q.empty()) {
        int x = q.front().first;
        int y = q.front().second;
```

```
q.pop();
        for (int i = 0; i < 4; i++) {</pre>
            int xx = x + dx[i], yy = y + dy[i];
            if (!check(xx, yy)) continue;
            if (grid[xx][yy] == 0 and vis[x][y] + 1 < vis[xx][
                vis[xx][yy] = vis[x][y] + 1;
                q.push({xx, yy});
// prepare grid and vis
// cin >> n >> m;
// grid.assign(n, vector < int > (m, 0));
// vis.assign(n, vector < int > (m, (int) 1e9));
DFS.h
Description: DFS
                                                      eb1ad9, 53 lines
// DFS
const int N = 2e5+25;
vector<int> g[N];
vector<bool> vis;
void clear_graph(int n) {
    for (auto& v : g) v.clear();
    vis.assign(n, false);
int dfs(int vertex) {
    vis[vertex] = true;
    for (int child : g[vertex]) {
        if (!vis[child]) {
            dfs(child);
big int
// Grid DFS
             U D R L 1 2 3 4
int dx[] = \{-1, 1, 0, 0, -1, -1, 1\};
int dy[] = \{ 0, 0, 1, -1, 1, -1, -1, 1\};
int n, m;
vector<vector<int>> grid, vis;
bool check(int x, int y) {
    return (x \ge 0 \text{ and } x < n) and (y \ge 0 \text{ and } y < m);
int dfs(int x, int y) {
    if (grid[x][y] == 0) return 0;
    vis[x][y] = 1;
    int count = 1;
    for (int i = 0; i < 4; i++) {
        int xx = x + dx[i], yy = y + dy[i];
```

if (!vis[xx][yy])

count += dfs(xx, yy);

```
return count;
// prepare grid and vis
// cin >> n >> m:
// grid.assign(n, vector < int > (m, 0));
// vis.assign(n, vector < int > (m, 0));
```

SCC.h

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

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

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

BiconnectedComponents.h

Description: Finds all biconnected components in an undirected graph, and runs a callback for the edges in each. In a biconnected component there are at least two distinct paths between any two nodes. Note that a node can be in several components. An edge which is not in a component is a bridge, i.e., not part of any cycle.

```
Usage: int eid = 0; ed.resize(N);
for each edge (a,b) {
ed[a].emplace_back(b, eid);
ed[b].emplace_back(a, eid++); }
bicomps([&](const vi& edgelist) {...});
Time: \mathcal{O}(E+V)
```

c6b7c7, 32 lines

```
vi num, st;
vector<vector<pii>> ed;
int Time;
template < class F>
int dfs(int at, int par, F& f) {
  int me = num[at] = ++Time, top = me;
  for (auto [y, e] : ed[at]) if (e != par) {
    if (num[v]) {
      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);</pre>
     else { /* e is a bridge */ }
 return top;
template<class F>
void bicomps (F f) {
 num.assign(sz(ed), 0);
 rep(i,0,sz(ed)) if (!num[i]) dfs(i, -1, f);
```

2sat.h

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

```
Usage: TwoSat ts(number of boolean variables);
ts.either(0, \sim3); // Var 0 is true or var 3 is false
ts.setValue(2); // Var 2 is true
ts.atMostOne(\{0, \sim 1, 2\}); // <= 1 of vars 0, \sim 1 and 2 are true
ts.solve(); // Returns true iff it is solvable
```

ts.values[0..N-1] holds the assigned values to the vars **Time:** $\mathcal{O}(N+E)$, where N is the number of boolean variables, and E is the number of clauses.

```
struct TwoSat {
```

```
int N;
vector<vi> gr;
vi values; // 0 = false, 1 = true
TwoSat(int n = 0) : N(n), gr(2*n) {}
int addVar() { // (optional)
  gr.emplace_back();
  gr.emplace_back();
  return N++;
void either(int f, int j) {
  f = \max(2 * f, -1 - 2 * f);
  j = \max(2*j, -1-2*j);
  gr[f].push_back(j^1);
  gr[j].push_back(f^1);
void setValue(int x) { either(x, x); }
void atMostOne(const vi& li) { // (optional)
  if (sz(li) <= 1) return;</pre>
  int cur = \simli[0];
  rep(i,2,sz(li)) {
    int next = addVar();
    either(cur, ~li[i]);
    either(cur, next);
    either(~li[i], next);
    cur = ~next;
  either(cur, ~li[1]);
```

vi val, comp, z; int time = 0;

```
int dfs(int i) {
    int low = val[i] = ++time, x; z.push_back(i);
    for(int e : gr[i]) if (!comp[e])
     low = min(low, val[e] ?: dfs(e));
    if (low == val[i]) do {
     x = z.back(); z.pop_back();
      comp[x] = low;
      if (values[x>>1] == -1)
       values[x>>1] = x&1;
    } while (x != i);
    return val[i] = low;
 bool solve() {
    values.assign(N, -1);
    val.assign(2*N, 0); comp = val;
    rep(i,0,2*N) if (!comp[i]) dfs(i);
    rep(i,0,N) if (comp[2*i] == comp[2*i+1]) return 0;
    return 1:
};
```

EulerWalk.h

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

```
780b64, 15 lines
vi eulerWalk (vector<vector<pii>>& gr, int nedges, int src=0) {
 int n = sz(qr);
 vi D(n), its(n), eu(nedges), ret, s = {src};
 D[src]++; // to allow Euler paths, not just cycles
  while (!s.emptv()) {
   int x = s.back(), y, e, &it = its[x], end = sz(gr[x]);
    if (it == end) { ret.push_back(x); s.pop_back(); continue; }
    tie(y, e) = qr[x][it++];
    if (!eu[e]) {
     D[x]--, D[y]++;
      eu[e] = 1; s.push_back(y);
 for (int x : D) if (x < 0 \mid | sz(ret) != nedges+1) return \{\};
  return {ret.rbegin(), ret.rend()};
```

7.5 Coloring

EdgeColoring.h

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

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

```
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]);
```

9775a0, 21 lines

```
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;
```

7.6 Heuristics

MaximalCliques.h

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

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

b0d5b1, 12 lines

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

MaximumClique.h

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

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

```
f7c0bc, 49 lines
typedef vector<bitset<200>> vb;
struct Maxclique {
  double limit=0.025, pk=0;
  struct Vertex { int i, d=0; };
  typedef vector<Vertex> vv;
  vb e;
  vv V;
  vector<vi> C;
  vi qmax, q, S, old;
  void init(vv& r) {
    for (auto \& v : r) v.d = 0;
   for (auto& v : r) for (auto j : r) v.d += e[v.i][j.i];
   sort(all(r), [](auto a, auto b) { return a.d > b.d; });
   int mxD = r[0].d;
   rep(i, 0, sz(r)) r[i].d = min(i, mxD) + 1;
  void expand(vv& R, int lev = 1) {
   S[lev] += S[lev - 1] - old[lev];
   old[lev] = S[lev - 1];
   while (sz(R)) {
     if (sz(q) + R.back().d <= sz(qmax)) return;</pre>
     q.push_back(R.back().i);
      for(auto v:R) if (e[R.back().i][v.i]) T.push_back({v.i});
```

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

MaximumIndependentSet.h

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

7.7 Trees

BinaryLifting.h

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

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

bfce85, 25 lines

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

LCA.h

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

```
Time: \mathcal{O}(N \log N + Q)
"../data-structures/RMQ.h"
                                                        0f62fb, 21 lines
struct LCA {
 int T = 0;
 vi time, path, ret;
  RMO<int> rmg;
  LCA(vector < vi > \& C) : time(sz(C)), rmq((dfs(C,0,-1), ret)) {}
  void dfs(vector<vi>& C, int v, int par) {
    time[v] = T++;
    for (int y : C[v]) if (y != par) {
      path.push_back(v), ret.push_back(time[v]);
      dfs(C, y, v);
 int lca(int a, int b) {
    if (a == b) return a;
    tie(a, b) = minmax(time[a], time[b]);
    return path[rmq.query(a, b)];
  //dist(a,b){return depth[a] + depth[b] - 2*depth[lca(a,b)];}
```

CompressTree.h

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

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

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

HLD I

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

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

```
"../data-structures/LazySegmentTree.h" 03139d, 46 lines

template <bool VALS_EDGES> struct HLD {
   int N, tim = 0;
   vector<vi> adj;
   vi par, siz, rt, pos;
   Node *tree;
```

```
HLD(vector<vi> adj )
    : N(sz(adj_)), adj(adj_), par(N, -1), siz(N, 1),
      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;
     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 (pos[rt[u]] > pos[rt[v]]) swap(u, v);
     op(pos[rt[v]], pos[v] + 1);
    if (pos[u] > pos[v]) swap(u, v);
   op(pos[u] + VALS_EDGES, pos[v] + 1);
  void modifyPath(int u, int v, int val) {
   process(u, v, [&] (int 1, int r) { tree->add(1, r, val); });
  int queryPath(int u, int v) { // Modify depending on problem
   int res = -1e9;
   process(u, v, [&](int 1, int r) {
       res = max(res, tree->query(1, r));
   });
    return res;
  int querySubtree(int v) { // modifySubtree is similar
    return tree->query(pos[v] + VALS EDGES, pos[v] + siz[v]);
};
```

LinkCutTree.h

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

```
Time: All operations take amortized \mathcal{O}(\log N).
                                                      0fb462, 90 lines
struct Node { // Splay tree. Root's pp contains tree's parent.
  Node *p = 0, *pp = 0, *c[2];
 bool flip = 0;
 Node() { c[0] = c[1] = 0; fix(); }
  void fix() {
   if (c[0]) c[0]->p = this;
   if (c[1]) c[1]->p = this;
   // (+ update sum of subtree elements etc. if wanted)
  void pushFlip() {
   if (!flip) return;
   flip = 0; swap(c[0], c[1]);
   if (c[0]) c[0]->flip ^= 1;
   if (c[1]) c[1]->flip ^= 1;
  int up() { return p ? p->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];
     y - > c[h ^ 1] = x;
    z->c[i ^ 1] = this;
    fix(); x->fix(); y->fix();
    if (p) p->fix();
    swap(pp, y->pp);
 void splay() {
    for (pushFlip(); p; ) {
     if (p->p) p->p->pushFlip();
      p->pushFlip(); pushFlip();
      int c1 = up(), c2 = p->up();
      if (c2 == -1) p->rot(c1, 2);
      else p->p->rot(c2, c1 != c2);
 Node* first() {
   pushFlip();
    return c[0] ? c[0]->first() : (splay(), this);
};
struct LinkCut {
 vector<Node> node;
 LinkCut(int N) : node(N) {}
 void link(int u, int v) { // add an edge (u, v)
    assert(!connected(u, v));
    makeRoot (&node[u]);
    node[u].pp = &node[v];
 void cut (int u, int v) { // remove an edge (u, v)
    Node *x = &node[u], *top = &node[v];
    makeRoot(top); x->splay();
    assert(top == (x->pp ?: x->c[0]));
    if (x->pp) x->pp = 0;
     x->c[0] = top->p = 0;
      x->fix();
 bool connected (int u, int v) { // are u, v in the same tree?
   Node* nu = access(&node[u])->first();
    return nu == access(&node[v])->first();
 void makeRoot(Node* u) {
    access(u);
    u->splav();
    if(u->c[0]) {
     u - c[0] - p = 0;
     u - c[0] - flip ^= 1;
     u - c[0] - pp = u;
     u - > c[0] = 0;
     u \rightarrow fix();
 Node* access(Node* u) {
   u->splay();
    while (Node* pp = u->pp) {
      pp->splay(); u->pp = 0;
      if (pp->c[1]) {
        pp \rightarrow c[1] \rightarrow p = 0; pp \rightarrow c[1] \rightarrow pp = pp; }
      pp->c[1] = u; pp->fix(); u = pp;
    return u:
};
```

DirectedMST.h

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

19

```
Time: \mathcal{O}\left(E\log V\right)
"../data-structures/UnionFindRollback.h"
                                                      39e620, 60 lines
struct Edge { int a, b; ll w; };
struct Node {
 Edge key;
  Node *1, *r;
  11 delta;
  void prop() {
    key.w += delta;
    if (1) 1->delta += delta;
    if (r) r->delta += delta;
    delta = 0:
  Edge top() { prop(); return key; }
Node *merge(Node *a, Node *b) {
  if (!a || !b) return a ?: b;
  a->prop(), b->prop();
  if (a->key.w > b->key.w) swap(a, b);
  swap(a->1, (a->r = merge(b, a->r)));
  return a;
void pop(Node*\& a) { a->prop(); a = merge(a->1, a->r); }
pair<11, vi> dmst(int n, int r, vector<Edge>& g) {
  RollbackUF uf(n);
  vector<Node*> heap(n);
  for (Edge e : q) heap[e.b] = merge(heap[e.b], new Node{e});
  11 \text{ res} = 0;
  vi seen(n, -1), path(n), par(n);
  seen[r] = r;
  vector<Edge> Q(n), in(n, \{-1,-1\}), comp;
  deque<tuple<int, int, vector<Edge>>> cycs;
  rep(s,0,n) {
    int u = s, qi = 0, w;
    while (seen[u] < 0) {</pre>
      if (!heap[u]) return {-1,{}};
      Edge e = heap[u]->top();
      heap[u]->delta -= e.w, pop(heap[u]);
      O[qi] = e, path[qi++] = u, seen[u] = s;
      res += e.w, u = uf.find(e.a);
      if (seen[u] == s) {
        Node * cvc = 0;
        int end = qi, time = uf.time();
        do cyc = merge(cyc, heap[w = path[--qi]]);
        while (uf.join(u, w));
        u = uf.find(u), heap[u] = cyc, seen[u] = -1;
        cycs.push_front({u, time, {&Q[qi], &Q[end]}});
    rep(i, 0, qi) in[uf.find(Q[i].b)] = Q[i];
  for (auto& [u,t,comp] : cycs) { // restore sol (optional)
    uf.rollback(t);
    Edge inEdge = in[u];
    for (auto& e : comp) in[uf.find(e.b)] = e;
    in[uf.find(inEdge.b)] = inEdge;
  rep(i,0,n) par[i] = in[i].a;
  return {res, par};
```

Math

7.8.1 Number of Spanning Trees

Create an $N \times N$ matrix mat, and for each edge $a \to b \in G$, do mat[a][b]--, mat[b][b]++ (and mat[b][a]--, mat[a][a]++ if G is undirected). Remove the *i*th row and column and take the determinant; this yields the number of directed spanning trees rooted at i (if G is undirected, remove any row/column). A simple graph with node degrees $d_1 \ge \cdots \ge d_n$ exists iff $d_1 + \cdots + d_n$ is even and for every $k = 1 \dots n$,

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

Geometry (8)

8.1 Geometric primitives

Point.h

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

```
template \langle class T \rangle int sgn(T x) \{ return (x > 0) - (x < 0); \}
template<class T>
struct Point {
  typedef Point P;
  explicit Point (T x=0, T y=0) : x(x), y(y) {}
  bool operator<(P p) const { return tie(x,y) < tie(p.x,p.y);</pre>
  bool operator==(P p) const { return tie(x,y)==tie(p.x,p.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) {</pre>
    return os << "(" << p.x << "," << p.y << ")"; }
};
```

lineDistance.h

Description:

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

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

SegmentDistance.h

Description:

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

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

5c88f4, 6 lines typedef Point < double > P;

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

SegmentIntersection.h

Description:

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

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

```
"Point.h", "OnSegment.h"
                                                      9d57f2, 13 lines
template < class P > vector < P > segInter (P a, P b, P c, P d) {
 auto oa = c.cross(d, a), ob = c.cross(d, b),
       oc = a.cross(b, c), od = a.cross(b, d);
  // Checks if intersection is single non-endpoint point.
 if (sqn(oa) * sqn(ob) < 0 && sqn(oc) * sqn(od) < 0)
   return { (a * ob - b * oa) / (ob - oa) };
 set<P> s;
 if (onSegment(c, d, a)) s.insert(a);
 if (onSegment(c, d, b)) s.insert(b);
 if (onSegment(a, b, c)) s.insert(c);
 if (onSegment(a, b, d)) s.insert(d);
 return {all(s)};
```

lineIntersection.h

Description:

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



```
Usage: auto res = lineInter(s1,e1,s2,e2);
            if (res.first == 1)
            cout << "intersection point at " << res.second << endl;</pre>
             "Point.h"
                                                                   a01f81, 8 lines
            template<class P>
            pair<int, P> lineInter(P s1, P e1, P s2, P e2) {
              auto d = (e1 - s1).cross(e2 - s2);
              if (d == 0) // if parallel
                return {-(s1.cross(e1, s2) == 0), P(0, 0)};
f6bf6b, 4 lines
              auto p = s2.cross(e1, e2), q = s2.cross(e2, s1);
              return {1, (s1 * p + e1 * q) / d};
```

sideOf.h

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

```
Usage: bool left = sideOf(p1,p2,q)==1;
"Point.h"
                                                            3af81c, 9 lines
```

```
template<class P>
int sideOf(P s, P e, P p) { return sqn(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 1 = (e-s).dist()*eps;
 return (a > 1) - (a < -1);
```

OnSegment.h

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

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

linearTransformation.h Description:

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

```
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();
```

Angle.h

"Point.h"

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

```
Usage: vector<Angle> v = \{w[0], w[0].t360() ...\}; // sorted
int j = 0; rep(i,0,n) { while (v[j] < v[i].t180()) ++j; }
// sweeps j such that (j-i) represents the number of positively
oriented triangles with vertices at 0 and i
                                                     0f0602, 35 lines
```

```
struct Angle {
  int x, y;
  Angle(int x, int y, int t=0) : x(x), y(y), t(t) {}
  Angle operator-(Angle b) const { return {x-b.x, y-b.y, t}; }
 int half() const {
    assert(x || y);
```

```
return v < 0 || (v == 0 && x < 0);
  Angle t90() const { return \{-y, x, t + (half() \&\& x >= 0)\}; \}
  Angle t180() const { return {-x, -y, t + half()}; }
 Angle t360() const { return {x, y, t + 1}; }
bool operator<(Angle a, Angle b) {</pre>
  // add a.dist2() and b.dist2() to also compare distances
  return make_tuple(a.t, a.half(), a.y * (ll)b.x) <</pre>
         make_tuple(b.t, b.half(), a.x * (ll)b.y);
// Given two points, this calculates the smallest angle between
// them, i.e., the angle that covers the defined line segment.
pair<Angle, Angle> segmentAngles(Angle a, Angle b) {
  if (b < a) swap(a, b);
  return (b < a.t180() ?
          make_pair(a, b) : make_pair(b, a.t360()));
Angle operator+(Angle a, Angle b) { // point \ a + vector \ b
  Angle r(a.x + b.x, a.y + b.y, a.t);
  if (a.t180() < r) r.t--;</pre>
  return r.t180() < a ? r.t360() : r;
Angle angleDiff(Angle a, Angle b) { // angle b- angle a
  int tu = b.t - a.t; a.t = b.t;
  return \{a.x*b.x + a.y*b.y, a.x*b.y - a.y*b.x, tu - (b < a)\};
```

8.2 Circles

CircleIntersection.h

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

```
84d6d3, 11 lines
typedef Point < double > P;
bool circleInter(P a, P b, double r1, double r2, pair < P, P > * out) {
  if (a == b) { assert(r1 != r2); return false; }
  P \text{ 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 + \text{vec*p}, per = \text{vec.perp}() * \text{sgrt}(\text{fmax}(0, h2) / d2);
  *out = {mid + per, mid - per};
  return true;
```

CircleTangents.h

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

```
"Point.h"
                                                      b0153d, 13 lines
template<class P>
vector<pair<P, P>> tangents(P c1, double r1, P c2, double r2) {
 P d = c2 - c1;
 double dr = r1 - r2, d2 = d.dist2(), h2 = d2 - dr * dr;
 if (d2 == 0 || h2 < 0) return {};</pre>
  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;
```

CirclePolygonIntersection.h

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

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

```
"../../content/geometry/Point.h"
                                                       a1ee63, 19 lines
typedef Point < double > P;
#define arg(p, q) atan2(p.cross(q), p.dot(q))
double circlePoly(P c, double r, vector<P> ps) {
 auto tri = [&] (P p, P q) {
   auto r2 = r * r / 2;
   P d = q - p;
    auto a = d.dot(p)/d.dist2(), b = (p.dist2()-r*r)/d.dist2();
    auto det = a * a - b;
    if (det <= 0) return arg(p, q) * r2;</pre>
    auto s = max(0., -a-sqrt(det)), t = min(1., -a+sqrt(det));
    if (t < 0 || 1 <= s) return arg(p, q) * r2;</pre>
   Pu = p + d * s, v = p + d * t;
    return arg(p,u) * r2 + u.cross(v)/2 + arg(v,g) * r2;
  auto sum = 0.0;
  rep(i, 0, sz(ps))
   sum += tri(ps[i] - c, ps[(i + 1) % sz(ps)] - c);
  return sum;
```

circumcircle.h

Description:

"Point.h"

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



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

MinimumEnclosingCircle.h

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

```
"circumcircle.h"
                                                     09dd0a, 17 lines
pair<P, double> mec(vector<P> ps) {
 shuffle(all(ps), mt19937(time(0)));
 P \circ = ps[0];
 double r = 0, EPS = 1 + 1e-8;
 rep(i, 0, sz(ps)) if ((o - ps[i]).dist() > r * EPS) {
   o = ps[i], r = 0;
   rep(j, 0, i) if ((o - ps[j]).dist() > r * EPS) {
     o = (ps[i] + ps[j]) / 2;
     r = (o - ps[i]).dist();
     rep(k, 0, j) if ((o - ps[k]).dist() > r * EPS) {
       o = ccCenter(ps[i], ps[j], ps[k]);
        r = (o - ps[i]).dist();
 return {o, r};
```

8.3 Polygons

InsidePolygon.h

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

```
Usage: vectorP> v = \{P\{4,4\}, P\{1,2\}, P\{2,1\}\};
bool in = inPolygon(v, P{3, 3}, false);
Time: \mathcal{O}(n)
"Point.h", "OnSegment.h", "SegmentDistance.h"
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) \le eps) return !strict;
   cnt ^= ((a.y<p[i].y) - (a.y<q.y)) * a.cross(p[i], q) > 0;
 return cnt;
```

PolygonArea.h

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

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

PolygonCenter.h

Description: Returns the center of mass for a polygon.

Time: $\mathcal{O}\left(n\right)$ "Point.h"

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

PolygonCut.h

Description:

thing to the left of the line going from s to e cut away.

```
Returns a vector with the vertices of a polygon with every-
Usage: vector<P> p = ...;
p = polygonCut(p, P(0,0), P(1,0));
"Point.h", "lineIntersection.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;</pre>
    if (side != (s.cross(e, prev) < 0))</pre>
      res.push_back(lineInter(s, e, cur, prev).second);
    if (side)
      res.push_back(cur);
 return res;
```

ConvexHull.h

Description:

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



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

HullDiameter.h

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

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

PointInsideHull.h

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

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

```
"Point.h", "sideOf.h", "OnSegment.h" 71446b, 14 lines
typedef Point<11> P;

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

LineHullIntersection.h

Description: Line-convex polygon intersection. The polygon must be ccw and have no collinear points. lineHull(line, poly) returns a pair describing the intersection of a line with the polygon: \bullet (-1,-1) if no collision, \bullet (i,-1)if touching the corner $i, \bullet (i, i)$ if along side $(i, i+1), \bullet (i, j)$ if crossing sides (i, i+1) and (j, j+1). In the last case, if a corner i is crossed, this is treated as happening on side (i, i + 1). The points are returned in the same order as the line hits the polygon. extrVertex returns the point of a hull with the max projection onto a line. Time: $\mathcal{O}(\log n)$ "Point.h" 7cf45b, 39 lines #define cmp(i,j) sgn(dir.perp().cross(poly[(i)%n]-poly[(j)%n])) #define extr(i) cmp(i + 1, i) >= 0 && cmp(i, i - 1 + n) < 0template <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 = (10 + hi) / 2;if (extr(m)) return m; **int** 1s = cmp(1o + 1, 1o), ms = cmp(m + 1, m); $(ls < ms \mid | (ls == ms \&\& ls == cmp(lo, m)) ? hi : lo) = m;$

```
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;</pre>
      (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;
```

Strings (9)

KMP.h

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

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

```
// kmp
// kmp
// Time Complexity: O(n)
vector<int> compute_prefix_function(const string& s) {
    vector<int> prefix_function(s.size());
    for (int i = 1; i < s.size(); i++) {
        int g = prefix_function[i - 1];
        while (g && s[i] != s[g]) g = prefix_function[g - 1];
        prefix_function[i] = g + (s[i] == s[g]);
    }
    return prefix_function;</pre>
```

Zfunc.h

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

```
0bf208, 35 lines
// Z-algorithm to compute the Z-array
vector<int> Z(const string& S) {
 int n = S.size();
 vector<int> z(n);
 int 1 = -1, r = -1;
  // Loop to compute the Z-array in linear time
  for (int i = 1; i < n; ++i) {
    z[i] = (i >= r) ? 0 : min(r - i, z[i - 1]);
    while (i + z[i] < n \&\& S[i + z[i]] == S[z[i]]) z[i]++;
    if (i + z[i] > r) \{ l = i; r = i + z[i]; \}
 return z;
// Time complexity of Z function: O(n), where n is the length
     of string S
vector<int> pattern_matching(const string& pattern, const
    string& text) {
 string combined = pattern + "#" + text; // Concatenate
       pattern, delimiter, and text
  vector<int> z = Z(combined); // Z-function on combined string
  vector<int> occurrences;
 int p_len = pattern.size();
  // Loop to find all occurrences of the pattern in the text
  for (int i = p_len + 1; i < combined.size(); ++i) {</pre>
    if (z[i] >= p_len) occurrences.push_back(i - p_len - 1);
  return occurrences;
// Time complexity of pattern_matching function:
// The Z-function runs in, where p_len is the length of the
```

Manacher.h

 $p_len + t_len)$ time.

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

// Iterating through the Z-array to find occurrences takes O(

pattern and t_len is the length of the text.

// Overall time complexity: $O(p_{-len} + t_{-len})$

vector<int> occ = pattern_matching(pattern, text);

MinRotation SuffixArray SuffixTree Hashing

```
Time: \mathcal{O}(N)
                                                     168d15, 33 lines
struct Manacher {
  array<vector<int>, 2> p;
  Manacher (const string& s) {
   int n = s.size();
    p = {vector<int>(n+1), vector<int>(n)};
    for (int z = 0; z < 2; z++) {
     int 1 = 0, r = 0;
     for (int i = 0; i < n; i++) {</pre>
       // use previous value if mirror value already
             calculated
        if (i < r) {
          int dist from r = r - i + !z;
          int mirror_idx = 1 + dist_from_r;
          p[z][i] = min(dist from r, p[z][mirror idx]);
        // expand palindrome around i
        int L = i - p[z][i], R = i + p[z][i] - !z;
       while (L > 0 and R < n-1 and s[L-1] != s[R+1]) { p[z][i]
            ]++; L--; R++; }
       if (R > r) \{ l = L; r = R; \}
 bool isPalindrome(int 1, int r) {
   int len = r - 1 + 1;
   int mid idx = (1 + r + 1) / 2;
   int radius = p[(len & 1)][mid_idx];
   return (radius * 2 + 1 >= len);
// Manacher mana(s);
// radius of palindrome at index i is p[i]
// print(mana.p[0]); // even len palindromes
// print(mana.p[1]);// odd len palindromes
```

MinRotation.h

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

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

SuffixArrav.h

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

```
384816, 45 lines
// #pragma once
struct SuffixArray {
  vector<int> sa, lcp;
  SuffixArray(string& s, int lim = 256) {
    int n = s.size() + 1, k = 0, a, b;
   vector<int> x(s.begin(), s.end()), y(n), ws(max(n, lim));
   x.push_back(0), sa = lcp = y;
   iota(sa.begin(), sa.end(), 0);
```

```
// Build suffix array using doubling approach
    for (int j = 0, p = 0; p < n; j = max(1, j * 2), lim = p) {
      p = j;
      iota(y.begin(), y.end(), n - j); // Initialize y with
           indices from n-j to n-1
      for (int i = 0; i < n; i++) if (sa[i] >= j) y[p++] = sa[i]
      fill(ws.begin(), ws.end(), 0); // Reset counting array
      for (int i = 0; i < n; i++) ws[x[i]]++; // Count</pre>
           occurrences of ranks
      for (int i = 1; i < lim; i++) ws[i] += ws[i - 1]; //</pre>
           Convert counts to positions
      for (int i = n; i--;) sa[--ws[x[y[i]]]] = y[i]; //
           Sorting suffixes based on 1st part
      swap(x, y);
      p = 1, x[sa[0]] = 0;
      for (int i = 1; i < n; i++) {</pre>
       a = sa[i - 1], b = sa[i];
       x[b] = (y[a] == y[b] && y[a + j] == y[b + j]) ? p - 1 :
              p++; // Compare suffixes
    }
    // Compute LCP array
    for (int i = 0, j; i < n - 1; lcp[x[i++]] = k)
     for (k \& \& k--, j = sa[x[i] - 1]; s[i + k] == s[j + k]; k
           ++);
};
void printSA(SuffixArray& sufa, string& s) {
 auto& lcp = sufa.lcp, sa = sufa.sa;
 for (int i = 1; i <= s.size(); i++)</pre>
    cout << lcp[i] << ' ' << sa[i] << ' ' << s.substr(sa[i]) <<
 cout << endl;
// // Create a SuffixArray object
// SuffixArray sufa(s);
// sufa.sa; // Suffix array 1 based
// sufa.lcp; // LCP array 1 based
// printSA(sufa, s); // prints SA, LCP, and substrings
```

SuffixTree.h

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

aae0b8, 50 lines

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

p[m]=p[v]; t[m][c]=m+1; t[m][toi(a[q])]=v;

l[v]=q; p[v]=m; t[p[m]][toi(a[l[m]])]=m;

```
v=s[p[m]]; q=l[m];
      while (q<r[m]) { v=t[v][toi(a[q])]; q+=r[v]-l[v]; }</pre>
      if (q==r[m]) s[m]=v; else s[m]=m+2;
      q=r[v]-(q-r[m]); m+=2; goto suff;
  SuffixTree(string a) : a(a) {
    fill(r,r+N,sz(a));
    memset(s, 0, sizeof s);
    memset(t, -1, sizeof t);
    fill(t[1],t[1]+ALPHA,0);
    s[0] = 1; 1[0] = 1[1] = -1; r[0] = r[1] = p[0] = p[1] = 0;
    rep(i,0,sz(a)) ukkadd(i, toi(a[i]));
  // example: find longest common substring (uses ALPHA = 28)
  pii best;
  int lcs(int node, int i1, int i2, int olen) {
    if (1[node] <= i1 && i1 < r[node]) return 1;</pre>
    if (1[node] <= i2 && i2 < r[node]) return 2;</pre>
    int mask = 0, len = node ? olen + (r[node] - 1[node]) : 0;
    rep(c, 0, ALPHA) if (t[node][c] != -1)
     mask |= lcs(t[node][c], i1, i2, len);
    if (mask == 3)
     best = max(best, {len, r[node] - len});
    return mask;
  static pii LCS(string s, string t) {
    SuffixTree st(s + (char) ('z' + 1) + t + (char) ('z' + 2));
    st.lcs(0, sz(s), sz(s) + 1 + sz(t), 0);
    return st.best;
Hashing.h
Description: Self-explanatory methods for string hashing.
class HashedString {
```

```
private:
// change M and B if you want
static const long long M = 1e9 + 9;
static const long long B = 9973;
// pow[i] contains B^i % M
static std::vector<long long> pow;
// p_hash[i] is the hash of the first i characters of the
     given string
std::vector<long long> p_hash;
std::vector<long long> s_hash;
public:
HashedString(const std::string &s) : p_hash(s.size() + 1) {
  // Ensure pow has enough elements to handle the string size
  while (pow.size() < s.size()) {</pre>
    pow.push_back((pow.back() * B) % M);
  // Compute forward hash
  p_hash[0] = 0;
  for (int i = 0; i < s.size(); i++) {</pre>
    p_hash[i + 1] = ((p_hash[i] * B) % M + s[i]) % M;
  // Compute backward hash
  s hash.resize(s.size() + 1);
  s_hash[s.size()] = 0;
```

for (int i = s.size() - 1; i >= 0; --i) {

AhoCorasick.h

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

Time: construction takes $\mathcal{O}(26N)$, where N= sum of length of patterns. find(x) is $\mathcal{O}(N)$, where N= length of x. findAll is $\mathcal{O}(NM)$. a91b91, 87 lines

```
struct AhoCorasick {
  enum {alpha = 26, first = 'A'}; // change this!
  struct Node {
   int back, next[alpha], start = -1, end = -1, nmatches = 0;
   Node(int v) { memset(next, v, sizeof(next)); }
  vector<Node> N:
  vector<int> backp;
  void insert(string& s, int j) {
    assert(!s.empty());
   int n = 0:
   for (char c : s) {
     int& m = N[n].next[c - first];
     if (m == -1) {
       n = m = static_cast<int>(N.size());
       N.emplace_back(-1);
     } else n = m;
   if (N[n].end == -1) N[n].start = j;
   backp.push_back(N[n].end);
   N[n].end = j; N[n].nmatches++;
  AhoCorasick(vector<string>& pat) : N(1, -1) {
    for (size_t i = 0; i < pat.size(); ++i)</pre>
     insert(pat[i], static_cast<int>(i));
   N[0].back = static_cast<int>(N.size());
   N.emplace_back(0);
    queue<int> q;
```

```
for (q.push(0); !q.empty(); q.pop()) {
      int n = q.front(), prev = N[n].back;
      for (int i = 0; i < alpha; ++i) {</pre>
        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);
  vector<int> find(string& word) {
    int n = 0;
    vector<int> res;
    for (char c : word) {
     n = N[n].next[c - first];
     res.push back(N[n].end);
    return res;
  vector<vector<int>> findAll(vector<string>& pat, string& word
    vector<int> r = find(word);
    vector<vector<int>> res(word.size());
    for (size_t i = 0; i < word.size(); ++i) {</pre>
     int ind = r[i];
      while (ind !=-1) {
       res[i - static_cast<int>(pat[ind].size()) + 1].
            push_back(ind);
        ind = backp[ind];
    return res;
};
vector<string> patterns;
AhoCorasick ac(patterns); // O (total length of all patterns)
ac.insert(newPattern, indexObasedPostion); // O(len(newPattern)
// // Find longest pattern that ends at each position in text
// // -1 if no pattern ends there, O(len(text))
vector<int> matches = ac.find(text);
// // Find all patterns that start at each position text
// // Time Complexity: O(word length * number of patterns)
vector<vector<int>> allMatches = ac.findAll(patterns, text);
```

Various (10)

10.1 Misc. algorithms

TernarySearch.h

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

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

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

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

LIS.h

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

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

FastKnapsack.h

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

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

b20ccc, 16 lines

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

10.2 Dynamic programming

KnuthDD1

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

DivideAndConquerDP FastMod

DivideAndConquerDP.h

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

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

d38d2b, 18 lines

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

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

10.3 Debugging tricks

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

10.4 Optimization tricks

__builtin_ia32_ldmxcsr(40896); disables denormals (which make floats 20x slower near their minimum value).

10.4.1 Bit hacks

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

10.4.2 Pragmas

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

FastMod.h

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

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