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template .bashrc .vimrc troubleshoot

Contest (1)

```
template.cpp
```

```
#include <bits/stdc++.h>
using namespace std;
#define rep(i, a, b) for(int i = a; i < (b); ++i)
#define trav(a, x) for(auto& a : x)
#define all(x) x.begin(), x.end()
#define sz(x) (int)(x).size()
typedef long long 11;
typedef pair<int, int> pii;
typedef vector<int> vi;
int main() {
 cin.sync_with_stdio(0); cin.tie(0);
  cin.exceptions(cin.failbit);
```

.bashrc

```
alias c='g++ -Wall -Wconversion -Wfatal-errors -g -std=c++14 \
 -fsanitize=undefined,address'
xmodmap -e 'clear lock' -e 'keycode 66=less greater' \#caps = \diamondsuit
```

.vimrc

set cin aw ai is ts=4 sw=4 tm=50 nu noeb bg=dark ru cul sy on | im jk <esc> | im kj <esc>

troubleshoot.txt

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

Wrong answer:

Any uninitialized variables?

Any assertions that might fail?

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

Are you reading or writing outside the range of any vector?

Any possible division by 0? (mod 0 for example)

Any possible infinite recursion? Invalidated pointers or iterators? Are you using too much memory? Debug with resubmits (e.g. remapped signals, see Various).

Time limit exceeded:

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

Memory limit exceeded:

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

Mathematics (2)

2.1 Equations

$$ax + by = e$$

$$cx + dy = f$$

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

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

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.

Recurrences

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

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

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

Trigonometry

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

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

 $(V+W)\tan(v-w)/2 = (V-W)\tan(v+w)/2$ where V, W are lengths of sides opposite angles v, w.

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

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

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

2.4 Geometry

2.4.1 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:
$$\frac{a+b}{a-b} = \frac{\tan \frac{\alpha+\beta}{2}}{\tan \frac{\alpha-\beta}{2}}$$

2.4.2 Quadrilaterals

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

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

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

2.4.3 Spherical coordinates



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

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

$$z = r \cos \theta \qquad \phi = a\tan(2y, x)$$

2.5 Linear algebra

2.5.1 Matrix inverse

The inverse of a 2x2 matrix:

$$\begin{bmatrix} a & b \\ c & d \end{bmatrix}^{-1} = \frac{1}{ad - bc} \begin{bmatrix} d & -b \\ -c & a \end{bmatrix}$$

In general:

$$A^{-1} = \frac{1}{\det(A)}A^*$$

where $A_{i,j}^* = (-1)^{i+j} * \Delta_{i,j}$ and $\Delta_{i,j}$ is the determinant of matrix A crossing out line i and column j.

2.6 Derivatives/Integrals

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

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

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

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

Integration by parts:

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

2.7 Sums

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

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

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

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

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

2.8 Series

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

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

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

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

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

2.9 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,

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

2.9.1 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 wich 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 $Po(\lambda)$, $\lambda = t\kappa$.

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

2.9.2 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.

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

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

50 lines

OrderStatisticTree SegmentTree LineContainer ConvexTree

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

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

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

Data structures (3)

OrderStatisticTree.h

Description: A set (not multiset!) with support for finding the n'th element, and finding the index of an element. **Time:** $\mathcal{O}(\log N)$

SegmentTree.h

Description: Very fast and quick segment tree. Only useful for easy invariants. 0-indexed. Range queries are half-open.

```
struct SegmTree {
  vector<int> T; int n;
  SegmTree(int n) : T(2 * n, (int)-2e9), n(n) {}

void Update(int pos, int val) {
  for (T[pos += n] = val; pos > 1; pos /= 2)
    T[pos / 2] = max(T[pos], T[pos ^ 1]);
}

int Query(int b, int e) {
  int res = -2e9;
  for (b += n, e += n; b < e; b /= 2, e /= 2) {
    if (b % 2) res = max(res, T[b++]);
    if (e % 2) res = max(res, T[--e]);
  }
  return res;
}
};</pre>
```

LineContainer.h

Description: Container where you can add lines of the form ax+b, and query maximum values at points x. For each line, also keeps a value p, which is the last (maximum) point for which the current line is dominant. (obviously, for the last line, p is infinity) Useful for dynamic programming. **Time:** $\mathcal{O}(\log N)$

```
_<bits/stdc++.h> 35 lines
using T = long long;
```

```
bool QUERY;
struct Line {
  mutable T a, b, p;
    T Eval(T x) const { return a * x + b; }
  bool operator<(const Line& o) const</pre>
    return QUERY ? p < o.p : a < o.a;
struct LineContainer : multiset<Line> {
  // for doubles, use kInf = 1/.0, div(a, b) = a / b
  const T kInf = numeric_limits<T>::max();
 T div(T a, T b) { // floored division
    return a / b - ((a ^ b) < 0 && a % b); }
 bool isect(iterator x, iterator y) {
    if (y == end()) { x->p = kInf; return false; }
    if (x->a == y->a) x->p = x->b > y->b ? kInf : -kInf;
    else x -> p = div(y -> b - x -> b, x -> a - y -> a);
    return x->p >= y->p;
  void InsertLine(T a, T b) {
    auto nx = insert({a, b, 0}), it = nx++, pv = it;
    while (isect(it, nx)) nx = erase(nx);
    if (pv != begin() && isect(--pv, it)) isect(pv, it = erase(
    while ((it = pv) != begin() && (--pv)->p >= it->p)
      isect(pv, erase(it));
 T EvalMax(T x) {
    assert(!emptv());
    OUERY = 1; auto it = lower bound(\{0,0,x\}); OUERY = 0;
    return it->Eval(x);
};
```

ConvexTree.h

Description: Container where you can add lines of the form a * x + b, and query maximum values at points x. Useful for dynamic programming. To change to minimum, either change the sign of all comparisons, the initialization of T and max to min, or just add lines of form $(-a)^*X + (-b)$ instead and negate the result.

```
};
const int64 kInf = 2e18; // Maximum abs(A * x + B)
const int kMin = -1e9, kMax = 1e9; // Bounds of query (x)

struct ConvexTree {
    struct Node { int 1, r; Line line; };
    vector<Node> T = { Node{0, 0, {0, -kInf}} };
    int root = 0;

int update(int node, int b, int e, Line upd) {
    if (node == 0) {
        T.push_back(Node{0, 0, upd});
        return T.size() - 1;
    }

    auto& cur = T[node].line;
    if (cur.Eval(b)>=upd.Eval(b) && cur.Eval(e)>=upd.Eval(e))
        return node;
    if (cur.Eval(b)<=upd.Eval(b) && cur.Eval(e)<=upd.Eval(e))
        return cur = upd, node;</pre>
```

Treap FenwickTree FenwickTree2d

```
int m = (b + e) / 2;
  if (cur.Eval(b) < upd.Eval(b)) swap(cur, upd);</pre>
 if (cur.Eval(m) >= upd.Eval(m)) {
   int res = update(T[node].r, m + 1, e, upd);
   T[node].r = res; // DO NOT ATTEMPT TO OPTIMIZE
   swap(cur, upd);
   int res = update(T[node].1, b, m, upd);
   T[node].1 = res; // DO NOT ATTEMPT TO OPTIMIZE
  return node;
void AddLine(Line 1) { root = update(root, kMin, kMax, 1); }
int64 query(int node, int b, int e, int x) {
 int64 ans = T[node].line.Eval(x);
  if (node == 0) return ans;
 int m = (b + e) / 2;
 if (x \le m) ans = max(ans, query(T[node].1, b, m, x));
 if (x > m) ans = max(ans, query(T[node].r, m + 1, e, x));
 return ans;
int64 QueryMax(int x) { return query(root, kMin, kMax, x); }
```

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. It can support several operations, including lazy propagation (sample reverse operation below). Can be made persistent, by making a copy at pull function To transform into ordered set, uncomment line at (A) and delete the subtraction logic at (B).

```
Time: \mathcal{O}(\log N) expected time per operation
                                                           109 lines
namespace Treap {
  struct Node {
    int val, pri;
    int left = 0, right = 0, subsize = 0, lazy = 0;
   Node(int val, int pri) : val(val), pri(pri) {}
  vector<Node> T(1, Node(-1, -1));
  int get_key(int node) {
   return T[T[node].left].subsize;
   // return T[node].val; (A)
  int pull(int node) {
    if (node == 0) return 0;
   T[node].subsize = T[T[node].left].subsize
     + T[T[node].right].subsize + 1;
    return node;
  int push (int node) {
    int& lazy = T[node].lazy;
    if (node == 0 or lazy == 0) return node;
    swap(T[node].left, T[node].right);
    T[T[node].left].lazy ^= lazy;
    T[T[node].right].lazy ^= lazy;
    lazy = 0;
    return node;
```

```
// Splits into < key and >= key
pair<int, int> Split(int node, int key) {
  push (node);
  if (node == 0) return {0, 0};
  int 1, r:
  if (get_key(node) < key) {</pre>
    tie(1, r) = Split(T[node].right, key-get_key(node)-1);
    T[node].right = 1;
    return {pull(node), r};
    tie(1, r) = Split(T[node].left, key);
    T[node].left = r;
    return {1, pull(node)};
}
// keys(node1) <= keys(node2) is REQUIRED
int Join(int node1, int node2) {
  push (node1); push (node2);
  if (!node1) return node2;
  if (!node2) return node1;
  if (T[node1].pri > T[node2].pri) {
    T[node1].right = Join(T[node1].right, node2);
    return pull(node1);
    T[node2].left = Join(node1, T[node2].left);
    return pull(node2);
// Can be any foreach function
void Dump(int node) {
  push (node);
  if (node == 0) return;
  Dump(T[node].left);
  cout << T[node].val << " ";
  Dump(T[node].right);
int Single(int value) {
  int node = T.size();
  T.push_back(Node(value, rand()));
  return pull (node);
// Only makes sense for cartesian tree
tuple<int, int, int> Slice(int node, int b, int e) {
  int 1, m, r;
  tie(m, r) = Split(node, e);
  tie(1, m) = Split(m, b);
  return make_tuple(1, m, r);
int Find(int node, int key) {
  int 1, m, r;
  tie(1, m, r) = Slice(node, key, key + 1);
  assert (node == Join(l, Join(m, r)));
  return m;
int Insert(int node, int key, int value) {
  int 1, r, m = Single(value);
  tie(1, r) = Split(node, key);
  return Join(l, Join(m, r));
```

```
int Reverse(int node) {
 T[node].lazy ^= 1;
  return push (node);
```

FenwickTree.h

Description: Adds a value to a (half-open) range and computes the sum on a (half-open) range. Beware of overflows!

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

```
#define int long long
struct FenwickTree {
 int n;
 vector<int> T1, T2;
 FenwickTree(int n): n(n), T1(n + 1, 0), T2(n + 1, 0) {}
 void Update(int b, int e, int val) {
   if (e !=-1)
     return Update(e, -1, -val), Update(b, -1, +val);
   int c1 = val, c2 = val * (b - 1);
    for (int pos = b + 1; pos <= n; pos += (pos & -pos)) {</pre>
     T1[pos] += c1; T2[pos] += c2;
 int Ouerv(int b, int e) {
   if (b != 0) return Query(0, e) - Query(0, b);
   int ans = 0;
   for (int pos = e; pos; pos -= (pos & -pos)) {
     ans += T1[pos] * (e - 1) - T2[pos];
   return ans;
};
```

FenwickTree2d.h

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

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

"FenwickTree.h" 32 lines

```
struct Fenwick2D {
 vector<vector<int>> ys;
 vector<vector<int>> T;
 Fenwick2D(int n) : ys(n + 1) {}
 void FakeUpdate(int x, int y) {
   for (++x; x < (int)ys.size(); x += (x & -x))
     ys[x].push_back(y);
 void Init() {
   for (auto& v : ys) {
     sort(v.begin(), v.end());
     T.emplace_back(v.size());
 int ind(int x, int y) {
   auto it = lower_bound(ys[x].begin(), ys[x].end(), y);
   return distance(ys[x].begin(), it);
 void Update(int x, int y, int val) {
   for (++x; x < (int)ys.size(); x += (x & -x))
   for (int i = ind(x,y); i < (int)T[x].size(); i += (i & -i))
```

```
trees[x][i] = trees[x][i] + val;
  int Query(int x, int y) {
    int sum = 0;
    for (; x > 0; x -= (x \& -x))
    for (int i = ind(x,y); i > 0; i -= (i & -i))
     sum = sum + T[x][i];
    return sum;
};
```

RMQ.h

Description: Range Minimum Queries on an array. Returns min(V[a], V[a + 1], ... V[b - 1]) in constant time. Set inf to something reasonable before

Usage: RMQ rmq(values); rmq.Query(inclusive, exclusive); Time: $\mathcal{O}(|V|\log|V|+Q)$

```
template <class T>
struct RMQ {
  const int kInf = numeric_limits<T>::max();
  vector<vector<T>> rmq;
  RMQ(const vector<T>& V) {
   int n = V.size(), on = 1, depth = 1;
    while (on < n) on *= 2, ++depth;
    rmq.assign(depth, V);
    for (int i = 0; i < depth - 1; ++i)
     for (int j = 0; j < n; ++j) {</pre>
        jmp[i + 1][j] = min(jmp[i][j],
          jmp[i][min(n - 1, j + (1 << i))]);
 T Ouery(int a, int b) {
   if (b <= a) return kInf;</pre>
   int dep = 31 - __builtin_clz(b - a); // log(b - a)
```

return min(rmq[dep][a], rmq[dep][b - (1 << dep)]);</pre>

Numerical (4)

FracBinarySearch.h

};

Description: Does binary search on fractions having an upper bound on the numerator, given a predicate pred. The predicate should be a monotonous function going from negative to 0 to positive. The function will find a RAN-DOM fraction f for which pred(f) = 0 or THROW exception 5.

```
Time: \mathcal{O}(\log((b-a)/\epsilon))
```

```
34 lines
using int64 = int64_t;
struct Frac { int64 a, b; };
template<typename Predicate>
Frac FracBinarySearch(int64 max_num, Predicate pred) {
  // Range is OPEN ON BOTH ENDS
  Frac lo{0, 1}, hi{1, 1}; // set to {1, 0} for (0...max_num)
  int sign = 1;
  // Number of tries should be >= 2 * log(max_num)
  for (int tries = 0; tries < 135; ++tries) {</pre>
    int64 adv = 0;
   bool down = false;
    for (int64 step = 1; step; down ? step \neq 2 : step \star= 2) {
     adv += step;
     Frac mid{lo.a * adv + hi.a, lo.b * adv + hi.b};
```

```
if (abs(mid.a) > kLim or mid.b > max_num) {
     adv -= step; down = true; continue;
    int64 res = pred(mid);
   if (res == 0) return mid;
   if (res * sign < 0) { adv -= step; down = true; }</pre>
 hi.a += lo.a * adv;
 hi.b += lo.b * adv;
 sign = -sign;
  swap(lo, hi);
throw 5;
```

GoldenSectionSearch.h

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

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

Time: $\mathcal{O}(\log((b-a)/\epsilon))$ 15 lines

```
template<typename Func>
double GoldenSectionSearch(double a, double b, Func f) {
 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 + b) / 2;
```

HillClimbing.h

Description: Poor man's optimization for unimodal functions. Finds minimum of a function func[Point] => double. To change with maximum, change the comparison at (*) 16 lines

```
struct Point { double x, y; };
template<typename Func>
pair < double, Point > HillClimb (Point p, Func func) {
 double best = func(p);
 for (double step = 1e9; step > 1e-20; step /= 2)
    for (int it = 0; it < 100; ++it)</pre>
      for (int dx = -1; dx \le 1; ++dx)
        for (int dy = -1; dy \le 1; ++dy) {
          Point q = p; q.x += dx * step; q.y += dy * step;
          double now = func(q);
          if (best > now) { best = now; p = q; } // (*)
 return make_pair(best, p);
```

Polynomial.h

```
field.
<br/>bits/stdc++.h>
using TElem = double;
using Poly = vector<TElem>;
TElem Eval(const Poly& P, TElem x) {
  TElem val = 0:
  for (int i = (int)P.size() - 1; i >= 0; --i)
   val = val * x + P[i];
 return val;
// Differentiation
Polv Diff(Polv P) {
 for (int i = 1; i < (int)P.size(); ++i)</pre>
   P[i - 1] = i * P[i];
 P.pop_back();
  return P:
// Integration
Poly Integrate (Poly p) {
 P.push_back(0);
  for (int i = (int)P.size() - 2; i >= 0; --i)
   P[i + 1] = P[i] / (i + 1);
  P[0] = 0;
  return P;
// Division by (X - x0)
Poly DivRoot (Poly P, TElem x0) {
 int n = P.size();
 TElem a = P.back(), b; P.back() = 0;
  for (int i = n--; i--; )
   b = P[i], P[i] = P[i + 1] * x0 + a, a = b;
 P.pop_back();
 return P:
// Multiplication modulo X^sz
Poly Multiply (Poly A, Poly B, int sz) {
 static FFTSolver fft;
 A.resize(sz, 0); B.resize(sz, 0);
 auto R = fft.Multiply(A, B);
 R.resize(sz, 0);
  return r;
// Scalar multiplication
Poly Scale (Poly P, TElem s) {
 for (auto& x : P)
   x = x * s;
 return P;
// Addition modulo X^sz
Poly Add (Poly A, Poly B, int sz) {
 A.resize(sz, 0); B.resize(sz, 0);
 for (int i = 0; i < sz; ++i)</pre>
   A[i] = A[i] + B[i];
 return A;
// ****************
// For Invert, Sqrt, size of argument should be 2 k
// ****************
Poly inv_step(Poly res, Poly P, int n) {
```

Description: Different operations on polynomials. Should work on any

```
auto res_sq = Multiply(res, res, n);
  auto sub = Multiply(res_sq, P, n);
  res = Add(Scale(res, 2), Scale(sub, -1), n);
  return res;
// Inverse modulo X^sz
// EXISTS ONLY WHEN P[0] IS INVERTIBLE
Poly Invert (Poly P) {
 assert(P[0].Get() == 1);
                           // i.e., P[0]^{(-1)}
  Poly res(1, 1);
  int n = P.size();
  for (int step = 2; step <= n; step *= 2) {
   res = inv step(res, P, step);
  // Optional, but highly encouraged
  auto check = Multiply(res, P, n);
  for (int i = 0; i < n; ++i) {
   assert(check[i].Get() == (i == 0));
 return res;
// Square root modulo X^sz
// EXISTS ONLY WHEN P[0] HAS SQUARE ROOT
Poly Sqrt (Poly P) {
  assert(P[0].Get() == 1);
                           // i.e., P[0]^{(-1)}
  Poly res(1, 1);
                           // i.e., P[0]^(1/2)
  Poly inv(1, 1);
  int n = P.size();
  for (int step = 2; step <= n; step *= 2) {
   auto now = inv_step(inv, res, step);
   now = Multiply(P, move(now), step);
   res = Add(res, now, step);
   res = Scale(res, (kMod + 1) / 2);
   inv = inv_step(inv, res, step);
  // Optional, but highly encouraged
  auto check = Multiply(res, res, n);
  for (int i = 0; i < n; ++i) {</pre>
   assert(check[i].Get() == P[i].Get());
 return res;
PolyRoots.h
Description: Finds the real roots to a polynomial.
Usage: Poly p = \{2, -3, 1\} // x^2 - 3x + 2 = 0
auto roots = GetRoots(p, -1e18, 1e18); // {1, 2}
<bits/stdc++.h>, "Polynomial.h"
                                                           26 lines
vector<double> GetRoots(Poly p, double xmin, double xmax) {
 if (p.size() == 2) { return {-p.front() / p.back()}; }
  else {
   Polv d = Diff(p);
   vector<double> dr = GetRoots(d, xmin, xmax);
   dr.push_back(xmin - 1);
   dr.push_back(xmax + 1);
    sort(dr.begin(), dr.end());
    vector<double> roots;
    for (auto i = dr.begin(), j = i++; i != dr.end(); j = i++) {
     double lo = *j, hi = *i, mid, f;
     bool sign = Eval(p, lo) > 0;
     if (sign ^ (Eval(p, hi) > 0)) {
```

```
// \ for \ (int \ it = 0; \ it < 60; ++it)  {
         while (hi - lo > 1e-8) {
          mid = (lo + hi) / 2, f = Eval(p, mid);
          if ((f <= 0) ^ sign) lo = mid;</pre>
          else hi = mid;
        roots.push_back((lo + hi) / 2);
    return roots;
PolyInterpolate.h
Description: Given n points (x[i], y[i]), computes an n-1-degree polyno-
mial p that passes through them: p(x) = a[0] * x^0 + ... + a[n-1] * x^{n-1}. For
numerical precision, pick x[k] = c * \cos(k/(n-1)*\pi), k = 0 \dots n-1.
Time: \mathcal{O}\left(n^2\right)
<bits/stdc++.h>, "Polynomial.h"
                                                                15 lines
Poly Interpolate (vector<TElem> x, vector<TElem> y) {
 int n = x.size();
 Poly res(n), temp(n);
 for (int k = 0; k < n; ++k)
    for (int i = k + 1; i < n; ++i)
   y[i] = (y[i] - y[k]) / (x[i] - x[k]);
  TElem last = 0; temp[0] = 1;
  for (int k = 0; k < n; ++k)
  for (int i = 0; i < n; ++i) {</pre>
    res[i] = res[i] + y[k] * temp[i];
    swap(last, temp[i]);
    temp[i] = temp[i] - last * x[k];
 return res;
BerlekampMassev.h
Description: Recovers any n-order linear recurrence relation from the first
2*n terms of the recurrence. Very useful for guessing linear recurrences af-
ter brute-force / backtracking the first terms. Should work on any field.
```

Numerical stability for floating-point calculations is not guaranteed. **Usage:** BerlekampMassey($\{0, 1, 1, 3, 5, 11\}$) => $\{1, 2\}$

 dits/stdc++.h>, "ModOps.h" vector<ModInt> BerlekampMassey(vector<ModInt> s) { int n = s.size(); vector<ModInt> C(n, 0), B(n, 0); C[0] = B[0] = 1;ModInt b = 1; int L = 0; for (int i = 0, m = 1; i < n; ++i) { ModInt d = s[i];for (int j = 1; j <= L; ++j)</pre> d = d + C[j] * s[i - j];if (d.get() == 0) { ++m; continue; } auto T = C; ModInt coef = d * inv(b); for (int j = m; j < n; ++j) C[j] = C[j] - coef * B[j - m];if (2 * L > i) { ++m; continue; } L = i + 1 - L; B = T; b = d; m = 1; C.resize(L + 1); C.erase(C.begin()); for (auto& x : C) x = ModInt(0) - x;

```
return C;
```

LinearRecurrence.h

Description: Generates the k-th term of a n-th order linear recurrence given the first n terms and the recurrence relation. Faster than matrix multiplication. Useful to use along with Berlekamp Massey.

Usage: LinearRec<double $>(\{0, 1\}, \{1, 1\}).Get(k)$ gives k-th Fibonacci number (0-indexed)

Time: $\mathcal{O}\left(n^2log(k)\right)$ per query

dits/stdc++.h>

```
template<typename T>
struct LinearRec {
  using Poly = vector<T>;
  int n; Poly first, trans;
  // Recurrence is S[i] = sum(S[i-j-1] * trans[j])
  // \ with \ S[0..(n-1)] = first
  LinearRec(const Poly &first, const Poly &trans) :
    n(first.size()), first(first), trans(trans) {}
  Poly combine (Poly a, Poly b) {
    Poly res(n * 2 + 1, 0);
    // You can apply constant optimization here to get a
    // \sim 10x speedup
    for (int i = 0; i <= n; ++i)</pre>
      for (int j = 0; j <= n; ++j)
        res[i + j] = res[i + j] + a[i] * b[j];
    for (int i = 2 * n; i > n; --i)
      for (int j = 0; j < n; ++j)
        res[i - 1 - j] = res[i - 1 - j] + res[i] * trans[j];
    res.resize(n + 1);
    return res:
  // Consider caching the powers for multiple queries
  T Get (int k) {
    Poly r(n + 1, 0), b(r);
    r[0] = 1; b[1] = 1;
    for (++k; k; k /= 2) {
     if (k % 2)
       r = combine(r, b);
      b = combine(b, b);
    T res = 0;
    for (int i = 0; i < n; ++i)</pre>
     res = res + r[i + 1] * first[i];
    return res;
};
```

FFT.h

Description: Fast Fourier transform. Also includes a function for convolution: conv(a, b) = c, where $c[x] = \sum a[i]b[x-i]$. a and b should be of roughly equal size. Does about 1.2s for 10^6 elements. Rounding the results of conv works if $(|a|+|b|) \max(a,b) < \sim 10^9$ (in theory maybe 10^6); you may want to use an NTT from the Number Theory chapter instead.

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

```
<bits/stdc++.h>
struct FFTSolver {
 using Complex = complex<double>;
 const double kPi = 4.0 * atan(1.0);
```

FST Integrate IntegrateAdaptive Simplex

```
vector<int> rev;
int __lq(int n) { return n == 1 ? 0 : 1 + __lq(n / 2); }
void compute_rev(int n, int lg) {
 rev.resize(n); rev[0] = 0;
 for (int i = 1; i < n; ++i) {</pre>
   rev[i] = (rev[i >> 1] >> 1) | ((i & 1) << (lq - 1));
vector<Complex> fft(vector<Complex> V, bool invert) {
  int n = V.size(), lg = lg(n);
  if ((int)rev.size() != n) compute_rev(n, lg);
  for (int i = 0; i < n; ++i) {</pre>
   if (i < rev[i])
     swap(V[i], V[rev[i]]);
  for (int step = 2; step <= n; step *= 2) {</pre>
   const double ang = 2 * kPi / step;
   Complex eps(cos(ang), sin(ang));
   if (invert) eps = conj(eps);
    for (int i = 0; i < n; i += step) {
     Complex w = 1:
     for (int a = i, b = i+step/2; b < i+step; ++a, ++b) {</pre>
       Complex aux = w * V[b];
       V[b] = V[a] - aux;
       V[a] = V[a] + aux;
       w *= eps;
  return V:
vector<Complex> transform(vector<Complex> V) {
 int n = V.size();
 vector<Complex> ret(n);
 Complex div_x = Complex(0, 1) * (4.0 * n);
  for (int i = 0; i < n; ++i) {
   int j = (n - i) % n;
   ret[i] = (V[i] + conj(V[j]))
     * (V[i] - conj(V[j])) / div_x;
  return ret;
vector<int> Multiply(vector<int> A, vector<int> B) {
 int n = A.size() + B.size() - 1;
 vector<int> ret(n);
 while (n != (n \& -n)) ++n;
 A.resize(n); B.resize(n);
 vector<Complex> V(n);
 for (int i = 0; i < n; ++i) {</pre>
   V[i] = Complex(A[i], B[i]);
 V = fft(move(V), false);
 V = transform(move(V));
 V = fft(move(V), true);
  for (int i = 0; i < (int)ret.size(); ++i)</pre>
```

```
ret[i] = round(real(V[i]));
    return ret;
};
FST.h
Description: Fast Subset transform. Useful for performing the following
convolution: R[a \text{ op } b] += A[a] * B[b], where op is either of AND, OR, XOR.
P has to have size N=2^n, for some n.
Time: \mathcal{O}(N \log N)
<br/>
<br/>
dits/stdc++.h>
vector<int> Transform(vector<int> P, bool inv) {
 int n = P.size();
  for (int step = 1; step < n; step *= 2) {</pre>
    for (int i = 0; i < n; i += 2 * step) {
      for (int j = i; j < i + step; ++j) {</pre>
        int u = P[j], v = P[j + step];
        tie(P[j], P[j + step]) =
        inv ? make_pair(v - u, u) : make_pair(v, u + v); // AND
        inv ? make_pair(v, u - v) : make_pair(u + v, u); // OR
        make_pair(u + v, u - v);
    }
 // if (inv) for (auto\& x : P) x /= n; // XOR only
 return P;
Integrate.h
Description: Simple integration of a function over an interval using Simp-
son'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
template<typename Func>
double Quad(Func f, double a, double b) {
  const int n = 1000;
  double h = (b - a) / 2 / n;
  double v = f(a) + f(b);
  for (int i = 1; i < 2 * n; ++i)
   v += f(a + i * h) * (i & 1 ? 4 : 2);
  return v * h / 3;
IntegrateAdaptive.h
Description: Fast integration using an adaptive Simpson's rule.
Usage: double z, v;
double h (double x) { return x*x + y*y + z*z <= 1; }
double g(double y) \{ :: y = y; return Quad(h, -1, 1); \}
double f(double z) \{ :: z = z; return Quad(q, -1, 1); \}
double sphereVol = Quad(f, -1, 1), pi = sphereVol*3/4;
                                                              23 lines
template<typename Func>
double simpson (Func f, double a, double b) {
 double c = (a + b) / 2;
 return (f(a) + 4 * f(c) + f(b)) * (b - a) / 6;
template<typename Func>
double recurse (Func f, double a, double b,
                double eps, double S) {
  double c = (a + b) / 2;
  double S1 = simpson(f, a, c);
 double S2 = simpson(f, c, b);
  double T = S1 + S2;
 if (abs(T - S) < 15 * eps || b - a < 1e-10)</pre>
    return T + (T - S) / 15;
```

return recurse (f, a, c, eps / 2, S1) +

```
recurse(f, c, b, eps / 2, S2);
}
template<typename Func>
double Quad(Func f, double a, double b, double eps = 1e-8) {
   return recurse(f, a, b, eps, simpson(f, 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: A = {{1,-1}, {-1,1}, {-1,-2}};
b = {1,1,-4}, c = {-1,-1}, x;
T val = LPSolver(A, b, c).solve(x);
```

pivot(r, s);

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

```
typedef double T; // long double, Rational, double + mod<P>...
typedef vector<T> vd;
typedef vector<vd> vvd;
const T eps = 1e-8, inf = 1/.0;
#define MP make_pair
#define ltj(X) if (s == -1 \mid \mid MP(X[j], N[j]) < MP(X[s], N[s])) s=j
#define rep(i, a, b) for(int i = a; i < (b); ++i)
#define sz(x) (int)(x).size()
struct LPSolver {
 int m, n; vi N, B; vvd D;
  LPSolver (const vvd& A, const vd& b, const vd& c) :
    m(sz(b)), n(sz(c)), N(n+1), B(m), D(m+2), vd(n+2)) {
      rep(i, 0, m) rep(j, 0, n) D[i][j] = A[i][j];
      rep(i,0,m) \{ B[i] = n+i; D[i][n] = -1; D[i][n+1] = b[i]; \}
      rep(j,0,n) \{ N[j] = j; D[m][j] = -c[j]; \}
      N[n] = -1; D[m+1][n] = 1;
 void pivot(int r, int s) {
    T *a = D[r].data(), inv = 1 / a[s];
    rep(i, 0, m+2) if (i != r && abs(D[i][s]) > eps) {
     T *b = D[i].data(), inv2 = b[s] * inv;
      rep(j, 0, n+2) b[j] = a[j] * inv2;
      b[s] = a[s] * inv2;
    rep(j,0,n+2) if (j != s) D[r][j] *= inv;
    rep(i,0,m+2) if (i != r) D[i][s] *= -inv;
    D[r][s] = inv;
    swap(B[r], N[s]);
 bool simplex(int phase) {
    int x = m + phase - 1;
    for (;;) {
      int s = -1;
      rep(j,0,n+1) if (N[j] != -phase) ltj(D[x]);
      if (D[x][s] >= -eps) return true;
      int r = -1;
      rep(i,0,m) {
       if (D[i][s] <= eps) continue;</pre>
        if (r == -1 || 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;
```

```
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;
};
```

SolveLinear.h

Description: Solves M * x = b. If there are multiple solutions, returns a solution which has all free variables set to 0. To compute rank, count the number of values in pivot. vector which are not -1. For inverse modulo prime powers, repeatedly set $A^{-1} = A^{-1}(2I - AA^{-1}) \pmod{p^k}$ where A^{-1} starts as the inverse of A mod p, and k is doubled in each step.

```
Time: \mathcal{O}\left(vars^2cons\right)
// Transforms a matrix into its row echelon form
// Returns a vector of pivots for each variable
// vars is the number of variables to do echelon for
vector<int> ToRowEchelon(vector<vector<double>> &M, int vars) {
  int n = M.size(), m = M[0].size();
  vector<int> pivots(vars, -1);
  int cur = 0;
  for (int var = 0; var < vars; ++var) {</pre>
   if (cur >= n) break;
   for (int con = cur + 1; con < n; ++con)
     if (sgn(M[con][var]) != 0)
       swap(M[con], M[cur]);
    if (sgn(M[cur][var]) != 0) {
     pivots[var] = cur;
      auto aux = M[cur][var];
      for (int i = 0; i < m; ++i)</pre>
       M[cur][i] = M[cur][i] / aux;
      for (int con = 0; con < n; ++con) {</pre>
       if (con != cur) {
          auto mul = M[con][var];
          for (int i = 0; i < m; ++i) {</pre>
            M[con][i] = M[con][i] - mul * M[cur][i];
      ++cur;
  return pivots;
// Computes the inverse of a nxn square matrix.
// Returns true if successful
bool Invert (vector<vector<double>> &M) {
```

```
int n = M.size();
 for (int i = 0; i < n; ++i) {</pre>
   M[i].resize(2 * n, 0); M[i][n + i] = 1;
 auto pivs = ToRowEchelon(M, n);
 for (auto x : pivs) if (x == -1) return false;
 for (int i = 0; i < n; ++i)</pre>
   M[i].erase(M[i].begin(), M[i].begin() + n);
 return true;
// Returns the solution of a system
// Will change matrix
// Throws 5 if inconsistent
vector<double> SolveSystem(vector<vector<double>> &M,
                           vector<double>& b) {
 int vars = M[0].size();
 for (int i = 0; i < (int)M.size(); ++i)</pre>
   M[i].push_back(b[i]);
  auto pivs = ToRowEchelon(M, vars);
 vector<double> solution(vars);
 for (int i = 0; i < vars; ++i) {</pre>
   solution[i] = (pivs[i] == -1) ? 0 : M[pivs[i]][vars];
  // Check feasible (optional)
  for (int i = 0; i < (int)M.size(); ++i) {</pre>
    double check = 0;
    for (int j = 0; j < vars; ++j)</pre>
     check = check + M[i][j] * solution[j];
    if (sgn(check - M[i][vars]) != 0)
      throw 5;
 return solution;
```

Tridiagonal.h

Description: Solves a linear equation system with a tridiagonal matrix with diagonal diag, subdiagonal sub and superdiagonal super, i.e., x = Tridiagonal(d, p, q, b) solves the equation system

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

The size of diag and b should be the same and super and sub should be one element shorter. T is intended to be double.

This is useful for solving problems on the type

```
a_i = b_i a_{i-1} + c_i a_{i+1} + d_i, 1 < i < 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, \dots, b_n, 0\}, \{a_0, d_1, d_2, \dots, d_n, a_{n+1}\}\}
```

```
Usage: int n = 1000000;
vector<double> diag(n,-1), sup(n-1,.5), sub(n-1,.5), b(n,1);
vector<double> x = tridiagonal(diag, super, sub, b);
```

```
Time: \mathcal{O}(N)
template <typename T>
vector<T> Tridiagonal(vector<T> diag, const vector<T>& super,
    const vector<T>& sub, vector<T> b) {
 for (int i = 0; i < b.size() - 1; ++i) {</pre>
    diag[i + 1] -= super[i] * sub[i] / diag[i];
    b[i + 1] = b[i] * sub[i] / diag[i];
 for (int i = b.size(); --i > 0;) {
   b[i] /= diag[i];
   b[i - 1] -= b[i] * super[i - 1];
 b[0] /= diag[0];
 return b;
```

Number theory (5)

5.1 Modular arithmetic

ModOps.h

Description: ModOps class and operations for easy modulo reduction. Quick to code, but not fast.

```
const int kMod = 1e9 + 7;
struct ModInt {
 long long n;
 ModInt (long long n = 0) : n(n % kMod) {}
 ModInt operator+(const ModInt& oth) { return n + oth.n; }
 ModInt operator-(const ModInt& oth) { return n - oth.n; }
 ModInt operator*(const ModInt& oth) { return n * oth.n; }
 long long get() { return n < 0 ? n + kMod : n; }</pre>
ModInt lgpow (ModInt b, int e) {
 ModInt r:
  for (r = 1; e; e /= 2, b = b * b)
   if (e % 2) r = r * b;
  return r;
ModInt inv(ModInt x) { return lgpow(x, kMod - 2); }
```

ModInverse.h

Description: Pre-computation of modular inverses. Assumes $\lim < k \text{Mod}$ and that kMod is a prime.

```
"ModOps.h"
                                                             7 lines
vector<ModInt> ComputeInverses(int lim) {
 vector<ModInt> inv(lim + 1); inv[1] = 1;
 for (int i = 2; i <= lim; ++i) {</pre>
   inv[i] = ModInt(0) - ModInt(kMod / i) * inv[kMod % i];
 return inv;
```

ModSum.h

Description: Sums of mod'ed arithmetic progressions.

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

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

```
21 lines
using ull = unsigned long long;
using 11 = long long;
ull SumSq(ull to) { return to / 2 * ((to-1) | 1); }
```

ModMulLL ModSqrt NTT MillerRabin phiFunction

```
ull DivSum(ull to, ull c, ull k, ull m) {
  ull res = k / m * SumSq(to) + c / m * to;
  k %= m; c %= m;
   ull to2 = (to * k + c) / m;
    res += to * to2;
    res -= DivSum(to2, m-1 - c, m, k) + to2;
  return res;
11 ModSum(ull to, 11 c, 11 k, 11 m) {
 c %= m; if (c < 0) c += m;
 k \% = m; if (k < 0) k += 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 large c.

Time: $\mathcal{O}(64/bits \cdot \log b)$, where bits = 64 - k, if we want to deal with k-bit

```
typedef unsigned long long ull;
const int bits = 10;
// if all numbers are less than 2^k, set bits = 64-k
const ull po = 1 << bits;</pre>
ull ModMul(ull a, ull b, ull &c) {
  ull x = a * (b & (po - 1)) % c;
  while ((b >>= bits) > 0) {
   a = (a << bits) % c;
   x += (a * (b & (po - 1))) % c;
 return x % c;
ull ModPow(ull a, ull b, ull mod) {
 if (b == 0) return 1;
  ull res = ModPow(a, b / 2, mod);
  res = ModMul(res, res, mod);
 if (b & 1) return ModMul(res, a, mod);
  return res;
```

ModSart.h

Description: Tonelli-Shanks algorithm for modular square roots. **Time:** $\mathcal{O}\left(\log^2 p\right)$ worst case, often $\mathcal{O}\left(\log p\right)$

```
30 lines
ll sqrt(ll a, ll p) {
 a %= p; if (a < 0) a += p;
 if (a == 0) return 0;
  assert (modpow(a, (p-1)/2, p) == 1);
  if (p % 4 == 3) return modpow(a, (p+1)/4, p);
  // a^{(n+3)/8} \text{ or } 2^{(n+3)/8} * 2^{(n-1)/4} \text{ works if } p \% 8 == 5
  11 s = p - 1;
  int r = 0;
  while (s % 2 == 0)
   ++r, s /= 2;
  11 n = 2; // find a non-square mod p
  while (modpow(n, (p - 1) / 2, p) != p - 1) ++n;
  11 x = modpow(a, (s + 1) / 2, p);
  11 b = modpow(a, s, p);
  11 q = modpow(n, s, p);
  for (;;) {
   11 t = b;
    int m = 0;
    for (; m < r; ++m) {
     if (t == 1) break;
     t = t * t % p;
```

```
if (m == 0) return x;
11 qs = modpow(q, 1 << (r - m - 1), p);
g = gs * gs % p;
x = x * gs % p;
b = b * q % p;
```

Number theoretic transform

NTT.h

Description: Number theoretic transform. Can be used for convolutions modulo specific nice primes of the form $2^a b + 1$, where the convolution result has size at most 2^a. For other primes/integers, use two different primes and combine with CRT. If NTT is not fast enough and you are multiplying a lot, consider doing naive solution for the small ones.

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

```
"ModPow.h"
const int kMod = (119 << 23) + 1, kRoot = 3; // = 998244353
// For p < 2^30 there is also e.g. (5 << 25, 3), (7 << 26, 3)
// (479 << 21, 3) and (483 << 21, 5). The last two are > 10^9.
struct FFTSolver
 vector<int> rev;
 int __lq(int n) { return n == 1 ? 0 : 1 + __lq(n / 2); }
 void compute_rev(int n, int lg) {
   rev.resize(n); rev[0] = 0;
   for (int i = 1; i < n; ++i) {
     rev[i] = (rev[i >> 1] >> 1) | ((i & 1) << (lq - 1));
 }
 vector<ModInt> fft(vector<ModInt> V, bool invert) {
   int n = V.size(), lg = __lg(n);
   if ((int)rev.size() != n) compute_rev(n, lg);
   for (int i = 0; i < n; ++i) {</pre>
     if (i < rev[i])
        swap(V[i], V[rev[i]]);
    for (int step = 2; step <= n; step *= 2) {
     ModInt eps = lgpow(kRoot, (kMod - 1) / step);
     if (invert) eps = inv(eps);
      for (int i = 0; i < n; i += step) {
       ModInt w = 1;
       for (int a = i, b = i+step/2; b < i+step; ++a, ++b) {</pre>
         ModInt aux = w * V[b];
         V[b] = V[a] - aux;
         V[a] = V[a] + aux;
         w = w * eps;
    return V;
 vector<ModInt> Multiply(vector<ModInt> A, vector<ModInt> B) {
   int n = A.size() + B.size() - 1, sz = n;
   while (n != (n \& -n)) ++n;
```

```
A.resize(n, 0); B.resize(n, 0);
A = fft (move(A), false);
B = fft(move(B), false);
vector<ModInt> ret(n);
ModInt inv_n = inv(n);
for (int i = 0; i < n; ++i) {</pre>
  ret[i] = A[i] * B[i] * inv n;
ret = fft(move(ret), true);
ret.resize(sz);
return ret;
```

Primality

Eratosthenes.h

MillerRabin.h

Description: Miller-Rabin primality probabilistic test. Probability of failing one iteration is at most 1/4. 15 iterations should be enough for 50-bit numbers.

Time: 15 times the complexity of $a^b \mod c$.

```
18 lines
```

```
using ull = unsigned long long;
bool IsPrime(ull p) {
 if (p == 2) return true;
 if (p == 1 || p % 2 == 0) return false;
 ull s = p - 1;
 while (s % 2 == 0) s /= 2;
  for (int i = 0; i < 15; ++i) {
   ull a = rand() % (p - 1) + 1, tmp = s;
   ull mod = ModPow(a, tmp, p);
    while (tmp != p - 1 && mod != 1 && mod != p - 1) {
     mod = ModMul(mod, mod, p);
     tmp *= 2;
   if (mod != p - 1 && tmp % 2 == 0) return false;
 return true;
```

Factor.h

5.4 Divisibility

Euclid.h

5.4.1 Bézout's identity

For $a \neq b \neq 0$, then d = qcd(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 totient or Euler's phi function is defined as $\phi(n) := \#$ of positive integers $\leq n$ that are coprime with n. The cototient is $n - \phi(n)$. $\phi(1) = 1$, 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 $\phi(n) = (p_1 - 1)p_1^{k_1 - 1} ... (p_r - 1)p_r^{k_r - 1}$. $\phi(n) = n \cdot \prod_{p|n} (1 - 1/p)$.

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

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

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

11 lines

5.5 Chinese remainder theorem

CRT.h

Description: Chinese Remainder Theorem.

Find z such that $z\%m_1 = r_1, z\%m_2 = r_2$. Here, z is unique modulo M = lcm(m1, m2). The vector version solves a system of equations of type $z\%m_i = p_i$. On output, return $\{0, -1\}$. Note that all numbers must be less than 2^{31} if you have type unsigned long long.

Time: $\log(m+n)$

```
pair<int, int> CRT(int m1, int r1, int m2, int r2) {
   int s, t;
   int g = Euclid(m1, m2, s, t);
   if (r1 % g != r2 % g) return make_pair(0, -1);
   int z = (s * r2 * m1 + t * r1 * m2) % (m1 * m2);
   if (z < 0) z += m1 * m2;
   return make_pair(m1 * m2 / g, z / g);
}

pair<int, int> CRT(vector<int> m, vector<int> r) {
   pair<int, int> ret = make_pair(m[0], r[0]);
   for (int i = 1; i < m.size(); i++) {
      ret = CRT(ret.first, ret.second, m[i], r[i]);
      if (ret.second == -1) break;
   }
   return ret;</pre>
```

5.6 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.

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

5.8 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.

Combinatorial (6)

6.1 The Twelvefold Way

Counts the # of functions $f: N \to K$, |N| = n, |K| = k. The elements in N and K can be distinguishable or indistinguishable, while f can be injective (one-to-one) of surjective (onto).

	N	K	none	injective	surjective
	dist	dist	k^n	$\frac{k!}{(k-n)!}$	k!S(n,k)
j	indist	dist	$\binom{n+k-1}{n}$	$\binom{k}{n}$	$\binom{n-1}{n-k}$
	dist	indist	$\sum_{t=0}^{k} S(n,t)$	$[n \le k]$	S(n,k)
j	indist	indist	$\sum_{t=1}^{k} p(n,t)$	$[n \le k]$	p(n,k)

Here, S(n, k) is the Stirling number of the second kind, and p(n, k) is the partition number.

6.2 Permutations

6.2.1 Factorial

n	1 2 3	4	5 6	7	8	9	10	
							3628800	_
							17	
$\overline{n!}$	4.0e7	7 4.8€	8 6.2e	9 8.7e	10 1.3e	e12 2.1e	13 3.6e14	_
							0 171	
n!	2e18	2e25	3e32	8e47 3	Be64 9e	157 6e2	$62 > DBL_1$	MAX

intperm.h

Description: Permutations to/from integers. The bijection is order preserving. **Time:** $\mathcal{O}\left(n^2\right)$

```
int factorial[] = {1, 1, 2, 6, 24, 120, 720, 5040}; // etc.
template <class Z, class It>
void perm_to_int(Z& val, It begin, It end) {
 int x = 0, n = 0;
 for (It i = begin; i != end; ++i, ++n)
   if (*i < *begin) ++x;
 if (n > 2) perm_to_int<Z>(val, ++begin, end);
  else val = 0;
 val += factorial[n-1] *x;
/* range [begin, end) does not have to be sorted. */
template <class Z, class It>
void int_to_perm(Z val, It begin, It end) {
 Z fac = factorial[end - begin - 1];
  // Note that the division result will fit in an integer!
 int x = val / fac;
 nth_element(begin, begin + x, end);
 swap(*begin, *(begin + x));
 if (end - begin > 2) int_to_perm(val % fac, ++begin, end);
```

6.2.2 Cycles

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

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

6.2.3 Derangements

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

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

derangements.h

Description: Generates the *i*:th derangement of S_n (in lexicographical order).

```
template <class T, int N>
struct derangements {
   T dgen[N][N], choose[N][N], fac[N];
   derangements() {
     fac[0] = choose[0][0] = 1;
     memset(dgen, 0, sizeof(dgen));
     rep(m,1,N) {
        fac[m] = fac[m-1] * m;
        choose[m][0] = choose[m][m] = 1;
        rep(k,1,m)
            choose[m][k] = choose[m-1][k-1] + choose[m-1][k];
     }
}
T DGen(int n, int k) {
   T ans = 0;
   if (dgen[n][k]) return dgen[n][k];
```

```
ans += (i&1?-1:1) * choose[k][i] * fac[n-i];
    return dgen[n][k] = ans;
  void generate(int n, T idx, int *res) {
    int vals[N];
    rep(i,0,n) vals[i] = i;
    rep(i,0,n) {
      int j, k = 0, m = n - i;
      rep(j, 0, m) if (vals[j] > i) ++k;
      rep(j,0,m)
        T p = 0;
        if (vals[j] > i) p = DGen(m-1, k-1);
        else if (vals[j] < i) p = DGen(m-1, k);
        if (idx <= p) break;</pre>
        idx -= p;
      res[i] = vals[i];
      memmove (vals + \dot{j}, vals + \dot{j} + 1, sizeof (int) * (m-\dot{j}-1));
};
```

6.2.4 Involutions

An involution is a permutation with maximum cycle length 2, and it is its own inverse.

$$a(n) = a(n-1) + (n-1)a(n-2)$$

 $a(0) = a(1) = 1$

 $1,\,1,\,2,\,4,\,10,\,26,\,76,\,232,\,764,\,2620,\,9496,\,35696,\\140152$

6.2.5 Stirling numbers of the first kind

$$s(n,k) = (-1)^{n-k}c(n,k)$$

c(n,k) is the unsigned Stirling numbers of the first kind, and they count the number of permutations on n items with k cycles.

$$s(n,k) = s(n-1,k-1) - (n-1)s(n-1,k)$$

$$s(0,0) = 1, s(n,0) = s(0,n) = 0$$

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

$$c(0,0) = 1, c(n,0) = c(0,n) = 0$$

6.2.6 Eulerian numbers

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

$$\pi(j) > \pi(j+1), \ k+1 \ j\text{:s s.t.} \ \pi(j) \geq j, \ k \ j\text{:s s.t.}$$

$$\pi(j) > j.$$

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

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

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

6.2.7 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

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

6.3 Partitions and subsets

6.3.1 Partition function

Partitions of n with exactly k parts, p(n, k), i.e., writing n as a sum of k positive integers, disregarding the order of the summands.

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

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

For partitions with any number of parts, p(n) obeys

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

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

binomial binomialModPrime RollingBinomial multinomial

6.3.2 Binomials

binomial.h

Description: The number of *k*-element subsets of an *n*-element set, $\binom{n}{k} = \frac{n!}{k!(n-k)!}$

```
Time: \mathcal{O}(\min(k, n-k))

11 choose(int n, int k) {
    11 c = 1, to = min(k, n-k);
    if (to < 0) return 0;
    rep(i,0,to) c = c * (n - i) / (i + 1);
    return c;
}
```

binomialModPrime.h

Description: Lucas' thm: Let n,m be non-negative integers and p a prime. Write $n=n_kp^k+\ldots+n_1p+n_0$ and $m=m_kp^k+\ldots+m_1p+m_0$. Then $\binom{n}{m}\equiv\prod_{i=0}^k\binom{n_i}{m_i}\pmod{p}$. fact and invfact must hold pre-computed factorials / inverse factorials, e.g. from ModInverse.h.

RollingBinomial.h

Description: $\binom{n}{k}$ (mod m) in time proportional to the difference between (n, k) and the previous (n, k).

```
const 11 mod = 1000000007;
vector<11> invs; // precomputed up to max n, inclusively
struct Bin {
  int N = 0, K = 0; 11 r = 1;
  void m(11 a, 11 b) { r = r * a % mod * invs[b] % mod; }
  11 choose(int n, int k) {
   if (k > n || k < 0) return 0;
   while (N < n) ++N, m(N, N-K);
   while (K < k) ++K, m(N-K+1, K);
   while (K > k) m(K, N-K+1), --K;
   while (N > n) m(N-K, N), --N;
   return r;
  }
};
```

multinomial.h

Description: $\binom{\sum k_i}{k_1, k_2, \dots, k_n} = \frac{(\sum k_i)!}{k_1! k_2! \dots k_n!}$ Time: $\mathcal{O}((\sum k_i) - k_1)$

```
Time: O((\sum k_i) - k_1) 6 lines

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

6.3.3 Stirling numbers of the second kind

Partitions of n distinct elements into exactly k groups.

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

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

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

6.3.4 Bell numbers

Total number of partitions of n distinct elements.

$$B(n) = \sum_{k=1}^{n} {n-1 \choose k-1} B(n-k) = \sum_{k=1}^{n} S(n,k)$$
$$B(0) = B(1) = 1$$

The first are 1, 1, 2, 5, 15, 52, 203, 877, 4140, 21147, 115975, 678570, 4213597. For a prime p

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

6.3.5 Triangles

Given rods of length $1, \ldots, n$,

$$T(n) = \frac{1}{24} \begin{cases} n(n-2)(2n-5) & n \text{ even} \\ (n-1)(n-3)(2n-1) & n \text{ odd} \end{cases}$$

is the number of distinct triangles (positive are) that can be constructed, i.e., the # of 3-subsets of [n] s.t. $x \le y \le z$ and $z \ne x + y$.

6.4 General purpose numbers

6.4.1 Catalan numbers

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

$$C_{n+1} = \frac{2(2n+1)}{n+2} C_n$$

$$C_0 = 1, C_{n+1} = \sum_{i=1}^{n} C_i C_{n-i}$$

First few are 1, 1, 2, 5, 14, 42, 132, 429, 1430, 4862, 16796, 58786, 208012, 742900.

- # of monotonic lattice paths of a $n \times n$ -grid which do not pass above the diagonal.
- # of expressions containing n pairs of parenthesis which are correctly matched.
- # of full binary trees with with n + 1 leaves (0 or 2 children).
- # of non-isomorphic ordered trees with n+1 vertices.
- # of ways a convex polygon with n + 2 sides can be cut into triangles by connecting vertices with straight lines.
- # of permutations of [n] with no three-term increasing subsequence.

6.4.2 Super Catalan numbers

The number of monotonic lattice paths of a $n \times n$ -grid that do not touch the diagonal.

$$S(n) = \frac{3(2n-3)S(n-1) - (n-3)S(n-2)}{n}$$
$$S(1) = S(2) = 1$$

1, 1, 3, 11, 45, 197, 903, 4279, 20793, 103049, 518859

6.4.3 Motzkin numbers

Number of ways of drawing any number of nonintersecting chords among n points on a circle. Number of lattice paths from (0,0) to (n,0) never going below the x-axis, using only steps NE, E, SE.

$$M(n) = \frac{3(n-1)M(n-2) + (2n+1)M(n-1)}{n+2}$$
$$M(0) = M(1) = 1$$

1, 1, 2, 4, 9, 21, 51, 127, 323, 835, 2188, 5798, 15511, 41835, 113634

6.4.4 Narayana numbers

Number of lattice paths from (0,0) to (2n,0) never going below the x-axis, using only steps NE and SE, and with k peaks.

$$N(n,k) = \frac{1}{n} \binom{n}{k} \binom{n}{k-1}$$
$$N(n,1) = N(n,n) = 1$$

EulerWalk DinicFlow EZFlow MinCostMaxFlow

```
\sum_{k=1}^{n} N(n,k) = C_n
```

1, 1, 1, 1, 3, 1, 1, 6, 6, 1, 1, 10, 20, 10, 1, 1, 15, 50

6.4.5 Schröder numbers

Number of lattice paths from (0,0) to (n,n) using only steps N,NE,E, never going above the diagonal. Number of lattice paths from (0,0) to (2n,0) using only steps NE, SE and double east EE, never going below the x-axis. Twice the Super Catalan number, except for the first term. 1, 2, 6, 22, 90, 394, 1806, 8558, 41586, 206098

Graph (7)

7.1 Euler walk

EulerWalk.h

Description: Eulerian undirected/directed path/cycle algorithm. For a directed / undirected graph. For each path/cycle found, calls a callback. You can check cycle by checking path endpoints. To transform into undirected, toggle comment on lines (*)

Time: $\mathcal{O}(E)$ where E is the number of edges.

```
struct EulerWalk {
  int n;
  vector<multiset<int>> G;
  vector<int> deg;
  EulerWalk(int n) : n(n), G(n + 1), deg(n + 1, 0) {}
  void AddEdge(int a, int b) {
   G[b].insert(a);
   deg[a] += 1; deg[b] -= 1;
   // G[a].insert(b); (*)
 vector<int> walk:
  void dfs(int node) {
    while (G[node].size()) {
      auto vec = *G[node].begin();
     G[node].erase(G[node].begin());
      // G[vec]. erase(G[vec]. find(node)); (*)
     dfs(vec);
    walk.push_back(node);
  template<typename Callback>
  void Solve (Callback cb) {
    for (int i = 1; i <= n; ++i) {</pre>
      while (deg[i] < 0) AddEdge(i, n); // (*)</pre>
      while (deg[i] > 0) AddEdge(n, i); // (*)
      // if (deg[i] \% 2) AddEdge(i, n); (*)
    // Paths
    vector<int> buff; dfs(n);
    for (auto node : walk) {
     if (node < n) buff.push_back(node);</pre>
      else if (buff.size()) {
        cb(buff); buff.clear();
```

```
// Cycles
    for (int i = 0; i < n; ++i) {</pre>
     walk.clear(); dfs(i);
      if (walk.size() > 1) cb(walk);
};
```

Network flow

```
DinicFlow.h
```

Description: Quick flow algorithm.

```
Time: \mathcal{O}(V^2 * E) or \mathcal{O}(E * sqrt(E)) on unit graphs
```

```
struct Dinic {
 struct Edge { int to, cap, flow, nxt; };
 vector<Edge> edges;
 vector<int> graph, at, dist;
 int src = 0, dest = 1;
 Dinic(int n): graph(n, -1), dist(n, -1) {}
 void add_edge(int from, int to, int cap) {
    edges.push_back(Edge {to, cap, 0, graph[from]});
   graph[from] = edges.size() - 1;
 void AddEdge(int from, int to, int cap) {
   add edge(from, to, cap);
   add_edge(to, from, 0);
 bool bfs() {
   queue<int> q;
   fill(dist.begin(), dist.end(), -1);
   dist[src] = 0; q.push(src);
    while (!q.empty()) {
     int node = q.front(); q.pop();
     for (int i = graph[node]; i >= 0; i = edges[i].nxt) {
       const auto &e = edges[i];
       if (dist[e.to] == -1 && e.flow < e.cap) {</pre>
         dist[e.to] = dist[node] + 1;
         q.push(e.to);
   return dist[dest] != -1;
 int dfs(int node, int flow) {
   if (flow == 0) return 0;
   if (node == dest) return flow;
    while (at[node] != -1) {
     int eid = at[node]; const auto &e = edges[eid];
     if (dist[e.to] == dist[node] + 1) {
       if (int ret = dfs(e.to, min(flow, e.cap - e.flow))) {
         edges[ eid ].flow += ret;
         edges[eid^1].flow -= ret;
         return ret;
     at[node] = e.nxt;
```

```
return 0;
  int Compute(int src, int dest) {
    this->src = src; this->dest = dest; int ret = 0;
    while (bfs()) {
      at = graph;
      while (int flow = dfs(src, 2e9))
        ret += flow;
    return ret;
};
```

EZFlow.h

Description: A slow, albeit very easy-to-implement flow algorithm. **Time:** $\mathcal{O}(EF)$ where E is the number of edges and F is the maximum flow.

```
struct EZFlow {
 vector<vector<int>> G;
  vector<bool> vis;
 int t;
  EZFlow(int n) : G(n), vis(n) {}
  bool dfs(int node) {
    if (node == t) return true;
    vis[node] = true;
    for (auto& vec : G[node])
      if (!vis[vec] && dfs(vec)) {
        G[vec].push back(node);
        swap (vec, G[node].back());
        G[node].pop back();
        return true;
    return false;
  void AddEdge(int a, int b) { G[a].push_back(b); }
  int ComputeFlow(int s, int t) {
    this->t = t; int ans = 0;
    while (dfs(s)) {++ans; fill(vis.begin(), vis.end(), false);}
    return ans;
};
```

MinCostMaxFlow.h

Description: Min-cost max-flow with potentials technique. If costs can be negative, call SetPi before Compute, but note that negative cost cycles are not allowed (that's NP-hard). To obtain the actual flow, look at positive values only.

```
Time: Approximately \mathcal{O}(E^2). Another upper bound is \mathcal{O}(FE \log E)
<br/>
<br/>
<br/>
dits/stdc++.h>, <bits/extc++.h>
```

```
using T = int;
const T kInf = numeric limits<T>::max() / 4;
struct MFMC {
 struct Edge { int to, nxt; T flow, cap, cost; };
 vector<Edge> edges;
 vector<T> dist, pi;
 vector<int> par, graph;
 MFMC(int n) :
```

CycleCancelFlow MinCut GlobalMinCut GomoryHu

```
n(n), dist(n), pi(n, 0), par(n), graph(n, -1) {}
void _addEdge(int from, int to, T cap, T cost) {
 edges.push_back(Edge{to, graph[from], 0, cap, cost});
  graph[from] = edges.size() - 1;
void AddEdge(int from, int to, T cap, T cost) {
  _addEdge(from, to, cap, cost);
  _addEdge(to, from, 0, -cost);
bool dijkstra(int s, int t) {
  fill(dist.begin(), dist.end(), kInf);
  fill(par.begin(), par.end(), -1);
  __gnu_pbds::priority_queue<pair<T, int>> q;
  vector<decltype(q)::point_iterator> its(n);
  dist[s] = 0; q.push({0, s});
  while (!q.empty()) {
   int node; T d;
   tie(d, node) = q.top(); q.pop();
   if (dist[node] != -d) continue;
    for (int i = graph[node]; i >= 0; ) {
     const auto &e = edges[i];
     T now = dist[node] + pi[node] - pi[e.to] + e.cost;
     if (e.flow < e.cap && now < dist[e.to]) {</pre>
       dist[e.to] = now;
       par[e.to] = i;
       if (its[e.to] == q.end())
         its[e.to] = q.push({-dist[e.to], e.to});
        else q.modify(its[e.to], {-dist[e.to], e.to});
      i = e.nxt;
  for (int i = 0; i < n; ++i)
 pi[i] = min(pi[i] + dist[i], kInf);
 return par[t] != -1;
pair<T, T> Compute(int s, int t) {
 T flow = 0, cost = 0;
  while (dijkstra(s, t)) {
   T \text{ now = kInf};
   for (int node = t; node != s; ) {
     int ei = par[node];
     now = min(now, edges[ei].cap - edges[ei].flow);
     node = edges[ei ^ 1].to;
    for (int node = t; node != s; ) {
     int ei = par[node];
     edges[ei].flow += now;
     edges[ei ^ 1].flow -= now;
     cost += edges[ei].cost * now;
     node = edges[ei ^ 1].to;
    flow += now;
  return {flow, cost};
// If some costs can be negative, call this before maxflow:
void SetPi(int s) { // (otherwise, leave this out)
  fill(pi.begin(), pi.end(), kInf); pi[s] = 0;
  int it = n, ch = 1; T v;
 while (ch-- && it--)
    for (int i = 0; i < n; ++i) if (pi[i] != kInf)</pre>
```

```
for (int ei = graph[i]; ei >= 0; ) {
          const auto& e = edges[ei];
          if (e.cap && (v = pi[i] + e.cost) < pi[e.to])</pre>
           pi[e.to] = v, ch = 1;
          ei = e.nxt;
    assert(it >= 0); // negative cost cycle
};
```

CvcleCancelFlow.h

Description: Cycle-cancelling algorithm for minimum cost circulation or minimum cost flow (uncomment lines for flow and remove for in iterate) lines

```
struct CCFlow {
 const int kInf = 1e9;
 struct Edge {
   int to, f, c, k;
   int res() { return c - f; }
 vector<Edge> es;
 vector<vector<int>> G;
 long long cost;
 CCFlow(int n) : G(n), in(n), dist(n) {}
 void add_edge(int a, int b, int c, int k) {
   G[a].push back(es.size());
   es.push_back(Edge{b, 0, c, k});
 void AddEdge(int a, int b, int c, int k) {
   add_edge(a, b, c, k);
   add_edge(b, a, 0, -k);
 int start, aug;
 vector<int> in;
 vector<long long> dist;
 int dfs(int node, int f) {
   if (in[nodel) {
     start = node;
     aug = f;
     return 1;
   in[node] = true;
   for (auto ei : G[node]) {
     auto& e = es[ei];
     if (dist[e.to] <= dist[node] + e.k or e.res() == 0)</pre>
       continue:
     dist[e.to] = dist[node] + e.k;
     int rec = dfs(e.to, min(f, e.res()));
     if (rec == 2) return 2;
     if (rec == 1) {
       es[ei].f += aug; es[ei ^ 1].f -= aug;
       cost += 1LL * aug * es[ei].k;
       return 1 + (node == start);
   in[node] = false;
   return 0;
 bool iterate() {
   bool ok = false;
   for (int s = 0; s < n; ++s) {
```

```
fill(in.begin(), in.end(), 0);
      fill(dist.begin(), dist.end(), 1LL * kInf * kInf);
      dist[s] = 0;
      ok \mid = dfs(s, kInf);
    return ok;
 long long Solve() {
    // AddEdge(t, s, kInf, -kInf);
    while (iterate());
    // int flow = es[G[t].back()].f;
    // cost += 1LL * flow * kInf;
    return cost;
};
```

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)
<bits/stdc++.h>
                                                            40 lines
using T = long long;
pair<T, vector<int>> GetMinCut(vector<vector<T>> weights) {
  int n = weights.size();
 vector<int> used(n), best_cut, cut;
 T best weight = numeric limits<T>::max();
  for (int phase = n - 1; phase > 0; phase--) {
    auto w = weights[0];
    auto added = used;
    int prev, k = 0;
    for (int i = 0; i < phase; ++i) {</pre>
      prev = k; k = -1;
      for (int j = 1; j < n; ++j)
        if (!added[j] && (k == -1 || w[j] > w[k]))
          k = j;
      if (i != phase - 1) {
        for (int j = 0; j < n; ++j)
          w[j] += weights[k][j];
        added[k] = true;
        continue;
      for (int j = 0; j < n; ++j)
          weights[prev][j] += weights[k][j];
      for (int j = 0; j < n; ++j)
          weights[j][prev] = weights[prev][j];
      used[k] = true; cut.push_back(k);
      if (w[k] < best_weight) {</pre>
        best_cut = cut;
        best weight = w[k];
 return {best_weight, best_cut};
```

GomoryHu.h

Description: Computes Gomory-Hu tree of a graph **Time:** $\mathcal{O}(V)$ calls of flow algorithm

14 lines

7.3 Matching

DFSMatching.h

Description: This is a simple matching algorithm but should be just fine in most cases. n is the size of the left partition and m is the size of the right partition. If you want to get the matched pairs, match[i] contains match for vertex i on the right side or -1 if it's not matched. Performance can be sigmificantly improved by splitting the condition at (*) and/or by randomizing the order at (**).

Time: $\mathcal{O}\left(EV\right)$ where E is the number of edges and V is the number of vertices

```
<bits/stdc++.h>
struct BipartiteMatcher {
 int n, m;
  vector<vector<int>> G;
  vector<int> L, R, vis;
  BipartiteMatcher(int n, int m) :
   n(n), m(m), G(n), L(n, -1), R(m, -1), vis(n) {}
  void AddEdge(int a, int b) {
   G[a].push_back(b);
  bool match (int node) {
    if (vis[node])
     return false:
    vis[node] = true;
    for (auto vec : G[node]) {
     if (R[vec] == -1 || match(R[vec])) { // (*)
       L[node] = vec; R[vec] = node; return true;
   return false;
  int Solve() {
    int ok = 1;
    while (ok--) {
      fill(vis.begin(), vis.end(), 0);
     for (int i = 0; i < n; ++i) // (**)
       if (L[i] == -1)
          ok \mid = match(i);
    return n - count (L.begin(), L.end(), -1);
```

```
// Only include if you want vertex cover
vector<bool> CL, CR;

void cover(int node) {
   for (auto vec : G[node]) if (!CR[vec]) {
      CR[vec] = true;
      CL[R[vec]] = false;
      cover(R[vec]);
   }
}

int VertexCover() {
   int ret = Solve();
   CL.assign(n, false); CR.assign(m, false);
   for (int i = 0; i < n; ++i) if (L[i] != -1) CL[i] = true;
   for (int i = 0; i < n; ++i) if (L[i] == -1) cover(i);
   return ret;
}
</pre>
```

WeightedMatching.h

auq:

Description: Min cost perfect bipartite matching. Negate costs for max cost.

```
Time: \mathcal{O}(N^3)
template<typename T>
int MinAssignment(const vector<vector<T>> &c) {
                                            // assert(n \le m);
 int n = c.size(), m = c[0].size();
                                            // v: potential
 vector<T> v(m), dist(m);
 vector<int> L(n, -1), R(m, -1);
                                            // matching pairs
 vector<int> index(m), prev(m);
 iota(index.begin(), index.end(), 0);
 auto residue = [&](int i, int j) { return c[i][j] - v[j]; };
 for (int f = 0; f < n; ++f) {
   for (int j = 0; j < m; ++j) {
     dist[j] = residue(f, j); prev[j] = f;
   T w; int j, 1;
   for (int s = 0, t = 0;;) {
     if (s == t) {
       l = s; w = dist[index[t++]];
       for (int k = t; k < m; ++k) {
         j = index[k]; T h = dist[j];
         if (h <= w) {
           if (h < w) { t = s; w = h; }
            index[k] = index[t]; index[t++] = j;
       for (int k = s; k < t; ++k) {
         j = index[k];
         if (R[j] < 0) goto aug;
     int q = index[s++], i = R[q];
     for (int k = t; k < m; ++k) {
       j = index[k];
       T h = residue(i, j) - residue(i, q) + w;
       if (h < dist[j]) {
         dist[j] = h; prev[j] = i;
         if (h == w) {
           if (R[j] < 0) goto aug;
            index[k] = index[t]; index[t++] = j;
```

```
for(int k = 0; k < 1; ++k)
    v[index[k]] += dist[index[k]] - w;
int i;
do {
    R[j] = i = prev[j];
    swap(j, L[i]);
} while (i != f);
}
T ret = 0;
for (int i = 0; i < n; ++i) {
    ret += c[i][L[i]]; // (i, L[i]) is a solution
}
return ret;</pre>
```

GeneralMatching.h

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

```
"../numerical/MatrixInverse-mod.h"
                                                            40 lines
vector<pii> generalMatching(int N, vector<pii>& ed) {
 vector<vector<ll>> mat(N, vector<ll>(N)), A;
 trav(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<11>(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]) {
       ll b = A[i][fj] * a % mod;
        rep(j, 0, M) A[i][j] = (A[i][j] - A[fi][j] * b) % mod;
      swap(fi,fj);
 return ret;
```

7.4 DFS algorithms

SCC.h

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

Usage: SCC(graph, [&](vi& v) $\{ \dots \}$) visits all components in reverse topological order. comp[i] holds the component

BiconnectedComponents 2sat LCA

```
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)
vector<int> val, comp, stk, cont;
int timer, ncomps;
template < class Graph, class Func>
int dfs(int node, Graph& G, Func f) {
  int low = val[node] = ++timer, x; stk.push_back(node);
  for (auto vec : G[node]) if (comp[vec] < 0)</pre>
    low = min(low, val[vec] ?: dfs(vec, G, f));
  if (low == val[node]) {
    do √
     x = stk.back(); stk.pop_back();
     comp[x] = ncomps;
     cont.push_back(x);
    } while (x != node);
    f(cont); cont.clear();
    ncomps++;
  return val[node] = low;
template < class Graph, class Func>
void SCC(Graph& G, Func f) {
  int n = G.size();
  val.assign(n, 0); comp.assign(n, -1);
  timer = ncomps = 0;
  for (int i = 0; i < n; ++i)
   if (comp[i] < 0)
      dfs(i, G, f);
```

BiconnectedComponents.h

Description: Finds all biconnected components in an undirected multigraph, 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. HOWEVER, note that we are outputting bridges as BCC's here, because we might be interested in vertex bcc's, not edge bcc's.

To get the articulation points, look for vertices that are in more than 1 BCC. To get the bridges, look for biconnected components with one edge

```
Time: \mathcal{O}(E+V)
                                                            54 lines
struct BCC {
  vector<pair<int, int>> edges;
  vector<vector<int>> G;
  vector<int> enter, low, stk;
  BCC(int n) : G(n), enter(n, -1) {}
  int AddEdge(int a, int b) {
    int ret = edges.size();
    edges.emplace back(a, b);
    G[a].push_back(ret);
   G[b].push_back(ret);
    return ret;
  template<typename Iter>
  void Callback (Iter bg, Iter en) {
    for (Iter it = bg; it != en; ++it) {
      auto edge = edges[*it];
      // Do something useful
```

```
void Solve() {
   for (int i = 0; i < (int)G.size(); ++i)</pre>
     if (enter[i] == -1) {
       dfs(i, -1);
 int timer = 0;
 int dfs(int node, int pei) {
   enter[node] = timer++;
    int ret = enter[node];
    for (auto ei : G[node]) if (ei != pei) {
     int vec = (edges[ei].first ^ edges[ei].second ^ node);
     if (enter[vec] != -1) {
       ret = min(ret, enter[vec]);
       if (enter[vec] < enter[node])</pre>
         stk.push_back(ei);
     } else {
       int sz = stk.size(), low = dfs(vec, ei);
       ret = min(ret, low);
       stk.push back(ei);
       if (low >= enter[node]) {
         Callback(stk.begin() + sz, stk.end());
         stk.resize(sz);
   return ret;
};
```

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 $(a \lor b) \land (la \lor c) \land (d \lor lb) \land \ldots$ becomes true, or reports that it is unsatisfiable. THROWS 5 IF NO SOLUTION Negated variables are represented by bit-inversions $(\sim x)$. Usage: TwoSat sat(4); // number of variables sat.Either(0, ~ 3); // Var 0 is true or var 3 is false sat.SetValue(2); // Var 2 is true sat.AtMostOne($\{0, \sim 1, 2\}$); // <= 1 of vars 0, \sim 1 and 2 are true sat.Solve(); // Returns solution or throws

Time: $\mathcal{O}\left(N+E\right)$, where N is the number of boolean variables, and E is the number of clauses.

```
struct TwoSat {
  int n:
  vector<vector<int>> G:
  vector<int> values; // 0 = false, 1 = true
  TwoSat(int n = 0) : n(n), G(2*n) {}
  int AddVar() { // (optional)
   G.emplace_back();
   G.emplace_back();
    return n++;
 void Implies(int a, int b) {
   a = (a >= 0 ? 2*a : -1-2*a);
   b = (b >= 0 ? 2*b : -1-2*b);
   G[a].push_back(b);
 void Either(int a, int b) {
    Implies (\sim a, b);
    Implies(~b, a);
```

```
void SetValue(int x) {
   Either (x, x);
 void AtMostOne(const vector<int>& vals) { // (optional)
   if (vals.size() <= 1) return;</pre>
   int cur = ~vals[0];
   for (int i = 2; i < (int)vals.size(); ++i) {</pre>
     int nxt = AddVar();
     Either(cur, ~vals[i]);
     Either(cur, nxt);
     Either(~vals[i], nxt);
     cur = ~nxt;
   Either(cur, ~vals[1]);
 vector<int> enter, comp, stk;
 int timer = 0;
 int dfs(int node) {
   int low = enter[node] = ++timer, x;
   stk.push back(node);
    for (auto vec : G[node]) if (!comp[vec])
     low = min(low, enter[vec] ?: dfs(vec));
    ++timer;
   if (low == enter[node]) do {
     x = stk.back(); stk.pop_back();
     comp[x] = timer;
     if (values[x >> 1] == -1)
     values[x >> 1] = 1 - x & 1;
   } while (x != node);
   return enter[node] = low;
 vector<int> Solve() {
   values.assign(n, -1);
   enter.assign(2 * n, 0); comp = enter;
   for (int i = 0; i < 2 * n; ++i) {
     if (!comp[i])
       dfs(i);
    for (int i = 0; i < n; ++i) {</pre>
     if (comp[2 * i] == comp[2 * i + 1])
       throw 5;
   return values;
};
```

7.5 Trees

LCA.h

Description: Lowest common ancestor. Finds the lowest common ancestor in a tree (with 0 as root). C should be an adjacency list of the tree, either directed or undirected. Can also find the distance between two nodes.

CompressTree HLD LinkCutTree

```
RMQ<pair<int, int>> rmq;
  int timer = 0;
  LCA(int n) : enter(n, -1), depth(n), G(n), linear(2 * n) {}
  void dfs (int node, int dep) {
   linear[timer] = {dep, node};
    enter[node] = timer++;
   depth[node] = dep;
    for (auto vec : G[node])
    if (enter[vec] == -1) {
     dfs(vec, dep + 1);
     linear[timer++] = {dep, node};
  void AddEdge(int a, int b) {
   G[a].push_back(b);
   G[b].push_back(a);
  void Build(int root) {
   dfs(root, 0);
    rmq.Build(linear);
  int Query(int a, int b) {
   a = enter[a], b = enter[b];
   return rmq.Query(min(a, b), max(a, b) + 1).second;
  int Distance(int a, int b) {
   return depth[a] + depth[b] - 2 * depth[Query(a, b)];
};
```

CompressTree.h

HLD.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 the nodes of the reduced tree, while at the same time populating a link array that stores the new parents. The root points to -1.

```
Time: \mathcal{O}(|S| * (\log |S| + LCA_{-}Q))
```

```
18 lines
vector<int> CompressTree(vector<int> v, LCA& lca,
                         vector<int>& link) {
  auto cmp = [&](int a, int b) {
   return lca.enter[a] < lca.enter[b];</pre>
  sort(v.begin(), v.end(), cmp);
  v.erase(unique(v.begin(), v.end()), v.end());
  for (int i = (int) v.size() - 1; i > 0; --i)
   v.push_back(lca.Query(v[i - 1], v[i]));
  sort(v.begin(), v.end(), cmp);
  v.erase(unique(v.begin(), v.end()), v.end());
 for (int i = 0; i < (int)v.size(); ++i)</pre>
   link[v[i]] = (i == 0 ? -1 : lca.Query(v[i - 1], v[i]));
  return v;
```

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.

```
struct HeavyLight {
 struct Node {
   int jump, subsize, depth, lin, parent;
   vector<int> leg;
 vector<Node> T;
 bool processed = false;
 HeavyLight(int n) : T(n) {}
 void AddEdge(int a, int b) {
   T[a].leg.push_back(b);
   T[b].leg.push_back(a);
 void Preprocess() {
   dfs_sub(0, -1); dfs_jump(0, 0);
   processed = true;
  // Gets the position in the HL linearization
 int GetPosition(int node) {
   assert (processed);
   return T[node].lin;
 // Gets an array of ranges of form [li...ri]
 // that correspond to the ranges you would need
 // to query in the underlying structure
 vector<pair<int, int>> GetPathRanges(int a, int b) {
   assert (processed);
   vector<pair<int, int>> ret;
    while (T[a].jump != T[b].jump) {
     if (T[T[a].jump].depth < T[T[b].jump].depth)</pre>
     swap(a, b);
      ret.emplace_back(T[T[a].jump].lin, T[a].lin + 1);
     a = T[T[a].jump].parent;
   if (T[a].depth < T[b].depth) swap(a, b);</pre>
   ret.emplace back(T[b].lin, T[a].lin + 1);
   return ret;
 int dfs_sub(int x, int par) {
   auto &node = T[x];
   node.subsize = 1; node.parent = par;
   if (par !=-1) {
     node.leg.erase(find(node.leg.begin(),
                          node.leg.end(), par));
     node.depth = 1 + T[par].depth;
   for (auto vec : node.leg)
     node.subsize += dfs_sub(vec, x);
   return node.subsize;
 int timer = 0;
 void dfs_jump(int x, int jump) {
   auto &node = T[x];
   node.jump = jump; node.lin = timer++;
   iter_swap(node.leg.begin(), max_element(node.leg.begin(),
     node.leg.end(), [&](int a, int b) {
       return T[a].subsize < T[b].subsize;</pre>
   }));
```

```
for (auto vec : node.leg)
      dfs_jump(vec, vec == node.leg.front() ? jump : vec);
};
```

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).
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 push flip() {
    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 \hat{b};
    Node *x = c[i], *y = b == 2 ? x : x -> c[h], *z = b ? y : x;
    if ((y->p = p)) p->c[up()] = y;
    c[i] = z -> c[i ^ 1];
    if (b < 2) {
      x - c[h] = y - c[h ^ 1];
      z \rightarrow c[h ^1] = b ? x : this:
    v - > c[i ^1] = b ? this : x;
    fix(); x->fix(); y->fix();
    if (p) p->fix();
    swap(pp, y->pp);
 void splay() {
    for (push_flip(); p; ) {
      if (p->p) p->p->push_flip();
      p->push_flip(); push_flip();
      int c1 = up(), c2 = p->up();
      if (c2 == -1) p->rot(c1, 2);
      else p->p->rot(c2, c1 != c2);
 Node* first() {
    push_flip();
    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));
    make_root(&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];
    make_root(top); x->splay();
```

```
assert(top == (x->pp ?: x->c[0]));
    if (x->pp) x->pp = 0;
    else {
     x->c[0] = top->p = 0;
      x->fix();
  bool connected (int u, int v) { // are u, v in the same tree?
   Node* nu = access(&node[u])->first();
    return nu == access(&node[v])->first();
  void make_root(Node* u) {
    access(u);
    u->splay();
    if(u->c[0]) {
     u - > c[0] - > p = 0;
     u - c[0] - flip ^= 1;
     u - c[0] - pp = u;
     u - > c[0] = 0;
     u->fix():
 Node* access(Node* u) {
   u->splay();
    while (Node* pp = u->pp) {
      pp \rightarrow splay(); u \rightarrow pp = 0;
      if (pp->c[1]) {
       pp - c[1] - p = 0; pp - c[1] - pp = pp; 
      pp - c[1] = u; pp - fix(); u = pp;
    return u;
};
```

7.6 Matrix tree theorem

MatrixTree.h

Description: To count the number of spanning trees in an undirected graph G: create an $N \times N$ matrix mat, and for each edge $(a,b) \in G$, do mat[a][a]++, mat[b][b]++, mat[a][b]--, mat[b][a]--. Remove the last row and column, and take the determinant.

Strings (8)

KMP.h

Description: pi[x] is the length of the longest prefix of s that ends at x (exclusively), other than s[0..x) itself. This is used by Match() to find all occurences of a string.

```
vector<int> ComputePi(string s) {
  int n = s.size();
  vector<int> pi(n + 1, -1);

for (int i = 0; i < n; ++i) {
  int j = pi[i];
  while (j!= -1 && s[j]!= s[i]) j = pi[j];
  pi[i + 1] = j + 1;
}</pre>
```

```
return pi;
vector<int> Match(string text, string pat) {
 vector<int> pi = ComputePi(pat), ret;
 int j = 0;
 for (int i = 0; i < (int)text.size(); ++i) {</pre>
    while (j != -1 && pat[j] != text[i]) j = pi[j];
    if (++; == pat.size())
      ret.push_back(i), j = pi[j];
 return ret;
ZFunction.h
Description: Given a string s, computes the length of the longest common
prefix of s[i..] and s[0..] for each i > 0!!
Usage: Zfunction("abacaba") => {0, 0, 1, 0, 3, 0, 1}
Time: \mathcal{O}(N)
                                                             21 lines
<bits/stdc++.h>
vector<int> ZFunction(string s) {
 int n = s.size();
 vector<int> z(n, 0);
 int L = 0, R = 0;
 for (int i = 1; i < n; i++) {</pre>
   if (i > R) {
     L = R = i;
      while (R < n \&\& s[R - L] == s[R]) R++;
      z[i] = R - L; R--;
    } else {
      int k = i-L;
      if (z[k] < R - i + 1) z[i] = z[k];
      else {
       L = i;
        while (R < n \&\& s[R - L] == s[R]) R++;
        z[i] = R - L; R--;
 return z;
```

Manacher.h

Description: Given a string s, computes the length of the longest palindromes centered in each position (for parity == 1) or between each pair of adjacent positions (for parity == 0).

```
Usage: Manacher("abacaba", 1) => \{0, 1, 0, 3, 0, 1, 0\} Manacher("aabbaa", 0) => \{1, 0, 3, 0, 1\} Time: \mathcal{O}(N)
```

PalindromicTree.h

Description: A trie-like structure for keeping track of palindromes of a string s. It has two roots, 0 (for even palindromes) and 1 (for odd palindromes). Each node stores the length of the palindrome, the count and a link to the longest "aligned" subpalindrome. Can be made online from left to right

```
Time: \mathcal{O}(N)
struct PalTree {
  struct Node {
    map<char, int> leg;
    int link, len, cnt;
  vector<Node> T;
  int nodes = 2;
  PalTree(string str) : T(str.size() + 2) {
    T[1].link = T[1].len = 0;
    T[0].link = T[0].len = -1;
    int last = 0;
    for (int i = 0; i < (int)str.size(); ++i) {</pre>
      char now = str[i];
      int node = last;
      while (now != str[i - T[node].len - 1])
        node = T[node].link;
      if (T[node].leg.count(now)) {
        node = T[node].leg[now];
        T[node].cnt += 1;
        last = node;
        continue;
      int cur = nodes++;
      T[cur].len = T[node].len + 2;
      T[node].leg[now] = cur;
      int link = T[node].link;
      while (link !=-1) {
        if (now == str[i - T[link].len - 1] &&
            T[link].leg.count(now)) {
          link = T[link].leg[now];
          break;
        link = T[link].link;
      if (link <= 0) link = 1;
      T[cur].link = link;
      T[cur].cnt = 1;
      last = cur;
    for (int node = nodes - 1; node > 0; --node) {
      T[T[node].link].cnt += T[node].cnt;
};
```

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 = s.size(); s += s;
  for (int b = 0; b < n; ++b)</pre>
```

SuffixArray SuffixAutomaton AhoCorasick

```
for (int i = 0; i < n; ++i) {</pre>
   if (a + i == b || s[a + i] < s[b + i]) {
     b += max(0, i - 1); break;
   if (s[a + i] > s[b + i]) { a = b; break; }
return a;
```

Suffix Array.h

Description: Builds suffix array for a string. a[i] is the starting index of the suffix which is i-th in the sorted suffix array. The lop function calculates longest common prefixes for indices. Can also sort cyclic permutations Memory: $\mathcal{O}(N)/\mathcal{O}(N \log N)$

Time: $\mathcal{O}(N \log N)$ where N is the length of the string for creation of the SA. $\mathcal{O}(log)$ for LCP.

```
struct SuffixArray {
 int n. csz;
  vector<vector<int>> classes;
  vector<int> cnt, order, oldc, newc, left;
  string str;
  SuffixArray(string s, bool cyclic) :
   n(s.size() + !cyclic), csz(max(n, 256)), cnt(csz),
   order(n), oldc(n), newc(n), left(n), str(s) {
     if (!cyclic) str += '\0';
  vector<int> Build() {
    for (int i = 0; i < n; ++i) {</pre>
     oldc[i] = newc[i] = str[i];
     order[i] = left[i] = i;
    for (int step = 1; step <= 2 * n; step *= 2) {</pre>
     // Counting sort (can be replaced by sort with left)
      // although not trivial
     fill(cnt.begin(), cnt.end(), 0);
      for (int i = 0; i < n; ++i) ++cnt[oldc[left[i]]];</pre>
      for (int i = 1; i < csz; ++i) cnt[i] += cnt[i - 1];</pre>
      for (int i = n - 1; i >= 0; --i)
       order[--cnt[oldc[left[i]]]] = left[i];
     newc[order[0]] = 0;
      for (int i = 1; i < n; ++i) {</pre>
       int now1 = order[i], last1 = order[i - 1],
            now2 = (now1 + step / 2) % n,
            last2 = (last1 + step / 2) % n;
        newc[now1] = newc[last1] + (oldc[now1] != oldc[last1]
               or oldc[now2] != oldc[last2]);
      classes.push_back(newc);
     swap(oldc, newc);
      for (int i = 0; i < n; ++i) {</pre>
       left[i] = (order[i] + n - step) % n;
    return order;
  int Compare(int i, int j, int len) {
   for (int step = 0; len; ++step, len /= 2) {
```

if (len % 2 == 0) continue;

```
int ret = classes[step][i] - classes[step][j];
     if (ret != 0) return ret < 0 ? -1 : 1;</pre>
     i = (i + (1 << step)) % n;
     j = (j + (1 << step)) % n;
    return 0;
 int GetLCP(int i, int j) {
   if (i == j) return str.back() == '\0' ? n - i - 1 : n;
    int ans = 0;
    for (int step = classes.size() - 1; step >= 0; --step) {
     if (classes[step][i] == classes[step][i]) {
       i = (i + (1 << step)) % n;
       j = (j + (1 << step)) % n;
       ans += (1 << step);
   return min(ans, n); // if cyclic
};
```

SuffixAutomaton.h

Description: Builds an automaton of all the suffixes of a given string (online from left to right). For each character c, do sa.ConsumeChar(c) You can change char to int and add negative numbers to support multiple strings.

Time: $\mathcal{O}(log(sigma))$ amortized per character added

dits/stdc++.h>

```
struct SuffixAutomaton {
 struct Node {
   int link, len;
   map<char, int> leg;
 };
 vector<Node> T;
 int last = 0, nodes = 1;
 SuffixAutomaton(int sz) : T(2 * sz + 1) {
   T[0].link = -1;
   T[0].len = 0;
 // Adds another character to the automaton
 // and returns the node of the whole new string
 // (the suffixes of that are parents in the link tree)
 int ConsumeChar(char c) {
    // Add state for whole string
   int cur = nodes++, node = last;
   T[cur].len = T[last].len + 1;
   T[cur].link = 0;
    // Add transitions to all suffixes which do not have one
    // already
    while (node != -1 && T[node].leg.count(c) == 0) {
     T[node].leg[c] = cur;
     node = T[node].link;
   if (node != −1) {
     int old = T[node].leg[c];
     if (T[old].len == T[node].len + 1) {
       T[cur].link = old;
     } else {
       int clone = nodes++;
       T[clone].leg = T[old].leg;
```

```
T[clone].len = T[node].len + 1;
       T[clone].link = T[old].link;
       T[old].link = T[cur].link = clone;
       while (node != -1 && T[node].leg[c] == old) {
         T[node].leg[c] = clone;
         node = T[node].link;
   return last = cur;
  // Runs through the automaton
 int Go(int node, char c) {
   while (node != -1 && T[node].leg.count(c) == 0)
     node = T[node].link;
   return (node == -1 ? 0 : T[node].leg[c]);
};
```

58 lines

```
AhoCorasick.h
Description: Aho-Corasick algorithm builds an automaton for multiple
pattern string matching
Time: \mathcal{O}(N * log(sigma)) where N is the total length
<br/>
<br/>
dits/stdc++.h>
                                                             48 lines
struct AhoCorasick {
  struct Node {
    int link;
    map<char, int> leg;
  vector<Node> T;
  int root = 0, nodes = 1;
  AhoCorasick(int sz) : T(sz) {}
  // Adds a word to trie and returns the end node
  int AddWord(const string &word) {
    int node = root;
    for (auto c : word) {
      auto &nxt = T[node].leg[c];
      if (nxt == 0) nxt = nodes++;
      node = nxt;
    return node;
  // Advances from a node with a character (like an automaton)
  int Advance(int node, char chr) {
    while (node != -1 && T[node].leg.count(chr) == 0)
      node = T[node].link;
    if (node == -1) return root;
    return T[node].leg[chr];
  // Builds links
  void BuildLinks() {
    queue<int> 0;
    Q.push (root);
    T[root].link = -1;
    while (!Q.empty()) {
      int node = 0.front();
      Q.pop();
      for (auto &p : T[node].leg) {
        int vec = p.second;
        char chr = p.first;
```

```
T[vec].link = Advance(T[node].link, chr);
        Q.push (vec);
};
```

Various (9)

9.1 Intervals

IntervalContainer.h

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

```
Time: \mathcal{O}(\log N)
struct IntervalContainer {
  map<int, int> s;
  using Iter = map<int, int>::iterator;
  Iter AddInterval(int 1, int r) {
   if (1 == r) return s.end();
   Iter it = s.lower bound(1);
   while (it != s.end() && it->first <= r) {
     r = max(r, it->second);
     it = s.erase(it);
    while (it != s.begin() && (--it)->second >= 1) {
     l = min(l, it->first);
     r = max(r, it->second);
     it = s.erase(it);
   return s.insert({1, r}).first;
  Iter FindInterval(int x) {
   auto it = s.upper bound(x);
   if (it == s.begin() or (--it)->second <= x)
     return s.end();
    return it:
  void RemoveInterval(int 1, int r) {
   if (1 == r) return;
```

IntervalCover.h

};

s.erase(it);

auto it = AddInterval(1, r);

if (1 != 12) s.insert({12, 1});

if (r != r2) s.insert({r, r2});

int 12 = it->first, r2 = it->second;

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

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

```
template < class T>
vi cover(pair<T, T> G, vector<pair<T, T>> I) {
 vi S(sz(I)), R;
  iota(all(S), 0);
  sort(all(S), [&](int a, int b) { return I[a] < I[b]; });</pre>
 T cur = G.first;
  int at = 0;
```

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

ConstantIntervals.h

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

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

```
template < class Func, class Callback, class T>
void recurse (int from, int to, Func f, Callback cb,
            int& i, T& p, T q) {
  if (p == q) return;
  if (from == to) {
    cb(i, to, p);
   i = to; p = q;
  } else {
    int mid = (from + to) / 2;
    recurse(from, mid, f, cb, i, p, f(mid));
   recurse (mid + 1, to, f, cb, i, p, q);
template < class Func, class Callback >
void ConstantIntervals(int from, int to, Func f, Callback cb) {
 if (to <= from) return;</pre>
 int i = from; auto p = f(i), q = f(to - 1);
 recurse(from, to - 1, f, cb, i, p, q);
 cb(i, to, q);
```

Misc. algorithms 9.2

TernarySearch.h

Time: $\mathcal{O}(\log(b-a))$

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 = TernarySearch(0, n-1, [&](int i) {return a[i];});

```
template < class Func>
int TernarySearch(int a, int b, Func 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;
 for (int i = a + 1; i <= b; ++i)</pre>
   if (f(a) < f(i)) a = i; // (B)
 return a:
```

AlphaBeta.h

Description: Uses the alpha-beta pruning method to find score values for states in games (minimax)

```
int AlphaBeta(state s, int alpha, int beta) {
 if (s.finished()) return s.score();
 for (state t : s.next()) {
   alpha = max(alpha, -AlphaBeta(t, -beta, -alpha));
   if (alpha >= beta) break;
 return alpha;
```

9.3 Dynamic programming

DivideAndConquerDP.h

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

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

```
struct DP { // Modify at will:
  int lo(int ind) { return 0; }
 int hi(int ind) { return ind; }
 11 f(int ind, int k) { return dp[ind][k]; }
 void store(int ind, int k, ll v) { res[ind] = pii(k, v); }
  void rec(int L, int R, int LO, int HI) {
    if (L >= R) return;
    int mid = (L + R) \gg 1;
    pair<11, int> best (LLONG_MAX, LO);
    rep(k, max(LO,lo(mid)), min(HI,hi(mid)))
     best = min(best, make_pair(f(mid, k), k));
    store (mid, best.second, best.first);
    rec(L, mid, LO, best.second+1);
    rec(mid+1, R, best.second, HI);
  void solve(int L, int R) { rec(L, R, INT_MIN, INT_MAX); }
```

KnuthDP.h

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

9.4 Debugging tricks

- signal(SIGSEGV, [](int) { _Exit(0); }); converts segfaults into Wrong Answers. Similarly one can catch SIGABRT (assertion failures) and SIGFPE (zero divisions). _GLIBCXX_DEBUG violations 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).

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9.5 Optimization tricks

9.5.1 Bit hacks

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

9.5.2 Pragmas

- #pragma GCC optimize ("Ofast") will make GCC auto-vectorize for loops and optimizes floating points better (assumes associativity and turns off denormals).
- #pragma GCC target ("avx,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).

Unrolling.h

```
5 line
```

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

Unrolling 21

Techniques (A)

techniques.txt

159 lines

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

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

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