

University of Virginia

Ground Zero

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1 Contest 1	001: 2002 0:::::::(14 4/ 00:::0 14 2/ (
2 Mathematics 1	<pre>return b < a ? a = b, 1 : 0; } // set a = min(a,b) tcT> bool ckmax(T& a, const T& b) { return a < b ? a = b, 1 : 0; } // set a = max(a,b)</pre>
3 Data structures 2	<pre>int main() { cin.tie(0)->sync_with_stdio(0); }</pre>
4 Numerical 4	
5 Number theory 8	alias clr="printf '\33c'" co() { g++ -std=c++17 -O2 -Wall -Wextra -Wshadow -Wconversion - → \$1 \$1.cpp; } run() { co \$1 && ./\$1; }
6 Combinatorial 10	hash.sh
7 Graph 11	cpp -dD -P -fpreprocessed tr -d '[:space:]' md5sum cut -c-6
8 Geometry 16	troubleshoot.txt 75 lines
9 Strings 20	General: Write down most of your thoughts, even if you're not sure whether they're useful.
10 Various 21	Give your variables (and files) meaningful names.
$\underline{\text{Contest}}$ (1)	Pre-submit: Write a few simple test cases if sample is not enough.
template.cpp 44 lines	Are time limits close? If so, generate max cases. Is the memory usage fine?
<pre>#include <bits stdc++.h=""> using namespace std;</bits></pre>	Could anything overflow? Remove debug output. Make sure to submit the right file.
<pre>using ll = long long; using db = long double; // or double if tight TL using str = string; using pi = pair<int,int>; #define mp make_pair #define f first #define s second #define tcT template<class t="" tct=""> using V = vector<t>; tcT, size_t SZ> using AR = array<t,sz>; using vi = V<int>; using vi = V<int>; using vpi = V<pi>; #define sz(x) int((x).size()) #define all(x) begin(x), end(x) #define rsz resize #define pb push_back</pi></int></int></t,sz></t></class></int,int></pre>	Wrong answer: Print your solution! Print debug output as well. Read the full problem statement again. Have you understood the problem correctly? Are you sure your algorithm works? Try writing a slow (but correct) solution. Can your algorithm handle the whole range of input? Did you consider corner cases (ex. n=1)? Is your output format correct? (including whitespace) Are you clearing all data structures between test cases? Any uninitialized variables? Any undefined behavior (array out of bounds)? Any overflows or NaNs (or shifting ll by >=64 bits)? Confusing N and M, i and j, etc.? Confusing ++i and i++? Return vs continue vs break? Are you sure the STL functions you use work as you think? Add some assertions, maybe resubmit. Create some test cases to run your algorithm on. Go through the algorithm for a simple case.
#define pb push_back #define ft front() #define bk back() #define FOR(i,a,b) for (int i = (a); i < (b); ++i) #define FOR(i,a) FOR(i,0,a) #define ROF(i,a,b) for (int i = (b)-1; i >= (a);i) #define ROF(i,a) ROF(i,0,a) #define rep(a) FOR(_,a) #define each(a,x) for (auto& a: x) const int MOD = le9+7; const db PI = acos((db)-1); mt19937 rng(0); // or mt19937_64	Go through the algorithm for a simple case. Go through this list again. Explain your algorithm to a teammate. Ask the teammate to look at your code. Go for a small walk, e.g. to the toilet. Rewrite your solution from the start or let a teammate do it. Geometry: Work with ints if possible. Correctly account for numbers close to (but not) zero. Related: for functions like acos make sure absolute val of input is not (slightly) greater than one. Correctly deal with vertices that are collinear, concyclic, coplanar (in 3D), etc. Subtracting a point from every other (but not itself)?

Runtime error: Have you tested all corner cases locally? Any uninitialized variables? Are you reading or writing outside the range of any vector? Any assertions that might fail? Any possible division by 0? (mod 0 for example) Any possible infinite recursion? Invalidated pointers or iterators? Are you using too much memory? Debug with resubmits (e.g. remapped signals, see Various). Time limit exceeded: Do you have any possible infinite loops? What's your complexity? Large TL does not mean that something simple (like NlogN) isn't intended. Are you copying a lot of unnecessary data? (References) Avoid vector, map. (use arrays/unordered_map) How big is the input and output? (consider FastIO) What do your teammates think about your algorithm? Calling count() on multiset? Memory limit exceeded: What is the max amount of memory your algorithm should need? Are you clearing all data structures between test cases? If using pointers try BumpAllocator.

Mathematics (2)

Cramer's Rule: given an equation Ax = b, the solution to a variable x_i is given by

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

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

2.1 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
$$r = \sqrt{a^2 + b^2}, \phi = \text{atan2}(b, a)$$
.

Geometry

2.2.1 Triangles

Side lengths: a, b, c

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

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

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

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

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

Length of bisector (divides angles in two):

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

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

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

2.2.2 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 $A = \sqrt{(p-a)(p-b)(p-c)(p-d)}$.

2.3 Derivatives/Integrals

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

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

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

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

Integration by parts:

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

2.4 Sums

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

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

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

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

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

Data structures (3)

HashMap.h

Description: Hash map with similar API as unordered_map. Initial capacity must be a power of 2 if provided.

Usage: ht<int,int> h({},{},{},{},{1<<16});

Memory: ~1.5x unordered map

Time: ~3x faster than unordered map

<ext/pb_ds/assoc_container.hpp> 5872b2, 9 lines using namespace __gnu_pbds; struct chash {

const uint64_t C = 11(4e18*acos(0))+71; // large odd number const int RANDOM = rng(); 11 operator()(11 x) const { return __builtin_bswap64((x^ \hookrightarrow RANDOM) *C); }

template < class K, class V > using ht = gp_hash_table < K, V, chash >; template < class K, class V > V get (ht < K, V > & u, K x) { auto it = u.find(x); return it == end(u) ? 0 : it->s; }

OrderStatisticTree.h

Description: order_of_key, find_by_order (order = num less)

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

#include <bits/extc++.h> using namespace __gnu_pbds;

template<class T> using Tree = tree<T, null_type, less<T>, rb_tree_tag, tree_order_statistics_node_update>;

LineContainer.h

void add(T a, T b) {

Description: Add lines of the form ax + b, query maximum y-coordinate for any x.

Time: $\mathcal{O}(\log N)$ df7386, 29 lines

using T = 11; const T INF = LLONG_MAX; // a/b rounded down // 11 fdiv(11 a, 11 b) { return a/b-((a^b)<0&&a%b); }

```
bool _{Q} = 0;
struct Line {
 T a, b; mutable T lst;
 T eval(T x) const { return a*x+b; }
 bool operator<(const Line&o)const{return 0?lst<o.lst:a<o.a;}</pre>
 T last_gre(const Line& o) const { assert(a <= o.a);</pre>
    // greatest x s.t. a*x+b >= o.a*x+o.b
    return lst=(a==o.a?(b>=o.b?INF:-INF):fdiv(b-o.b,o.a-a));}
struct LineContainer: multiset<Line> {
```

bool isect(iterator it) { auto n_it = next(it);

if (n_it == end()) return it->lst = INF, 0;

return it->last_gre(*n_it) >= n_it->lst; }

```
auto it = ins({a,b,0}); while (isect(it)) erase(next(it));
  if (it == begin()) return;
  if (isect(--it)) erase(next(it)), isect(it);
  while (it != begin()) {
    --it; if (it->lst < next(it)->lst) break;
    erase(next(it)); isect(it); }
T qmax(T x) { assert(!empty());
  _Q = 1; T res = lb(\{0,0,x\}) \rightarrow eval(x); _Q = 0;
  return res: }
```

LineContainerDeque.h

Description: LineContainer assuming both slopes and queries monotonic. Time: $\mathcal{O}(1)$

```
"LineContainer.h"
struct LCdeque : deque<Line> {
  void addBack(Line L) { // assume nonempty
      auto a = bk; pop_back(); a.lst = a.last_gre(L);
      if (size() && bk.lst >= a.lst) continue;
    L.lst = INF; pb(L);
  void addFront(Line L) {
    while (1) {
      if (!size()) { L.lst = INF; break; }
      if ((L.lst = L.last_gre(ft)) >= ft.lst) pop_front();
    push front (L);
  void add(T a, T b) { // line goes to one end of deque
    if (!size() || a <= ft.a) addFront({a,b,0});</pre>
    else assert(a >= bk.a), addBack({a,b,0});
  int ord = 0; // 1 = x's come in increasing order, -1 =
     ⇔decreasing order
  T query(T x) {
    assert (ord);
    if (ord == 1) {
      while (ft.lst < x) pop_front();</pre>
      return ft.eval(x);
      while (size () >1&&prev (prev (end ())) ->1st>=x) pop_back ();
      return bk.eval(x);
};
```

RMQ.h

cd2981, 6 lines

Description: 1D range minimum query. If TL is an issue, use arrays instead of vectors and store values instead of indices.

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

```
Time: \mathcal{O}(1)
                                                       a3f881, 19 lines
tcT> struct RMQ { // floor(log_2(x))
  int level(int x) { return 31- builtin clz(x); }
  V<T> v; V<vi> jmp;
  int cmb(int a, int b) {
    return v[a] == v[b]?min(a,b):(v[a] < v[b]?a:b); }
  void init(const V<T>& _v) {
    v = _v; jmp = {vi(sz(v))};
    iota(all(jmp[0]),0);
    for (int j = 1; 1 << j <= sz(v); ++j) {
      jmp.pb(vi(sz(v)-(1<< j)+1));
      FOR(i, sz(jmp[j])) jmp[j][i] = cmb(jmp[j-1][i],
        jmp[j-1][i+(1<<(j-1))]);
```

```
}
int index(int 1, int r) { // kat is rex instead
  assert(1 <= r); int d = level(r-l+1);
  return cmb(jmp[d][1],jmp[d][r-(1<<d)+1]); }
T query(int 1, int r) { return v[index(l,r)]; }
::</pre>
```

SegmentTree.h

Description: 1D point update and range query where cmb is any associative operation. seg[1] = query(0, N-1).

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

1630f3, 18 line

```
tcT> struct SegTree { // cmb(ID,b) = b
 const T ID{}; T cmb(T a, T b) { return a+b; }
  int n: V<T> seq;
  void init(int _n) { // upd, query also work if n = _n
   for (n = 1; n < _n; ) n *= 2;
   seq.assign(2*n,ID); }
  void pull(int p) { seq[p] = cmb(seq[2*p], seq[2*p+1]); }
  void upd(int p, T val) { // set val at position p
   seq[p += n] = val; for (p /= 2; p; p /= 2) pull(p); }
  T query(int 1, int r) { // zero-indexed, inclusive
   T ra = ID, rb = ID;
    for (1 += n, r += n+1; 1 < r; 1 /= 2, r /= 2) {
     if (1&1) ra = cmb(ra, seq[1++]);
     if (r&1) rb = cmb(seq[--r],rb);
    return cmb(ra,rb);
};
```

LazySegmentTree.h

Description: 1D range increment and sum query. **Time:** $\mathcal{O}(\log N)$

78a06d, 26 lines

```
tcT, int SZ> struct LazySeq {
  static_assert(pct(SZ) == 1); // SZ must be power of 2
  const T ID{}; T cmb(T a, T b) { return a+b; }
  T seg[2*SZ], lazy[2*SZ];
  LazySeg() { FOR(i,2*SZ) seg[i] = lazy[i] = ID; }
  void push (int ind, int L, int R) {
    seg[ind] += (R-L+1)*lazy[ind]; // dependent on operation
   if (L != R) FOR(i,2) lazy[2*ind+i] += lazy[ind];
   lazy[ind] = 0;
  } // recalc values for current node
  void pull(int ind){seg[ind]=cmb(seg[2*ind],seg[2*ind+1]);}
  void build() { ROF(i,1,SZ) pull(i); }
  void upd(int lo,int hi,T inc,int ind=1,int L=0, int R=SZ-1) {
    push(ind,L,R); if (hi < L || R < lo) return;</pre>
    if (lo <= L && R <= hi) {
     lazy[ind] = inc; push(ind, L, R); return; }
    int M = (L+R)/2; upd(lo,hi,inc,2*ind,L,M);
    upd(lo,hi,inc,2*ind+1,M+1,R); pull(ind);
  T query (int lo, int hi, int ind=1, int L=0, int R=SZ-1) {
    push(ind,L,R); if (lo > R || L > hi) return ID;
    if (lo <= L && R <= hi) return seg[ind];</pre>
   int M = (L+R)/2; return cmb(query(lo,hi,2*ind,L,M),
      query(lo,hi,2*ind+1,M+1,R));
};
```

PSeg.h

Description: Persistent min segtree with lazy updates, no propagation. If making d a vector then save the results of upd and build in local variables first to avoid issues when vector resizes in C++14 or lower.

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

8f37fa, 45 lines

```
tcT, int SZ> struct pseq {
  static const int LIM = 2e7;
 struct node {
   int 1, r; T val = 0, lazy = 0;
   void inc(T x) { lazy += x; }
   T get() { return val+lazy; }
 };
 node d[LIM]; int nex = 0;
 int copy(int c) { d[nex] = d[c]; return nex++; }
 T cmb(T a, T b) { return min(a,b); }
 void pull(int c) { d[c].val =
    cmb(d[d[c].1].get(), d[d[c].r].get()); }
 T query(int c, int lo, int hi, int L, int R) {
    if (lo <= L && R <= hi) return d[c].get();</pre>
    if (R < lo || hi < L) return MOD;</pre>
    int M = (L+R)/2;
    return d[c].lazy+cmb(query(d[c].l,lo,hi,L,M),
              query(d[c].r,lo,hi,M+1,R));
  int upd(int c, int lo, int hi, T v, int L, int R) {
   if (R < lo || hi < L) return c;</pre>
    int x = copy(c);
    if (lo <= L && R <= hi) { d[x].inc(v); return x; }</pre>
    int M = (L+R)/2;
   d[x].1 = upd(d[x].1, lo, hi, v, L, M);
   d[x].r = upd(d[x].r,lo,hi,v,M+1,R);
   pull(x); return x;
 int build(const V<T>& arr, int L, int R) {
   int c = nex++;
   if (L == R) {
     if (L < sz(arr)) d[c].val = arr[L];
     return c;
    int M = (L+R)/2;
    d[c].l = build(arr, L, M), d[c].r = build(arr, M+1, R);
   pull(c); return c;
 void upd(int lo, int hi, T v) {
   loc.pb(upd(loc.bk, lo, hi, v, 0, SZ-1)); }
 T query(int ti, int lo, int hi) {
   return query(loc[ti],lo,hi,0,SZ-1); }
 void build(const V<T>&arr) {loc.pb(build(arr, 0, SZ-1));}
};
```

Treap.h

Description: Easy BBST. Use split and merge to implement insert and delete.

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

bdb758, 65 lines

```
using pt = struct tnode*;
struct thode {
  int pri, val; pt c[2]; // essential
  int sz; ll sum; // for range queries
  bool flip = 0; // lazy update
  tnode(int _val) {
    pri = rng(); sum = val = _val;
    sz = 1; c[0] = c[1] = nullptr;
  ~tnode() { FOR(i,2) delete c[i]; }
int getsz(pt x) { return x?x->sz:0; }
11 getsum(pt x) { return x?x->sum:0; }
pt prop(pt x) { // lazy propagation
  if (!x || !x->flip) return x;
  swap (x->c[0], x->c[1]);
  x \rightarrow flip = 0; FOR(i,2) if (x \rightarrow c[i]) x \rightarrow c[i] \rightarrow flip ^= 1;
  return x;
```

```
pt calc(pt x) {
  pt a = x - c[0], b = x - c[1];
  assert(!x->flip); prop(a), prop(b);
  x->sz = 1+getsz(a)+getsz(b);
  x->sum = x->val+getsum(a)+getsum(b);
  return x;
void tour(pt x, vi& v) { // print values of nodes,
  if (!x) return: // inorder traversal
  prop(x); tour(x->c[0],v); v.pb(x->val); tour(x->c[1],v);
pair<pt,pt> split(pt t, int v) { // >= v goes to the right
  if (!t) return {t,t};
  prop(t);
  if (t->val >= v) {
    auto p = split(t->c[0], v); t->c[0] = p.s;
    return {p.f,calc(t)};
    auto p = split(t->c[1], v); t->c[1] = p.f;
    return {calc(t),p.s};
pair<pt, pt> splitsz(pt t, int sz) { // sz nodes go to left
  if (!t) return {t,t};
  prop(t);
  if (\text{getsz}(t->c[0]) >= sz) {
    auto p = splitsz(t->c[0],sz); t->c[0] = p.s;
    return {p.f,calc(t)};
  } else {
    auto p=splitsz(t->c[1],sz-getsz(t->c[0])-1); t->c[1]=p.f;
    return {calc(t),p.s};
pt merge(pt l, pt r) { // keys in l < keys in r
  if (!1 || !r) return 1?:r;
  prop(l), prop(r); pt t;
  if (l->pri > r->pri) l->c[1] = merge(l->c[1],r), t = 1;
  else r - > c[0] = merge(1, r - > c[0]), t = r;
  return calc(t);
pt ins(pt x, int v) { // insert v
  auto a = split(x,v), b = split(a.s,v+1);
  return merge(a.f, merge(new tnode(v), b.s)); }
pt del(pt x, int v) { // delete v
  auto a = split(x, v), b = split(a.s, v+1);
  return merge(a.f,b.s); }
```

BIT2DOff.h

Description: point update and rectangle sum with offline 2D BIT. For each of the points to be updated, $x \in (0, SZ)$ and $y \neq 0$.

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

962052, 34 lines

```
template<class T, int SZ> struct OffBIT2D {
  bool mode = 0; // mode = 1 -> initialized
  vpi todo; // locations of updates to process
  int cnt[SZ], st[SZ];
  vi val; vector<T> bit; // store all BITs in single vector
  void init() { assert(!mode); mode = 1;
    int lst[SZ]; F0R(i,SZ) lst[i] = cnt[i] = 0;
    sort(all(todo),[](const pi& a, const pi& b) {
      return a.s < b.s; });
    each(t,todo) for (int x = t.f; x < SZ; x += x&-x)
      if (lst[x] != t.s) lst[x] = t.s, cnt[x] ++;
    int sum = 0; F0R(i,SZ) lst[i] = 0, st[i] = (sum += cnt[i]);
    val.rsz(sum); bit.rsz(sum); reverse(all(todo));
    each(t,todo) for (int x = t.f; x < SZ; x += x&-x)
      if (lst[x] != t.s) lst[x] = t.s, val[--st[x]] = t.s;</pre>
```

```
int rank(int y, int 1, int r) {
  return ub(begin(val)+l,begin(val)+r,y)-begin(val)-l; }
void UPD(int x, int y, T t) {
 for (y = rank(y, st[x], st[x]+cnt[x]); y \le cnt[x]; y += y&-y
   bit[st[x]+y-1] += t; }
void upd(int x, int y, T t) {
 if (!mode) todo.pb({x,y});
 else for (;x<SZ;x+=x\&-x) UPD (x,y,t); }
int QUERY(int x, int y) { T res = 0;
  for (y = rank(y, st[x], st[x]+cnt[x]); y; y -= y&-y) res +=
     \hookrightarrowbit[st[x]+y-1];
  return res; }
T query(int x, int y) { assert(mode);
  T res = 0; for (;x;x-=x\&-x) res += QUERY(x,y);
T query(int xl, int xr, int yl, int yr) {
  return query(xr,yr)-query(xl-1,yr)
    -query(xr,yl-1)+query(xl-1,yl-1); }
```

Numerical (4)

4.1 Matrices

Matrix.h

```
Description: 2D matrix operations.
```

```
"../../number-theory (11.1)/Modular Arithmetic/ModInt.h"
                                                      b18e29, 21 lines
using T = mi;
using Mat = V<V<T>>; // use array instead if tight TL
Mat makeMat(int r, int c) { return Mat(r, V<T>(c)); }
Mat makeId(int n) {
 Mat m = makeMat(n,n); FOR(i,n) m[i][i] = 1;
  return m;
Mat operator*(const Mat& a, const Mat& b) {
  int x = sz(a), y = sz(a[0]), z = sz(b[0]);
  assert (y == sz(b)); Mat c = makeMat(x,z);
  FOR(i,x) FOR(j,y) FOR(k,z) c[i][k] += a[i][j]*b[j][k];
  return c:
Mat& operator*=(Mat& a, const Mat& b) { return a = a*b; }
Mat pow(Mat m, 11 p) {
  int n = sz(m); assert(n == sz(m[0]) && p >= 0);
  Mat res = makeId(n);
 for (; p; p /= 2, m \star= m) if (p&1) res \star= m;
  return res;
```

Determinant.h

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

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

```
return res;
```

IntDeterminant.h

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

```
Time: \mathcal{O}(N^3)
                                                       3313dc, 18 lines
const 11 mod = 12345;
11 det(vector<vector<11>>& a) {
 int n = sz(a); ll ans = 1;
 rep(i,0,n) {
    rep(j,i+1,n) {
      while (a[j][i] != 0) { // gcd step
        ll t = a[i][i] / a[j][i];
        if (t) rep(k,i,n)
         a[i][k] = (a[i][k] - a[j][k] * t) % mod;
        swap(a[i], a[j]);
        ans \star = -1;
    ans = ans * a[i][i] % mod;
    if (!ans) return 0;
 return (ans + mod) % mod;
```

| SolveLinear.h

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

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

```
SolveLinear2.h
```

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

SolveLinearBinary.h

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

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

MatrixInv.h

Description: Uses gaussian elimination to convert into reduced row echelon form and calculates determinant. For determinant via arbitrary modulos, use a modified form of the Euclidean algorithm because modular inverse may not exist. If you have computed $A^{-1} \pmod{p^k}$, then the inverse $\pmod{p^{2k}}$ is $A^{-1}(2I-AA^{-1})$.

Time: $\mathcal{O}\left(N^3\right)$, determinant of 1000×1000 matrix of modints in 1 second if you reduce # of operations by half

```
const db EPS = 1e-9; // adjust?
int getRow(V<V<db>>& m, int R, int i, int nex) {
  pair<db,int> bes{0,-1}; // find row with max abs value
  FOR(j,nex,R) ckmax(bes,{abs(m[j][i]),j});
  return bes.f < EPS ? -1 : bes.s; }
int getRow(V<vmi>& m, int R, int i, int nex) {
```

```
FOR(j, nex,R) if (m[j][i] != 0) return j;
  return -1; }
pair<T, int > gauss (Mat& m) { // convert to reduced row echelon
  \hookrightarrow form
  if (!sz(m)) return {1,0};
  int R = sz(m), C = sz(m[0]), rank = 0, nex = 0;
  T prod = 1; // determinant
  F0R(i,C) {
    int row = getRow(m,R,i,nex);
    if (row == -1) { prod = 0; continue; }
   if (row != nex) prod \star= -1, swap(m[row], m[nex]);
    prod *= m[nex][i]; rank++;
    T x = 1/m[nex][i]; FOR(k,i,C) m[nex][k] *= x;
   FOR(j,R) if (j != nex) {
     T v = m[j][i]; if (v == 0) continue;
     FOR(k,i,C) m[j][k] \rightarrow v*m[nex][k];
   nex++;
  return {prod, rank};
Mat inv(Mat m) {
  int R = sz(m); assert(R == sz(m[0]));
  Mat x = makeMat(R, 2*R);
  F0R(i,R) {
   x[i][i+R] = 1;
   FOR(j,R) \times [i][j] = m[i][j];
  if (gauss(x).s != R) return Mat();
  Mat res = makeMat(R,R);
  FOR(i,R) FOR(j,R) res[i][j] = x[i][j+R];
  return res;
```

MatrixTree.h

Description: Kirchhoff's Matrix Tree Theorem. Given adjacency matrix. calculates # of spanning trees.

```
"MatrixInv.h"
                                                      48363d, 11 lines
T numSpan (const Mat& m) {
  int n = sz(m); Mat res = makeMat(n-1,n-1);
  FOR(i,n) FOR(j,i+1,n) {
   mi ed = m[i][j]; res[i][i] += ed;
   if (j != n−1) {
     res[j][j] += ed;
      res[i][j] -= ed, res[j][i] -= ed;
 return gauss (res).f;
```

ShermanMorrison.h

Description: Calculates $(A + uv^T)^{-1}$ given $B = A^{-1}$. Not invertible if sum=0.

```
void ad(Mat& B, const V<T>& u, const V<T>& v) {
 int n = sz(A); V < T > x(n), y(n);
 F0R(i,n) F0R(j,n)
   x[i] += B[i][j]*u[j], y[j] += v[i]*B[i][j];
  T sum = 1; FOR(i,n) FOR(j,n) sum += v[i]*B[i][j]*u[j];
 FOR(i,n) FOR(j,n) B[i][j] = x[i]*y[j]/sum;
```

Tridiagonal.h

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

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

This is useful for solving problems on the type

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

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

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

Fails if the solution is not unique.

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

Time: $\mathcal{O}(N)$ 8f9fa8, 26 lines typedef double T: vector<T> tridiagonal(vector<T> diag, const vector<T>& super, const vector<T>& sub, vector<T> b) { int n = sz(b); vi tr(n); rep(i, 0, n-1) { **if** (abs(diag[i]) < 1e-9 * abs(super[i])) { // diag[i] == 0b[i+1] = b[i] * diag[i+1] / super[i];if (i+2 < n) b[i+2] -= b[i] * sub[i+1] / super[i];</pre> diag[i+1] = sub[i]; tr[++i] = 1;diag[i+1] -= super[i]*sub[i]/diag[i]; b[i+1] -= b[i] * sub[i] / diag[i]; **for** (**int** i = n; i--;) { **if** (tr[i]) { swap(b[i], b[i-1]);diag[i-1] = diag[i];b[i] /= super[i-1];} else { b[i] /= diag[i]; if (i) b[i-1] -= b[i]*super[i-1]; return b;

4.2 Polynomials and recurrences

Polv.h

Description: Basic poly ops including division. Can replace T with double,

```
"../../number-theory (11.1)/Modular Arithmetic/ModInt.h"
                                                      cd218a, 73 lines
using T = mi; using poly = V<T>;
void remz(poly& p) { while (sz(p)&&p.bk==T(0)) p.pop_back(); }
poly REMZ(poly p) { remz(p); return p; }
polv rev(polv p) { reverse(all(p)); return p; }
poly shift (poly p, int x) {
  if (x \ge 0) p.insert(begin(p), x, 0);
  else assert(sz(p)+x \ge 0), p.erase(begin(p),begin(p)-x);
  return p;
poly RSZ(const poly& p, int x) {
  if (x <= sz(p)) return poly(begin(p), begin(p)+x);</pre>
  poly q = p; q.rsz(x); return q; }
T eval(const poly& p, T x) { // evaluate at point x
 T res = 0; R0F(i,sz(p)) res = x*res+p[i];
```

```
return res; }
poly dif(const poly& p) { // differentiate
 polv res; FOR(i,1,sz(p)) res.pb(T(i)*p[i]);
  return res; }
poly integ(const poly& p) { // integrate
  static poly invs{0,1};
  for (int i = sz(invs); i \le sz(p); ++i)
   invs.pb(-MOD/i*invs[MOD%i]);
  poly res(sz(p)+1); F0R(i,sz(p)) res[i+1] = p[i]*invs[i+1];
 return res;
poly& operator+= (poly& 1, const poly& r) {
 1.rsz(max(sz(1),sz(r))); F0R(i,sz(r)) 1[i] += r[i];
 return 1; }
poly& operator = (poly& 1, const poly& r) {
 1.rsz(max(sz(1),sz(r))); FOR(i,sz(r)) l[i] -= r[i];
poly& operator *= (poly& 1, const T& r) { each(t,1) t *= r;
 return 1; }
poly& operator/=(poly& 1, const T& r) { each(t,1) t /= r;
 return 1; }
poly operator+(poly 1, const poly& r) { return 1 += r; }
poly operator-(poly 1, const poly& r) { return 1 -= r; }
poly operator-(poly 1) { each(t,1) t *= -1; return 1; }
poly operator* (poly 1, const T& r) { return 1 *= r; }
poly operator* (const T& r, const poly& 1) { return 1*r; }
poly operator/(poly 1, const T& r) { return 1 /= r; }
poly operator* (const poly& 1, const poly& r) {
 if (!min(sz(l),sz(r))) return {};
 poly x(sz(1)+sz(r)-1);
 FOR(i, sz(1)) FOR(j, sz(r)) x[i+j] += l[i]*r[j];
 return x;
poly& operator*=(poly& 1, const poly& r) { return 1 = 1*r; }
pair<poly, poly> quoRemSlow(poly a, poly b) {
 remz(a); remz(b); assert(sz(b));
 T lst = b.bk, B = T(1)/lst; each(t,a) t *= B;
 each(t,b) t \star= B;
 poly q(max(sz(a)-sz(b)+1,0));
  for (int dif; (dif=sz(a)-sz(b)) >= 0; remz(a)) {
   q[dif] = a.bk; FOR(i,sz(b)) a[i+dif] -= q[dif]*b[i]; }
  each(t,a) t *= lst;
 return {q,a}; // quotient, remainder
poly operator% (const poly& a, const poly& b) {
 return quoRemSlow(a,b).s; }
T resultant (poly a, poly b) \{ // R(A,B) \}
 // =b_m^n*prod_{j=1}^mA(mu_j)
  // =b_m^na_n^m*prod_{i=1}^nprod_{j=1}^m (mu_j-lambda_i)
  // = (-1) ^{mn}a_n^m*prod_{i=1}^nB(lambda_i)
  // = (-1) ^{nm}R(B, A)
  // Also, R(A,B) = b_m^{(A-CB,B)} R(A-CB,B)
  int ad = sz(a)-1, bd = sz(b)-1;
  if (bd <= 0) return bd < 0 ? 0 : pow(b.bk.ad);</pre>
  int pw = ad; a = a\%b; pw -= (ad = sz(a) -1);
  return resultant(b,a)*pow(b.bk,pw)*T((bd&ad&1)?-1:1);
PolyRoots.h
Description: Finds the real roots of a polynomial.
```

Usage: poly_roots ($\{\{2, -3, 1\}\}, -1e9, 1e9\}$) // solve $x^2-3x+2=0$

Time: $\mathcal{O}\left(N^2\log(1/\epsilon)\right)$ "Poly.h" c9127a, 20 lines

```
typedef db T;
poly polyRoots(poly p, T xmin, T xmax) {
 if (sz(p) == 2) { return {-p[0]/p[1]}; }
 auto dr = polyRoots(dif(p), xmin, xmax);
```

dr.pb(xmin-1); dr.pb(xmax+1); sort(all(dr));

```
poly ret;
  FOR(i,sz(dr)-1) {
   T l = dr[i], h = dr[i+1];
   bool sign = eval(p,1) > 0;
   if (sign^(eval(p,h) > 0)) {
     FOR(it, 60) { // while (h-1 > 1e-8)
        auto m = (1+h)/2, f = eval(p, m);
        if ((f \le 0) ^ sign) 1 = m;
        else h = m;
      ret.pb((1+h)/2);
  return ret;
PolyInterpolate.h
Description: n points determine unique polynomial of degree \leq n-1. For
numerical precision pick v[k].f = c * \cos(k/(n-1)*\pi), k = 0...n-1.
Time: \mathcal{O}(n^2)
"Polv.h"
polv interpolate(V<pair<T,T>> v) {
  poly res, tmp{1};
  FOR(i, sz(v)) { T prod = 1; // add one point at a time
   FOR(j,i) v[i].s = prod*v[j].s, prod *= v[i].f-v[j].f;
   v[i].s \neq prod; res += v[i].s*tmp; tmp *= poly{-v[i].f,1};
  \} // add multiple of (x-v[0].f)*(x-v[1].f)*...*(x-v[i-1].f)
  return res;
LinearRecurrence.h
Description: Berlekamp-Massey. Computes linear recurrence C of order
N for sequence s of 2N terms. C[0] = 1 and for all i \geq sz(C) - 1,
\sum_{j=0}^{sz(C)-1} C[j]s[i-j] = 0.
Usage: LinRec L; L.init({0,1,1,2,3}); L.eval(5); L.eval(6); //
Time: init \Rightarrow \mathcal{O}(N|C|), eval \Rightarrow \mathcal{O}(|C|^2 \log p) or faster with FFT
"Poly.h"
struct LinRec {
  poly s, C, rC;
  void BM() {
    int x = 0; T b = 1;
    poly B; B = C = \{1\}; // B is fail vector
    FOR(i,sz(s)) { // update C after adding a term of s
     ++x; int L = sz(C), M = i+3-L;
     T d = 0; FOR(j,L) d += C[j]*s[i-j]; // [D^i]C*s
     if (d.v == 0) continue; // [D^i]C*s=0
     poly _{C} = C; T coef = d*inv(b);
     C.rsz(max(L,M)); FOR(j,sz(B)) C[j+x] -= coef*B[j];
     if (L < M) B = \_C, b = d, x = 0;
  void init(const poly& _s) {
   s = _s; BM();
    rC = C; reverse(all(rC));
    C.erase(begin(C)); each(t,C) t \star = -1;
  poly getPow(ll p) { // get x^p mod rC
    if (p == 0) return {1};
   poly r = getPow(p/2); r = (r*r) %rC;
   return p&1?(r*poly{0,1})%rC:r;
  T dot(poly v) { // dot product with s
   T ans = 0; FOR(i,sz(v)) ans += v[i]*s[i];
    return ans; } // get p-th term of rec
  T eval(ll p) { assert(p >= 0); return dot(getPow(p)); }
```

```
PolyInvSimpler.h
Description: computes A^{-1} such that AA^{-1} \equiv 1 \pmod{x^n}. New-
ton's method: If you want F(x) = 0 and F(Q_k) \equiv 0 \pmod{x^a} then
Q_{k+1}=Q_k-rac{F(Q_k)}{F'(Q_k)}\ (\mathrm{mod}\ x^{2a}) satisfies F(Q_{k+1})\equiv 0\ (\mathrm{mod}\ x^{2a}). Ap-
plication: if f(n), g(n) are the #s of forests and trees on n nodes then
\sum_{n=0}^{\infty} f(n)x^n = \exp\left(\sum_{n=1}^{\infty} \frac{g(n)}{n!}\right).
Usage: vmi v\{1,5,2,3,4\}; ps(exp(2*log(v,9),9)); // squares v
Time: \mathcal{O}(N \log N). For N = 5 \cdot 10^5, inv~270ms, \log \sim 350ms, \exp \sim 550ms
poly inv(poly A, int n) { // Q-(1/Q-A)/(-Q^{-2})
  poly B{inv(A[0])};
  for (int x = 2; x/2 < n; x *= 2)
   B = 2*B-RSZ (conv(RSZ(A,x),conv(B,B)),x);
  return RSZ(B,n);
poly sqrt (const poly & A, int n) { //Q-(Q^2-A)/(2Q)
  assert (A[0].v == 1); poly B{1};
  for (int x = 2; x/2 < n; x *= 2)
    B = inv(T(2)) *RSZ(B+conv(RSZ(A,x),inv(B,x)),x);
  return RSZ(B,n);
// return {quotient, remainder}
pair<poly, poly> quoRem(const poly& f, const poly& g) {
  if (sz(f) < sz(q)) return {{}, f};</pre>
  poly q = conv(inv(rev(q), sz(f) - sz(q) + 1), rev(f));
  q = rev(RSZ(q, sz(f) - sz(q) + 1));
  poly r = RSZ(f-conv(q,q),sz(q)-1); return \{q,r\};
poly log(poly A, int n) { assert(A[0].v == 1); // (ln A)' = A'/
  A.rsz(n); return integ(RSZ(conv(dif(A),inv(A,n-1)),n-1)); }
poly exp(poly A, int n) { assert(A[0].v == 0);
  polv B{1}, IB{1}; // inverse of B
  for (int x = 1; x < n; x *= 2) {
    IB = 2*IB-RSZ(conv(B,conv(IB,IB)),x);
    poly Q = dif(RSZ(A,x)); Q += RSZ(conv(IB, dif(B)-conv(B,Q))
       \hookrightarrow, 2*x-1);
    B = B+RSZ (conv(B,RSZ(A,2*x)-integ(Q)),2*x);
  return RSZ(B,n);
PolyMultipoint.h
Description: Multipoint evaluation and interpolation
Time: \mathcal{O}\left(N\log^2 N\right)
"PolyInv.h", "PolyConv.h"
                                                          9f6b18, 29 lines
void segProd(V<poly>& stor, poly& v, int ind, int 1, int r) {
  \hookrightarrow // v -> places to evaluate at
  if (1 == r) { stor[ind] = {-v[1],1}; return; }
  int m = (1+r)/2; seqProd(stor, v, 2*ind, 1, m); seqProd(stor, v, 2*
     \hookrightarrow ind+1,m+1,r);
  stor[ind] = conv(stor[2*ind],stor[2*ind+1]);
void evalAll(V<poly>& stor, poly& res, poly v, int ind = 1) {
 v = quoRem(v,stor[ind]).s;
  if (sz(stor[ind]) == 2) { res.pb(sz(v)?v[0]:0); return; }
  evalAll(stor,res,v,2*ind); evalAll(stor,res,v,2*ind+1);
// evaluate polynomial v at points in p
poly multiEval(poly v, poly p) {
  V < poly > stor(4 * sz(p)); segProd(stor, p, 1, 0, sz(p) - 1);
  poly res; evalAll(stor,res,v); return res; }
poly combAll(V<poly>& stor, poly& dems, int ind, int 1, int r)
  if (l == r) return {dems[l]};
```

4.3 Optimization

GoldenSectionSearch.h

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

Usage: gss(-1000,1000,[](db x) { return 4+x+.3*x*x; }); // -5/3 **Time:** $\mathcal{O}(\log((b-a)/\epsilon))$

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

HillClimbing.h

Description: Poor man's optimization for unimodal functions

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

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

Integrate.h

return tot*dif/3;

Description: Integration of a function over an interval using Simpson's rule, exact for polynomials of degree up to 3. The error should be proportional to dif^4 , although in practice you will want to verify that the result is stable to desired precision when epsilon changes.

```
Usage: quad([] (db x) { return x*x+3*x+1; }, 2, 3) // 14 833, 6 nies

template < class F > db quad(F f, db a, db b) {
  const int n = 1000;
  db dif = (b-a)/2/n, tot = f(a)+f(b);
  FOR(i,1,2*n) tot + = f(a+i*dif)*(i&1?4:2);
```

IntegrateAdaptive.h

```
Description: Unused. Fast integration using adaptive Simpson's rule, exact
for polynomials of degree up to 5.
Usage: db z, y;
```

```
db h(db x) { return x*x + y*y + z*z \le 1; }
db g(db y) \{ :: y = y; \text{ return quad}(h, -1, 1); \}
db f(db z) \{ :: z = z; return quad(q, -1, 1); \}
db sphereVol = quad(f, -1, 1), pi = sphereVol*3/4;
                                                       3b316e, 10 lines
template < class F > db simpson (F f, db a, db b) {
  db c = (a+b)/2; return (f(a)+4*f(c)+f(b))*(b-a)/6; }
template < class F > db rec(F& f, db a, db b, db eps, db S) {
  db c = (a+b)/2;
  db S1 = simpson(f,a,c), S2 = simpson(f,c,b), T = S1+S2;
  if (abs(T-S) <=15*eps || b-a<1e-10) return T+(T-S)/15;</pre>
  return rec(f,a,c,eps/2,S1)+rec(f,c,b,eps/2,S2);
template < class F > db quad (F f, db a, db b, db eps = 1e-8) {
  return rec(f,a,b,eps,simpson(f,a,b)); }
```

Simplex.h

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

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

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

```
using T = db; // double probably suffices
using vd = V<T>; using vvd = V<vd>;
const T eps = 1e-8, inf = 1/.0;
struct LPSolver {
```

#define ltj(X) if $(s==-1 \mid | mp(X[j],N[j]) < mp(X[s],N[s]))$ s=jint m, n; // # m = contraints, # n = variables vi N, B; // N[j] = non-basic variable (j-th column), = 0 vvd D; // B[i] = basic variable (i-th row) 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)) { FOR(i,m) FOR(j,n) D[i][j] = A[i][j];FOR(i,m) B[i] = n+i, D[i][n] = -1, D[i][n+1] = b[i]; // B[i]: basic variable for each constraint // D[i][n]: artificial variable for testing feasibility FOR(j,n) N[j] = j, D[m][j] = -c[j];// D[m] stores negation of objective, // which we want to minimize N[n] = -1; D[m+1][n] = 1; // to find initial feasible } // solution, minimize artificial variable void pivot(int r, int s) { // swap B[r] (row) T inv = 1/D[r][s]; // with N[r] (column) FOR(i,m+2) if (i != r && abs(D[i][s]) > eps) { T binv = D[i][s]*inv;FOR(j, n+2) if (j != s) D[i][j] -= D[r][j] *binv;D[i][s] = -binv;D[r][s] = 1; F0R(j,n+2) D[r][j] *= inv; // scale r-th row swap(B[r],N[s]);bool simplex(int phase) { **int** x = m+phase-1;

while (1) { // if phase=1, ignore artificial variable

if (D[x][s] >= -eps) return 1;

int s = -1; FOR(j, n+1) if (N[j] != -phase) ltj(D[x]);

// find most negative col for nonbasic (NB) variable

```
// can't get better sol by increasing NB variable
    int r = -1:
    F0R(i,m) {
      if (D[i][s] <= eps) continue;</pre>
      if (r == -1 \mid | mp(D[i][n+1] / D[i][s], B[i])
             < mp(D[r][n+1] / D[r][s], B[r])) r = i;
      // find smallest positive ratio
    } // -> max increase in NB variable
    if (r == -1) return 0; // objective is unbounded
    pivot(r,s);
T solve(vd& x) { // 1. check if x=0 feasible
  int r = 0; FOR(i,1,m) if (D[i][n+1] < D[r][n+1]) r = i;
  if (D[r][n+1] < -eps) { // if not, find feasible start
    pivot(r,n); // make artificial variable basic
    assert(simplex(2)); // I think this will always be true??
    if (D[m+1][n+1] < -eps) return -inf;</pre>
    // D[m+1][n+1] is max possible value of the negation of
    // artificial variable, optimal value should be zero
    // if exists feasible solution
    FOR(i,m) if (B[i] == -1) { // artificial var basic
      int s = 0; FOR(j,1,n+1) ltj(D[i]); // -> nonbasic
      pivot(i,s);
  bool ok = simplex(1); x = vd(n);
  FOR(i,m) if (B[i] < n) x[B[i]] = D[i][n+1];
  return ok ? D[m][n+1] : inf;
```

4.4 Fourier transforms

FastFourierTransform.h

vector<C> in(n), out(n);

copy(all(a), begin(in));

c99f9c, 67 lines

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

```
Time: \mathcal{O}(N \log N) with N = |A| + |B| (~1s for N = 2^{22})
                                                     00ced6, 35 lines
typedef complex<double> C;
typedef vector<double> vd;
void fft(vector<C>& a) {
 int n = sz(a), L = 31 - __builtin_clz(n);
  static vector<complex<long double>> R(2, 1);
  static vector<C> rt(2, 1); // (^ 10% faster if double)
  for (static int k = 2; k < n; k *= 2) {
   R.resize(n); rt.resize(n);
    auto x = polar(1.0L, acos(-1.0L) / k);
    rep(i,k,2*k) rt[i] = R[i] = i&1 ? R[i/2] * x : R[i/2];
  rep(i,0,n) rev[i] = (rev[i / 2] | (i & 1) << L) / 2;
  rep(i,0,n) if (i < rev[i]) swap(a[i], a[rev[i]]);
  for (int k = 1; k < n; k *= 2)
    for (int i = 0; i < n; i += 2 * k) rep(j, 0, k) {
     Cz = rt[j+k] * a[i+j+k]; // (25% faster if hand-rolled)
      a[i + j + k] = a[i + j] - z;
      a[i + j] += z;
vd conv(const vd& a, const vd& b) {
 if (a.empty() || b.empty()) return {};
 vd res(sz(a) + sz(b) - 1);
  int L = 32 - __builtin_clz(sz(res)), n = 1 << L;</pre>
```

```
rep(i,0,sz(b)) in[i].imag(b[i]);
fft(in):
for (C& x : in) x *= x;
rep(i, 0, n) out[i] = in[-i & (n - 1)] - conj(in[i]);
fft(out);
rep(i, 0, sz(res)) res[i] = imag(out[i]) / (4 * n);
return res;
```

FastFourierTransformMod.h

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

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

```
typedef vector<11> v1;
template<int M> vl convMod(const vl &a, const vl &b) {
 if (a.emptv() || b.emptv()) return {};
 vl res(sz(a) + sz(b) - 1);
 int B=32-__builtin_clz(sz(res)), n=1<<B, cut=int(sqrt(M));</pre>
  vector<C> L(n), R(n), outs(n), outl(n);
  rep(i,0,sz(a)) L[i] = C((int)a[i] / cut, (int)a[i] % cut);
  rep(i,0,sz(b)) R[i] = C((int)b[i] / cut, (int)b[i] % cut);
  fft(L), fft(R);
  rep(i, 0, n) {
    int j = -i \& (n - 1);
    outl[j] = (L[i] + conj(L[j])) * R[i] / (2.0 * n);
    outs[j] = (L[i] - conj(L[j])) * R[i] / (2.0 * n) / 1i;
  fft(outl), fft(outs);
  rep(i, 0, sz(res)) {
    ll av = ll(real(outl[i])+.5), cv = ll(imag(outs[i])+.5);
    11 \text{ bv} = 11(\text{imag}(\text{outl}[i]) + .5) + 11(\text{real}(\text{outs}[i]) + .5);
    res[i] = ((av % M * cut + bv) % M * cut + cv) % M;
  return res;
```

FFT.h

Description: Multiply polynomials of ints for any modulus $< 2^{31}$. For XOR convolution ignore m within fft.

Time: $\mathcal{O}(N \log N)$. For $N = 10^6$, conv ~0.13ms, conv_general ~320ms.

```
"ModInt.h"
// const int MOD = 998244353:
tcT> void fft(V<T>& A, bool invert = 0) { // NTT
  int n = sz(A); assert((T::mod-1)%n == 0); V<T> B(n);
  for (int b = n/2; b; b /= 2, swap (A, B)) { // w = n/b'th root
    T w = pow(T::rt(), (T::mod-1)/n*b), m = 1;
    for (int i = 0; i < n; i += b*2, m *= w) FOR (j,b) {
     T u = A[i+j], v = A[i+j+b] *m;
      B[i/2+j] = u+v; B[i/2+j+n/2] = u-v;
  if (invert) { reverse(1+all(A));
    T z = inv(T(n)); each(t,A) t *= z; }
} // for NTT-able moduli
tcT> V<T> conv(V<T> A, V<T> B) {
  if (!min(sz(A),sz(B))) return {};
  int s = sz(A) + sz(B) - 1, n = 1; for (; n < s; n *= 2);
  A.rsz(n), fft(A); B.rsz(n), fft(B);
  FOR(i,n) A[i] \star= B[i];
  fft(A,1); A.rsz(s); return A;
template < class M, class T> V < M> mulMod(const V < T> & x, const V < T
  auto con = [](const V<T>& v) {
```

V<M> w(sz(v)); FOR(i,sz(v)) w[i] = (int)v[i];

return w; };

19a793, 24 lines

```
return conv(con(x), con(y));
} // arbitrary moduli
tcT> V<T> conv_general(const V<T>& A, const V<T>& B) {
    using m0 = mint<(119<<23)+1,62>; auto c0 = mulMod<m0>(A,B);
    using m1 = mint<(5<<25)+1, 62>; auto c1 = mulMod<m1>(A,B);
    using m2 = mint<(7<<26)+1, 62>; auto c2 = mulMod<m2>(A,B);
    int n = sz(c0); V<T> res(n); m1 r01 = inv(m1(m0::mod));
    m2 r02 = inv(m2(m0::mod)), r12 = inv(m2(m1::mod));
    FOR(i,n) { // a=remainder mod m0::mod, b fixes it mod m1::mod
    int a = c0[i].v, b = ((c1[i]-a)*r01).v,
        c = (((c2[i]-a)*r02-b)*r12).v;
    res[i] = (T(c)*m1::mod+b)*m0::mod+a; // c fixes m2::mod
}
return res;
}
```

FastSubsetTransform.h

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

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

464cf3, 16 lines

Number theory (5)

5.1 Modular arithmetic

Modular Arithmetic.h

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

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

ModInverse.h

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

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

ModPow.h

b83e45, 8 lines

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

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

ModLog.h

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

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

ModSum.h

Description: Sums of mod'ed arithmetic progressions.

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

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

5c5bc5, 16 lines

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

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

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

ModMulLL.h

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

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

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

ModSqrt.h

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

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

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

5.2 Primality

FastEratosthenes.h

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

Time: LIM=1e9 ≈ 1.5 s

6b2912, 20 lines

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

MillerRabin.h

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

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

"ModMulLL.h" 60dcd1, 12 lines

bool isPrime(ull n) {

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

Factor.h

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

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

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

5.3 Divisibility

euclid.h

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

11 euclid(11 a, 11 b, 11 &x, 11 &y) {
 if (!b) return x = 1, y = 0, a;
 11 d = euclid(b, a % b, y, x);
 return y -= a/b * x, d;
}

CRT.h

 $\textbf{Description:} \ \ \text{Chinese Remainder Theorem.}$

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

5.3.1 Bézout's identity

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

$$ax + by = d$$

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

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

phiFunction.h

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

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

cf7d6d, 8 lis

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

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

5.4 Fractions

ContinuedFractions.h

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

For consecutive convergents, $p_{k+1}q_k - q_{k+1}p_k = (-1)^k$. $(p_k/q_k$ alternates between > x and < x.) If x is rational, y eventually becomes ∞ ; if x is the root of a degree 2 polynomial the a's eventually become cyclic.

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

```
typedef double d; // for N \sim 1e7; long double for N \sim 1e9
pair<ll, ll> approximate(d x, ll N) {
  11 LP = 0, LO = 1, P = 1, O = 0, inf = LLONG MAX; d v = x;
  for (::) {
    11 \lim_{N \to \infty} = \min(P ? (N-LP) / P : \inf, O ? (N-LO) / O : \inf),
       a = (11) floor(v), b = min(a, lim),
       NP = b*P + LP, NQ = b*Q + LQ;
    if (a > b) {
      // If b > a/2, we have a semi-convergent that gives us a
      // better approximation; if b = a/2, we *may* have one.
      // Return {P, O} here for a more canonical approximation.
      return (abs(x - (d)NP / (d)NO) < abs(x - (d)P / (d)O)) ?
        make_pair(NP, NQ) : make_pair(P, Q);
    if (abs(y = 1/(y - (d)a)) > 3*N) {
      return {NP, NQ};
    LP = P; P = NP;
    LO = O; O = NO;
```

FracBinarySearch.h

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

```
Usage: fracBS([](Frac f) { return f.p>=3*f.q; }, 10); // {1,3}
Time: \mathcal{O}(\log(N))
struct Frac { 11 p, q; };
template<class F>
Frac fracBS(F f, 11 N) {
 bool dir = 1, A = 1, B = 1;
 Frac 10\{0, 1\}, hi\{1, 1\}; // Set hi to 1/0 to search (0, N)
 if (f(lo)) return lo;
  assert(f(hi));
  while (A | | B) {
    11 adv = 0, step = 1; // move hi if dir, else lo
    for (int si = 0; step; (step *= 2) >>= si) {
      Frac mid{lo.p * adv + hi.p, lo.g * adv + hi.g};
      if (abs(mid.p) > N || mid.q > N || dir == !f(mid)) {
        adv -= step; si = 2;
    hi.p += lo.p * adv;
    hi.q += lo.q * adv;
    dir = !dir;
    swap(lo, hi);
    A = B; B = !!adv;
```

5.5 Pythagorean Triples

return dir ? hi : lo;

The Pythagorean triples are uniquely generated by

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

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

5.6 Primes

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

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

5.7 Estimates

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

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

5.8 Mobius Function

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

Mobius Inversion:

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

Other useful formulas/forms:

IntPerm multinomial

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

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

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

Combinatorial (6)

6.1 Permutations

6.1.1 Factorial

n						9		
n!	1 2 6	24 1	20 720	5040	40320	362880	3628800	
n							17	
n!	4.0e7	4.8e	8 6.2e	9 8.7e	10 1.3e	12 2.1e	13 3.6e14	
n	20	25	30	40	50 1	00 - 15	0 171	
n!	2e18	2e25	3e32	8e47 3	Be64 9e	157 6e2	$62 > DBL_M$	AX

IntPerm.h

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

Time: $\mathcal{O}(n)$ 044568, 6 lines int permToInt(vi& v) {

```
int use = 0, i = 0, r = 0;
for(int x:v) r = r * ++i + \_builtin_popcount(use & -(1<<x)),
                                     // (note: minus, not ~!)
return r;
```

6.1.2 Cycles

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

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

6.1.3 Derangements

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

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

6.1.4 Burnside's lemma

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

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

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

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

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

Partitions and subsets

6.2.1 Partition function

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

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

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

6.2.2 Lucas' Theorem

Let n, m be non-negative integers and p a prime. Write $n = n_k p^k + ... + n_1 p + n_0$ and $m = m_k p^k + ... + m_1 p + m_0$. Then $\binom{n}{m} \equiv \prod_{i=0}^{k} \binom{n_i}{m_i} \pmod{p}.$ **6.2.3** Binomials

multinomial.h

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

11 c = 1, m = v.emptv() ? 1 : v[0];rep(i,1,sz(v)) rep(j,0,v[i]) c = c * ++m / (j+1);return c;

General purpose numbers

6.3.1 Bernoulli numbers

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

Sums of powers:

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

Euler-Maclaurin formula for infinite sums:

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

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

6.3.2 Stirling numbers of the first kind

Number of permutations on n items with k cycles.

$$c(n,k) = c(n-1,k-1) + (n-1)c(n-1,k), \ c(0,0) = 1$$

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

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

6.3.3 Eulerian numbers

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

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

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

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

6.3.4 Stirling numbers of the second kind

Partitions of n distinct elements into exactly k groups.

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

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

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

6.3.5 Bell numbers

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

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

6.3.6 Labeled unrooted trees

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

6.3.7 Catalan numbers

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

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

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

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

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

Graph (7)

7.1 Fundamentals

BellmanFord.h

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

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

FlovdWarshall.h

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

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

531245, 12 lines

```
const 11 inf = 1LL << 62;</pre>
void floydWarshall (vector<vector<ll>>& m) {
  int n = sz(m);
  rep(i, 0, n) m[i][i] = min(m[i][i], 0LL);
  rep(k,0,n) rep(i,0,n) rep(j,0,n)
   if (m[i][k] != inf && m[k][j] != inf) {
      auto newDist = max(m[i][k] + m[k][j], -inf);
     m[i][j] = min(m[i][j], newDist);
  rep(k, 0, n) if (m[k][k] < 0) rep(i, 0, n) rep(j, 0, n)
    if (m[i][k] != inf && m[k][j] != inf) m[i][j] = -inf;
```

TopoSort.h

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

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

66a137, 14 lines

```
vi topoSort(const vector<vi>& gr) {
```

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

7.2 Network flow

PushRelabel.h

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

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

0ae1d4, 48 lines

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

```
MinCostMaxFlow.h
```

Description: Min-cost max-flow. cap[i][j] != cap[j][i] is allowed; double edges are not. If costs can be negative, call setpi before maxflow, but note that negative cost cycles are not supported. To obtain the actual flow, look at positive values only.

```
Time: Approximately \mathcal{O}\left(E^2\right)
#include <bits/extc++.h>
const 11 INF = numeric_limits<11>::max() / 4;
typedef vector<ll> VL;
struct MCMF {
 int N;
  vector<vi> ed, red;
  vector<VL> cap, flow, cost;
 vi seen;
 VL dist, pi;
  vector<pii> par;
  MCMF (int N) :
    N(N), ed(N), red(N), cap(N, VL(N)), flow(cap), cost(cap),
    seen(N), dist(N), pi(N), par(N) {}
  void addEdge(int from, int to, ll cap, ll cost) {
    this->cap[from][to] = cap;
    this->cost[from][to] = cost;
    ed[from].push_back(to);
    red[to].push_back(from);
  void path(int s) {
    fill(all(seen), 0);
    fill(all(dist), INF);
    dist[s] = 0; ll di;
    __gnu_pbds::priority_queue<pair<ll, int>> q;
    vector<decltype(q)::point_iterator> its(N);
    q.push({0, s});
    auto relax = [&](int i, ll cap, ll cost, int dir) {
      11 val = di - pi[i] + cost;
      if (cap && val < dist[i]) {
        dist[i] = val;
        par[i] = \{s, dir\};
        if (its[i] == q.end()) its[i] = q.push({-dist[i], i});
        else q.modify(its[i], {-dist[i], i});
    };
    while (!q.empty()) {
      s = q.top().second; q.pop();
      seen[s] = 1; di = dist[s] + pi[s];
      for (int i : ed[s]) if (!seen[i])
        relax(i, cap[s][i] - flow[s][i], cost[s][i], 1);
      for (int i : red[s]) if (!seen[i])
        relax(i, flow[i][s], -cost[i][s], 0);
    rep(i, 0, N) pi[i] = min(pi[i] + dist[i], INF);
  pair<11, 11> maxflow(int s, int t) {
    11 totflow = 0, totcost = 0;
    while (path(s), seen[t]) {
     11 fl = INF;
      for (int p,r,x = t; tie(p,r) = par[x], x != s; x = p)
        fl = min(fl, r ? cap[p][x] - flow[p][x] : flow[x][p]);
```

```
totflow += fl;
      for (int p,r,x = t; tie(p,r) = par[x], x != s; x = p)
       if (r) flow[p][x] += fl;
        else flow[x][p] -= fl;
    rep(i, 0, N) rep(j, 0, N) totcost += cost[i][j] * flow[i][j];
    return {totflow, totcost};
  // If some costs can be negative, call this before maxflow:
  void setpi(int s) { // (otherwise, leave this out)
    fill(all(pi), INF); pi[s] = 0;
    int it = N, ch = 1; l1 v;
    while (ch-- && it--)
     rep(i,0,N) if (pi[i] != INF)
        for (int to : ed[i]) if (cap[i][to])
          if ((v = pi[i] + cost[i][to]) < pi[to])</pre>
           pi[to] = v, ch = 1;
    assert(it >= 0); // negative cost cycle
};
```

EdmondsKarp.h

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

```
template<class T> T edmondsKarp(vector<unordered_map<int, T>>&
  assert (source != sink);
 T flow = 0;
 vi par(sz(graph)), q = par;
  for (;;) {
   fill(all(par), -1);
   par[source] = 0;
   int ptr = 1;
   q[0] = source;
   rep(i,0,ptr) {
     int x = q[i];
     for (auto e : graph[x]) {
       if (par[e.first] == -1 && e.second > 0) {
         par[e.first] = x;
         q[ptr++] = e.first;
         if (e.first == sink) goto out;
   return flow;
   T inc = numeric_limits<T>::max();
   for (int y = sink; y != source; y = par[y])
     inc = min(inc, graph[par[y]][y]);
    flow += inc;
    for (int y = sink; y != source; y = par[y]) {
     int p = par[v];
     if ((graph[p][y] -= inc) <= 0) graph[p].erase(y);</pre>
     graph[y][p] += inc;
```

MinCut.h

Description: After running max-flow, the left side of a min-cut from s to t is 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}\left(V^3\right)
                                                        8b0e19, 21 lines
pair<int, vi> globalMinCut(vector<vi> mat) {
 pair<int, vi> best = {INT MAX, {}};
 int n = sz(mat);
 vector<vi> co(n);
 rep(i, 0, n) co[i] = {i};
 rep(ph,1,n) {
    vi w = mat[0];
    size_t s = 0, t = 0;
    rep(it,0,n-ph) { // O(V^2) \rightarrow O(E \log V) with prio. queue
      w[t] = INT MIN;
      s = t, t = max_element(all(w)) - w.begin();
      rep(i,0,n) w[i] += mat[t][i];
    best = min(best, \{w[t] - mat[t][t], co[t]\});
    co[s].insert(co[s].end(), all(co[t]));
    rep(i, 0, n) mat[s][i] += mat[t][i];
    rep(i, 0, n) mat[i][s] = mat[s][i];
    mat[0][t] = INT_MIN;
 return best;
```

GomoryHu.h

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

7.3 Matching

hopcroftKarp.h

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

int hopcroftKarp(vector<vi>& g, vi& btoa) {

```
int res = 0;
vi A(g.size()), B(btoa.size()), cur, next;
for (;;) {
  fill(all(A), 0);
  fill(all(B), \theta);
  cur.clear();
  for (int a : btoa) if (a != -1) A[a] = -1;
  rep(a, 0, sz(g)) if(A[a] == 0) cur.push_back(a);
  for (int lay = 1;; lay++) {
   bool islast = 0;
    next.clear();
    for (int a : cur) for (int b : g[a]) {
      if (btoa[b] == -1) {
       B[b] = lav;
       islast = 1;
      else if (btoa[b] != a && !B[b]) {
       B[b] = lay;
        next.push_back(btoa[b]);
    if (islast) break;
    if (next.empty()) return res;
    for (int a : next) A[a] = lay;
    cur.swap(next);
  rep(a, 0, sz(q))
    res += dfs(a, 0, g, btoa, A, B);
```

DFSMatching.h

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

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

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

MinimumVertexCover.h

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

```
"DFSMatching.h" da4196, 20 lines

vi cover(vector<vi>& g, int n, int m) {

vi match(m, -1);
```

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

WeightedMatching.h

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

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

GeneralMatching.h

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

```
"../numerical/MatrixInverse-mod.h"
                                                       cb1912, 40 lines
vector<pii> generalMatching(int N, vector<pii>& ed) {
 vector<vector<ll>> mat(N, vector<ll>(N)), A;
  for (pii pa : ed) {
    int a = pa.first, b = pa.second, r = rand() % mod;
    mat[a][b] = r, mat[b][a] = (mod - r) % mod;
```

```
int r = matInv(A = mat), M = 2*N - r, fi, fj;
assert (r % 2 == 0);
if (M != N) do {
  mat.resize(M, vector<ll>(M));
  rep(i,0,N) {
    mat[i].resize(M);
    rep(j,N,M) {
      int r = rand() % mod;
      mat[i][j] = r, mat[j][i] = (mod - r) % mod;
} while (matInv(A = mat) != M);
vi has (M, 1); vector<pii> ret;
rep(it,0,M/2) {
  rep(i,0,M) if (has[i])
    rep(j,i+1,M) if (A[i][j] && mat[i][j]) {
      fi = i; fj = j; goto done;
  } assert(0); done:
  if (fj < N) ret.emplace_back(fi, fj);</pre>
  has[fi] = has[fi] = 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

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

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

```
BiconnectedComponents.h
```

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

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

2965e5, 33 lines

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

2sat.h

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

```
Usage: TwoSat ts(number of boolean variables);
ts.either(0, \sim3); // Var 0 is true or var 3 is false
ts.setValue(2); // Var 2 is true
ts.atMostOne(\{0, \sim 1, 2\}); // \ll 1 of vars 0, \sim 1 and 2 are true
ts.solve(); // Returns true iff it is solvable
ts.values[0..N-1] holds the assigned values to the vars
```

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

```
5f9706, 56 lines
```

```
struct TwoSat {
 int N:
 vector<vi> gr;
 vi values; // 0 = false, 1 = true
 TwoSat (int n = 0) : N(n), gr(2*n) {}
```

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

EulerWalk.h

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

```
vi eulerWalk(vector<vector<pii>% gr, int nedges, int src=0) {
  int n = sz(gr);
  vi D(n), its(n), eu(nedges), ret, s = {src};
  D[src]++; // to allow Euler paths, not just cycles
  while (!s.empty()) {
    int x = s.back(), y, e, &it = its[x], end = sz(gr[x]);
    if (it == end) { ret.push_back(x); s.pop_back(); continue; }
    tie(y, e) = gr[x][it++];
  if (!eu[e]) {
    D[x]--, D[y]++;
```

```
eu[e] = 1; s.push_back(y);
    }
for (int x : D) if (x < 0 || sz(ret) != nedges+1) return {};
return {ret.rbegin(), ret.rend()};
}</pre>
```

7.5 Coloring

EdgeColoring.h

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

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

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

7.6 Heuristics

MaximalCliques.h

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

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

b0d5b1, 12 lines

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

MaximumClique.h

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

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

```
f7c0bc, 49
```

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

MaximumIndependentSet.h

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

7.7 Trees

BinaryLifting.h

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

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

bfce85, 25 lines

LCA CompressTree HLD LinkCutTree

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

LCA.h

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

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

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

CompressTree.h

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

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

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

```
rep(i,0,m) {
  int a = li[i], b = li[i+1];
  li.push back(lca.lca(a, b));
sort(all(li), cmp);
li.erase(unique(all(li)), li.end());
rep(i, 0, sz(li)) rev[li[i]] = i;
vpi ret = {pii(0, li[0])};
rep(i, 0, sz(li)-1) {
  int a = li[i], b = li[i+1];
  ret.emplace_back(rev[lca.lca(a, b)], b);
return ret;
```

template <bool VALS EDGES> struct HLD {

HLD.h

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

values are stored in the edges, as opposed to the nodes. All values initialized to the segtree default. Root must be 0. Time: $\mathcal{O}\left((\log N)^2\right)$

```
"../data-structures/LazySegmentTree.h"
                                                                        6f34db, 46 lines
```

```
int N, tim = 0;
vector<vi> adj;
vi par, siz, depth, rt, pos;
HLD(vector<vi> adi )
  : N(sz(adj_)), adj(adj_), par(N, -1), siz(N, 1), depth(N),
    rt(N),pos(N),tree(new Node(0, N)) { dfsSz(0); dfsHld(0); }
void dfsSz(int v) {
  if (par[v] != -1) adj[v].erase(find(all(adj[v]), par[v]));
  for (int& u : adj[v]) {
    par[u] = v, depth[u] = depth[v] + 1;
    dfsSz(u);
   siz[v] += siz[u];
   if (siz[u] > siz[adj[v][0]]) swap(u, adj[v][0]);
void dfsHld(int v) {
  pos[v] = tim++;
  for (int u : adj[v]) {
   rt[u] = (u == adj[v][0] ? rt[v] : u);
    dfsHld(u);
template <class B> void process(int u, int v, B op) {
  for (; rt[u] != rt[v]; v = par[rt[v]]) {
    if (depth[rt[u]] > depth[rt[v]]) swap(u, v);
    op(pos[rt[v]], pos[v] + 1);
  if (depth[u] > depth[v]) swap(u, v);
  op(pos[u] + VALS_EDGES, pos[v] + 1);
void modifyPath(int u, int v, int val) {
  process(u, v, [&](int 1, int r) { tree->add(1, r, val); });
int queryPath(int u, int v) { // Modify depending on problem
  int res = -1e9;
  process(u, v, [&] (int 1, int r) {
     res = max(res, tree->query(1, r));
  return res;
int querySubtree(int v) { // modifySubtree is similar
  return tree->query(pos[v] + VALS_EDGES, pos[v] + siz[v]);
```

```
};
```

LinkCutTree.h

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

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

```
struct Node { // Splay tree. Root's pp contains tree's parent.
 Node *p = 0, *pp = 0, *c[2];
 bool flip = 0;
 Node() { c[0] = c[1] = 0; fix(); }
 void fix() {
   if (c[0]) c[0]->p = this;
   if (c[1]) c[1]->p = this;
    // (+ update sum of subtree elements etc. if wanted)
 void pushFlip() {
   if (!flip) return;
    flip = 0; swap(c[0], c[1]);
   if (c[0]) c[0]->flip ^= 1;
   if (c[1]) c[1]->flip ^= 1;
 int up() { return p ? p->c[1] == this : -1; }
 void rot(int i, int b) {
   int h = i ^ b;
   Node *x = c[i], *v = b == 2 ? x : x -> c[h], *z = b ? v : 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 (pushFlip(); p; ) {
     if (p->p) p->p->pushFlip();
     p->pushFlip(); pushFlip();
     int c1 = up(), c2 = p->up();
     if (c2 == -1) p->rot(c1, 2);
      else p->p->rot(c2, c1 != c2);
 Node* first() {
   pushFlip();
   return c[0] ? c[0]->first() : (splay(), this);
};
struct LinkCut {
 vector<Node> node;
 LinkCut(int N) : node(N) {}
 void link(int u, int v) { // add an edge (u, v)
   assert(!connected(u, v));
   makeRoot(&node[u]);
   node[u].pp = &node[v];
 void cut(int u, int v) { // remove an edge (u, v)
   Node *x = &node[u], *top = &node[v];
   makeRoot(top); x->splay();
   assert(top == (x->pp ?: x->c[0]));
    if (x->pp) x->pp = 0;
    else {
```

```
x \rightarrow c[0] = top \rightarrow p = 0;
      x->fix();
  bool connected(int u, int v) { // are u, v in the same tree?
   Node* nu = access(&node[u])->first();
    return nu == access(&node[v])->first();
  void makeRoot(Node* u) {
    access (II):
    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;
};
```

DirectedMST.h

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

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

39e620, 60 lines

```
"../data-structures/UnionFindRollback.h"
struct Edge { int a, b; ll w; };
struct Node {
  Edge kev;
  Node *1, *r;
  11 delta:
  void prop() {
   kev.w += delta:
   if (1) 1->delta += delta;
   if (r) r->delta += delta;
   delta = 0:
  Edge top() { prop(); return key; }
Node *merge(Node *a, Node *b) {
 if (!a || !b) return a ?: b;
  a->prop(), b->prop();
  if (a->key.w > b->key.w) swap(a, b);
  swap (a->1, (a->r = merge(b, a->r)));
  return a;
void pop (Node \star \& a) { a->prop(); a = merge(a->1, a->r); }
pair<ll, vi> dmst(int n, int r, vector<Edge>& g) {
  RollbackUF uf(n);
  vector<Node*> heap(n);
  for (Edge e : g) heap[e.b] = merge(heap[e.b], new Node{e});
  11 \text{ res} = 0;
  vi seen(n, -1), path(n), par(n);
  seen[r] = r;
  vector<Edge> Q(n), in(n, \{-1,-1\}), comp;
  deque<tuple<int, int, vector<Edge>>> cycs;
  rep(s, 0, n) {
   int u = s, qi = 0, w;
```

```
while (seen[u] < 0) {</pre>
    if (!heap[u]) return {-1,{}};
    Edge e = heap[u]->top();
    heap[u]->delta -= e.w, pop(heap[u]);
    Q[qi] = e, path[qi++] = u, seen[u] = s;
    res += e.w, u = uf.find(e.a);
    if (seen[u] == s) {
      Node* cvc = 0;
      int end = qi, time = uf.time();
      do cyc = merge(cyc, heap[w = path[--qi]]);
      while (uf.join(u, w));
      u = uf.find(u), heap[u] = cyc, seen[u] = -1;
      cycs.push_front({u, time, {&Q[qi], &Q[end]}});
  rep(i, 0, qi) in[uf.find(Q[i].b)] = Q[i];
for (auto& [u,t,comp] : cycs) { // restore sol (optional)
  uf.rollback(t);
  Edge inEdge = in[u];
  for (auto& e : comp) in[uf.find(e.b)] = e;
  in[uf.find(inEdge.b)] = inEdge;
rep(i,0,n) par[i] = in[i].a;
return {res, par};
```

7.8 Math

7.8.1 Number of Spanning Trees

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

7.8.2 Erdős–Gallai theorem

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

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

Geometry (8)

8.1 Geometric primitives

Point.h

Description: Class to handle points in the plane. T can be e.g. double or long long. (Avoid int.) 47ec0a, 28 lines

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

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

lineDistance.h

Description:

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



f6bf6b, 4 lines

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

SegmentDistance.h

Description:

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

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



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

SegmentIntersection.h

Description:

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



```
Usage: vector<P> inter = segInter(s1,e1,s2,e2);
if (sz(inter) == 1)
cout << "segments intersect at " << inter[0] << endl;</pre>
"Point.h", "OnSegment.h"
template < class P > vector < P > segInter (P a, P b, P c, P d) {
  auto oa = c.cross(d, a), ob = c.cross(d, b),
       oc = a.cross(b, c), od = a.cross(b, d);
```

// Checks if intersection is single non-endpoint point.

```
if (sgn(oa) * sgn(ob) < 0 && sgn(oc) * sgn(od) < 0)
  return { (a * ob - b * oa) / (ob - oa) };
set<P> s;
if (onSegment(c, d, a)) s.insert(a);
if (onSegment(c, d, b)) s.insert(b);
if (onSegment(a, b, c)) s.insert(c);
if (onSegment(a, b, d)) s.insert(d);
return {all(s)};
```

lineIntersection.h

Description:

If a unique intersection point of the lines going through s1,e1 and s2,e2 exists {1, point} is returned. If no intersection point exists $\{0, (0,0)\}$ is returned and if infinitely many exists $\{-1, e^2\}$ (0,0)} is returned. The wrong position will be returned if P is Point<|l> and the intersection point does not have integer coordinates. Products of three coordinates are used in inter- \(^{\sigma}\) mediate steps so watch out for overflow if using int or ll.



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

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

sideOf.h

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

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

```
3af81c, 9 lines
template<class P>
int sideOf(P s, P e, P p) { return sgn(s.cross(e, p)); }
template<class P>
int sideOf(const P& s, const P& e, const P& p, double eps) {
 auto a = (e-s).cross(p-s);
 double 1 = (e-s).dist()*eps;
 return (a > 1) - (a < -1);
```

OnSegment.h

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

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

linearTransformation.h Description:

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



```
03a306, 6 lines
typedef Point < double > P;
```

```
P linearTransformation (const P& p0, const P& p1,
    const P& q0, const P& q1, const P& r) {
```

```
P dp = p1-p0, dq = q1-q0, num(dp.cross(dq), dp.dot(dq));
return q0 + P((r-p0).cross(num), (r-p0).dot(num))/dp.dist2();
```

Angle.h

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

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

```
struct Angle {
  int x, y;
  int t;
  Angle(int x, int y, int t=0) : x(x), y(y), t(t) {}
  Angle operator-(Angle b) const { return {x-b.x, y-b.y, t}; }
  int half() const {
    assert(x || y);
    return y < 0 || (y == 0 && x < 0);
  Angle t90() const { return \{-y, x, t + (half() \&\& x \ge 0)\}; }
  Angle t180() const { return \{-x, -y, t + half()\}; }
  Angle t360() const { return {x, y, t + 1}; }
bool operator<(Angle a, Angle b) {</pre>
  // add a.dist2() and b.dist2() to also compare distances
  return make_tuple(a.t, a.half(), a.y * (11)b.x) <</pre>
         make_tuple(b.t, b.half(), a.x * (ll)b.y);
// Given two points, this calculates the smallest angle between
// them, i.e., the angle that covers the defined line segment.
pair<Angle, Angle> segmentAngles(Angle a, Angle b) {
  if (b < a) swap(a, b);
  return (b < a.t180() ?
          make_pair(a, b) : make_pair(b, a.t360()));
Angle operator+(Angle a, Angle b) { // point a + vector b
  Angle r(a.x + b.x, a.y + b.y, a.t);
  if (a.t180() < r) r.t--;
  return r.t180() < a ? r.t360() : r;
Angle angleDiff(Angle a, Angle b) { // angle b - angle a
  int tu = b.t - a.t; a.t = b.t;
  return \{a.x*b.x + a.y*b.y, a.x*b.y - a.y*b.x, tu - (b < a)\};
```

8.2 Circles

CircleIntersection.h

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

```
"Point.h"
                                                      84d6d3, 11 lines
typedef Point<double> P;
bool circleInter(P a, P b, double r1, double r2, pair < P, P >* out) {
  if (a == b) { assert(r1 != r2); return false; }
  P \text{ vec} = b - a;
  double d2 = vec.dist2(), sum = r1+r2, dif = r1-r2,
         p = (d2 + r1*r1 - r2*r2)/(d2*2), h2 = r1*r1 - p*p*d2;
  if (sum*sum < d2 || dif*dif > d2) return false;
  P mid = a + vec*p, per = vec.perp() * sqrt(fmax(0, h2) / d2);
  *out = {mid + per, mid - per};
  return true;
```

CircleTangents.h

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

```
"Point.h"
```

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

CirclePolygonIntersection.h

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

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

```
"../../content/geometry/Point.h"
                                                                         alee63, 19 lines
```

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

circumcircle.h

Description:

"Point.h"

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



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

7cf45b, 39 lines

MinimumEnclosingCircle.h

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

8.3 Polygons

InsidePolygon.h

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

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

"Point.h", "OnSegment.h", "SegmentDistance.h" 2bf504, 11 lines

```
template < class P >
bool inPolygon(vector < P > & p, P a, bool strict = true) {
  int cnt = 0, n = sz(p);
  rep(i,0,n) {
    P q = p[(i + 1) % n];
    if (onSegment(p[i], q, a)) return !strict;
        //or: if (segDist(p[i], q, a) < = eps) return !strict;
    cnt ^= ((a.y < p[i].y) - (a.y < q.y)) * a.cross(p[i], q) > 0;
  }
  return cnt;
}
```

PolygonArea.h

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

"Point.h"

f12300, 6 lines

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

PolygonCenter.h

Description: Returns the center of mass for a polygon.

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

"Point.h"

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

PolygonCut.h

Description:

Returns a vector with the vertices of a polygon with everything to the left of the line going from s to e cut away.

```
Usage: vector<P> p = ...;
p = polygonCut(p, P(0,0), P(1,0));
```



```
"Point.h", "lineIntersection.h" f2b7d4, 13 lines
typedef Point<double> P;
vector<P> polygonCut(const vector<P>& poly, P s, P e) {
  vector<P> res;
  rep(i,0,sz(poly)) {
    P cur = poly[i], prev = i ? poly[i-1] : poly.back();
  bool side = s.cross(e, cur) < 0;
  if (side != (s.cross(e, prev) < 0))
    res.push_back(lineInter(s, e, cur, prev).second);
  if (side)
    res.push_back(cur);
}
return res;</pre>
```

ConvexHull.h

Description:

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



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

HullDiameter.h

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

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

9706dc, 9 lines

oint.h" c571b8, 12 lines

```
typedef Point<11> P;
array<P, 2> hullDiameter(vector<P> S) {
  int n = sz(S), j = n < 2 ? 0 : 1;
  pair<11, array<P, 2>> res({0, {S[0], S[0]}});
  rep(i,0,j)
  for (;; j = (j + 1) % n) {
    res = max(res, {(S[i] - S[j]).dist2(), {S[i], S[j]}});
    if ((S[(j + 1) % n] - S[j]).cross(S[i + 1] - S[i]) >= 0)
        break;
  }
  return res.second;
}
```

PointInsideHull.h

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

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

```
"Point.h", "sideOf.h", "OnSegment.h" 71446b, 14 lines
```

```
typedef Point<11> P;
```

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

LineHullIntersection.h

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

```
#define cmp(i, j) sqn(dir.perp().cross(poly[(i)%n]-poly[(j)%n]))
#define extr(i) cmp(i + 1, i) >= 0 && cmp(i, i - 1 + n) < 0
template <class P> int extrVertex(vector<P>& poly, P dir) {
 int n = sz(poly), lo = 0, hi = n;
  if (extr(0)) return 0;
  while (10 + 1 < hi) {
    int m = (10 + hi) / 2;
    if (extr(m)) return m;
    int ls = cmp(lo + 1, lo), ms = cmp(m + 1, m);
    (1s < ms \mid | (1s == ms \&\& 1s == cmp(1o, m)) ? hi : 1o) = m;
  return lo;
#define cmpL(i) sgn(a.cross(poly[i], b))
template <class P>
array<int, 2> lineHull(P a, P b, vector<P>& poly) {
 int endA = extrVertex(poly, (a - b).perp());
  int endB = extrVertex(poly, (b - a).perp());
  if (cmpL(endA) < 0 || cmpL(endB) > 0)
    return {-1, -1};
  array<int, 2> res;
  rep(i, 0, 2) {
    int lo = endB, hi = endA, n = sz(poly);
    while ((lo + 1) % n != hi) {
      int m = ((lo + hi + (lo < hi ? 0 : n)) / 2) % n;</pre>
      (cmpL(m) == cmpL(endB) ? lo : hi) = m;
    res[i] = (lo + !cmpL(hi)) % n;
    swap (endA, endB);
 if (res[0] == res[1]) return {res[0], -1};
 if (!cmpL(res[0]) && !cmpL(res[1]))
    switch ((res[0] - res[1] + sz(poly) + 1) % sz(poly)) {
      case 0: return {res[0], res[0]};
      case 2: return {res[1], res[1]};
 return res;
```

8.4 Misc. Point Set Problems

```
ClosestPair.h
```

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

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

ac41a6, 17 lines

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

kdTree.h

Description: KD-tree (2d, can be extended to 3d)

bac5b0, 63 lines typedef long long T; typedef Point<T> P; const T INF = numeric limits<T>::max(); bool on_x(const P& a, const P& b) { return a.x < b.x; }</pre> bool on_y(const P& a, const P& b) { return a.y < b.y; }</pre> struct Node { P pt; // if this is a leaf, the single point in it

```
T x\theta = INF, x1 = -INF, y\theta = INF, y1 = -INF; // bounds
Node *first = 0, *second = 0;
T distance (const P& p) { // min squared distance to a point
  T x = (p.x < x0 ? x0 : p.x > x1 ? x1 : p.x);
  T y = (p.y < y0 ? y0 : p.y > y1 ? y1 : p.y);
  return (P(x,y) - p).dist2();
Node (vector<P>&& vp) : pt(vp[0]) {
  for (P p : vp) {
    x\theta = min(x\theta, p.x); x1 = max(x1, p.x);
    y0 = min(y0, p.y); y1 = max(y1, p.y);
```

```
if (vp.size() > 1) {
     // split on x if width >= height (not ideal...)
     sort (all (vp), x1 - x0 >= y1 - y0 ? on_x : on_y);
     // divide by taking half the array for each child (not
     // best performance with many duplicates in the middle)
     int half = sz(vp)/2;
     first = new Node({vp.begin(), vp.begin() + half});
      second = new Node({vp.begin() + half, vp.end()});
};
struct KDTree {
 Node* root:
  KDTree(const vector<P>& vp) : root(new Node({all(vp)})) {}
  pair<T, P> search(Node *node, const P& p) {
   if (!node->first) {
      // uncomment if we should not find the point itself:
```

```
// if (p == node->pt) return {INF, P()};
     return make_pair((p - node->pt).dist2(), node->pt);
   Node *f = node->first, *s = node->second;
   T bfirst = f->distance(p), bsec = s->distance(p);
    if (bfirst > bsec) swap(bsec, bfirst), swap(f, s);
    // search closest side first, other side if needed
    auto best = search(f, p);
    if (bsec < best.first)</pre>
     best = min(best, search(s, p));
   return best:
 // find nearest point to a point, and its squared distance
 // (requires an arbitrary operator< for Point)</pre>
 pair<T, P> nearest (const P& p) {
   return search(root, p);
};
```

FastDelaunav.h

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

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

```
"Point.h"
                                                         eefdf5, 88 lines
typedef Point<11> P;
typedef struct Quad* Q;
typedef __int128_t 111; // (can be 11 if coords are < 2e4)</pre>
P arb(LLONG MAX, LLONG MAX); // not equal to any other point
struct Onad {
  Q rot, o; P p = arb; bool mark;
 P& F() { return r()->p; }
  O& r() { return rot->rot; }
  0 prev() { return rot->o->rot; }
  Q next() { return r()->prev(); }
bool circ(P p, P a, P b, P c) { // is p in the circumcircle?
  111 p2 = p.dist2(), A = a.dist2()-p2,
      B = b.dist2()-p2, C = c.dist2()-p2;
  return p.cross(a,b) *C + p.cross(b,c) *A + p.cross(c,a) *B > 0;
Q makeEdge(P orig, P dest) {
  Q r = H ? H : new Quad{new Quad{new Quad{new Quad{0}}}};
  H = r -> 0; r -> r() -> r() = r;
  rep(i, 0, 4) r = r \rightarrow rot, r \rightarrow p = arb, r \rightarrow o = i & 1 ? <math>r : r \rightarrow r();
  r->p = orig; r->F() = dest;
  return r;
void splice(Q a, Q b) {
  swap(a->o->rot->o, b->o->rot->o); swap(a->o, b->o);
Q connect(Q a, Q b) {
  Q = makeEdge(a->F(), b->p);
  splice(q, a->next());
  splice(q->r(), b);
  return q;
pair<Q,Q> rec(const vector<P>& s) {
  if (sz(s) <= 3) {
```

Q = makeEdge(s[0], s[1]), b = makeEdge(s[1], s.back());

if (sz(s) == 2) **return** { a, a->r() };

```
splice(a->r(), b);
    auto side = s[0].cross(s[1], s[2]);
    0 c = side ? connect(b, a) : 0;
    return {side < 0 ? c->r() : a, side < 0 ? c : b->r() };
#define H(e) e->F(), e->p
#define valid(e) (e->F().cross(H(base)) > 0)
 Q A, B, ra, rb;
 int half = sz(s) / 2;
 tie(ra, A) = rec({all(s) - half});
 tie(B, rb) = rec({sz(s) - half + all(s)});
  while ((B->p.cross(H(A)) < 0 && (A = A->next())) ||
         (A->p.cross(H(B)) > 0 && (B = B->r()->o)));
  Q base = connect(B->r(), A);
  if (A->p == ra->p) ra = base->r();
  if (B->p == rb->p) rb = base;
#define DEL(e, init, dir) Q e = init->dir; if (valid(e)) \
    while (circ(e->dir->F(), H(base), e->F())) {
     Q t = e->dir; \setminus
      splice(e, e->prev()); \
      splice(e->r(), e->r()->prev()); \
      e->0 = H; H = e; e = t; \setminus
  for (;;) {
    DEL(LC, base->r(), o); DEL(RC, base, prev());
    if (!valid(LC) && !valid(RC)) break;
    if (!valid(LC) || (valid(RC) && circ(H(RC), H(LC))))
     base = connect(RC, base->r());
      base = connect(base->r(), LC->r());
  return { ra, rb };
vector<P> triangulate(vector<P> pts) {
  sort(all(pts)); assert(unique(all(pts)) == pts.end());
 if (sz(pts) < 2) return {};
 Q e = rec(pts).first;
 vector<Q> q = \{e\};
 int qi = 0;
  while (e->o->F().cross(e->F(), e->p) < 0) e = e->o;
#define ADD { 0 c = e; do { c \rightarrow mark = 1; pts.push back(c \rightarrow p); \
  q.push_back(c->r()); c = c->next(); } while (c != e); }
  ADD; pts.clear();
  while (qi < sz(q)) if (!(e = q[qi++]) \rightarrow mark) ADD;
  return pts;
```

$8.5 \quad 3D$

PolyhedronVolume.h

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

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

Point3D.h

Description: Class to handle points in 3D space. T can be e.g. double or long long.

```
template < class T > struct Point3D {
 typedef Point3D P;
 typedef const P& R;
```

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

3dHull.h

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

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

```
"Point3D.h" 5b45fc, 49 lines
```

```
typedef Point3D<double> P3;
struct PR {
  void ins(int x) { (a == -1 ? a : b) = x; }
  void rem(int x) { (a == x ? a : b) = -1; }
  int cnt() { return (a !=-1) + (b !=-1); }
  int a, b;
struct F { P3 q; int a, b, c; };
vector<F> hull3d(const vector<P3>& A) {
  assert (sz(A) >= 4);
  vector<vector<PR>>> E(sz(A), vector<PR>(sz(A), {-1, -1}));
#define E(x,y) E[f.x][f.y]
  vector<F> FS:
  auto mf = [&](int i, int j, int k, int l) {
   P3 q = (A[j] - A[i]).cross((A[k] - A[i]));
   if (q.dot(A[1]) > q.dot(A[i]))
     q = q * -1;
   F f{q, i, j, k};
   E(a,b).ins(k); E(a,c).ins(j); E(b,c).ins(i);
   FS.push back(f);
  rep(i,0,4) rep(j,i+1,4) rep(k,j+1,4)
   mf(i, j, k, 6 - i - j - k);
  rep(i, 4, sz(A)) {
    rep(j, 0, sz(FS)) {
     F f = FS[j];
     if(f.q.dot(A[i]) > f.q.dot(A[f.a])) {
       E(a,b).rem(f.c);
       E(a,c).rem(f.b);
```

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

sphericalDistance.h

Description: Returns the shortest distance on the sphere with radius radius between the points with azimuthal angles (longitude) f1 (ϕ_1) and f2 (ϕ_2) from x axis and zenith angles (latitude) t1 (θ_1) and t2 (θ_2) from z axis (0 = north pole). All angles measured in radians. The algorithm starts by converting the spherical coordinates to cartesian coordinates so if that is what you have you can use only the two last rows. dx*radius is then the difference between the two points in the x direction and d*radius is the total distance between the points.

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

$\underline{\text{Strings}}$ (9)

KMP h

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

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

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

Zfunc.h

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

```
Time: O(n)

vi Z(const string& S) {

vi Z(sz(S));
```

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

Manacher.h

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

```
Imme: O(N)

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

MinRotation.h

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

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

SuffixArrav.h

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

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

SuffixTree Hashing AhoCorasick IntervalContainer

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

SuffixTree.h

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

Time: $\mathcal{O}\left(26N\right)$

aae0b8, 50 lines

Hashing.h

```
struct SuffixTree {
  enum { N = 200010, ALPHA = 26 }; // N \sim 2*maxlen+10
  int toi(char c) { return c - 'a'; }
  string a; // v = cur node, q = cur position
  int t[N][ALPHA],1[N],r[N],p[N],s[N],v=0,q=0,m=2;
  void ukkadd(int i, int c) { suff:
   if (r[v]<=q) {
     if (t[v][c]==-1) { t[v][c]=m; l[m]=i;
       p[m++]=v; v=s[v]; q=r[v]; goto suff; }
      v=t[v][c]; q=l[v];
    if (g==-1 || c==toi(a[g])) g++; else {
     l[m+1]=i; p[m+1]=m; l[m]=l[v]; r[m]=q;
     p[m]=p[v]; t[m][c]=m+1; t[m][toi(a[q])]=v;
     l[v]=q; p[v]=m; t[p[m]][toi(a[l[m]])]=m;
     v=s[p[m]]; q=l[m];
      while (q < r[m]) \{ v = t[v][toi(a[q])]; q + = r[v] - l[v]; \}
     if (q==r[m]) s[m]=v; else s[m]=m+2;
      q=r[v]-(q-r[m]); m+=2; qoto suff;
  SuffixTree(string a) : a(a) {
    fill(r,r+N,sz(a));
   memset(s, 0, sizeof s);
    memset(t, -1, sizeof t);
    fill(t[1],t[1]+ALPHA,0);
    s[0] = 1; 1[0] = 1[1] = -1; r[0] = r[1] = p[0] = p[1] = 0;
    rep(i,0,sz(a)) ukkadd(i, toi(a[i]));
  // example: find longest common substring (uses ALPHA = 28)
  pii best;
  int lcs(int node, int i1, int i2, int olen) {
   if (l[node] <= i1 && i1 < r[node]) return 1;</pre>
   if (1[node] <= i2 && i2 < r[node]) return 2;</pre>
    int mask = 0, len = node ? olen + (r[node] - 1[node]) : 0;
    rep(c, 0, ALPHA) if (t[node][c] != -1)
     mask |= lcs(t[node][c], i1, i2, len);
    if (mask == 3)
     best = max(best, {len, r[node] - len});
    return mask;
  static pii LCS(string s, string t) {
    SuffixTree st(s + (char) ('z' + 1) + t + (char) ('z' + 2));
    st.lcs(0, sz(s), sz(s) + 1 + sz(t), 0);
    return st.best;
};
```

```
Description: Self-explanatory methods for string hashing.
// Arithmetic mod 2^64-1. 2x slower than mod 2^64 and more
// code, but works on evil test data (e.g. Thue-Morse, where
// ABBA... and BAAB... of length 2^10 hash the same mod 2^64).
// "typedef ull H;" instead if you think test data is random,
// or work mod 10^9+7 if the Birthday paradox is not a problem.
typedef uint64 t ull;
struct H {
 ull x; H(ull x=0) : x(x) \{ \}
  H operator+(H \circ) { return x + \circ.x + (x + \circ.x < x); }
  H operator-(H o) { return *this + ~o.x; }
 H operator*(H o) { auto m = (__uint128_t)x * o.x;
    return H((ull)m) + (ull)(m >> 64); }
  ull get() const { return x + !~x; }
  bool operator==(H o) const { return get() == o.get(); }
 bool operator<(H o) const { return get() < o.get(); }</pre>
static const H C = (11)1e11+3; // (order \sim 3e9; random also ok)
struct HashInterval {
  vector<H> ha, pw;
 HashInterval(string& str) : ha(sz(str)+1), pw(ha) {
    pw[0] = 1;
    rep(i, 0, sz(str))
     ha[i+1] = ha[i] * C + str[i],
      pw[i+1] = pw[i] * C;
 H hashInterval(int a, int b) { // hash [a, b)
    return ha[b] - ha[a] * pw[b - a];
};
vector<H> getHashes(string& str, int length) {
 if (sz(str) < length) return {};</pre>
 H h = 0, pw = 1;
 rep(i,0,length)
   h = h * C + str[i], pw = pw * C;
  vector<H> ret = {h};
 rep(i,length,sz(str)) {
    ret.push_back(h = h * C + str[i] - pw * str[i-length]);
 return ret;
H hashString(string& s){H h{}; for(char c:s) h=h*C+c;return h;}
```

AhoCorasick.h

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

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

```
struct AhoCorasick {
  enum {alpha = 26, first = 'A'}; // change this!
  struct Node {
    // (nmatches is optional)
    int back, next[alpha], start = -1, end = -1, nmatches = 0;
    Node(int v) { memset(next, v, sizeof(next)); }
};
  vector<Node> N;
  vi backp;
```

```
void insert(string& s, int j) {
  assert(!s.empty());
  int n = 0;
  for (char c : s) {
    int& m = N[n].next[c - first];
    if (m == -1) { n = m = sz(N); N.emplace_back(-1); }
    else n = m;
  if (N[n].end == -1) N[n].start = j;
  backp.push_back(N[n].end);
  N[n].end = j;
  N[n].nmatches++;
AhoCorasick(vector<string>& pat) : N(1, -1) {
  rep(i,0,sz(pat)) insert(pat[i], i);
  N[0].back = sz(N);
  N.emplace_back(0);
  queue<int> q;
  for (q.push(0); !q.empty(); q.pop()) {
    int n = q.front(), prev = N[n].back;
    rep(i,0,alpha) {
      int &ed = N[n].next[i], y = N[prev].next[i];
      if (ed == -1) ed = y;
      else {
        N[ed].back = y;
        (N[ed].end == -1 ? N[ed].end : backp[N[ed].start])
          = N[v].end;
        N[ed].nmatches += N[y].nmatches;
        q.push(ed);
vi find(string word) {
  int n = 0;
  vi res; // 11 count = 0;
  for (char c : word) {
    n = N[n].next[c - first];
    res.push_back(N[n].end);
    // count += N[n].nmatches;
  return res;
vector<vi> findAll(vector<string>& pat, string word) {
  vi r = find(word);
  vector<vi> res(sz(word));
  rep(i, 0, sz(word)) {
    int ind = r[i];
    while (ind !=-1) {
      res[i - sz(pat[ind]) + 1].push_back(ind);
      ind = backp[ind];
  return res;
```

Various (10)

10.1 Intervals

IntervalContainer.h

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

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

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

IntervalCover.h

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

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

9e9d8d, 19 lines

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

ConstantIntervals.h

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

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

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

10.2 Misc. algorithms

TernarySearch.h

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

Usage: int ind = ternSearch(0,n-1,[&](int i){return a[i];}); Time: $\mathcal{O}(\log(b-a))$

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

LIS.h

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

```
2932a0, 17 lines
template < class I > vi lis(const vector < I > & S) {
 if (S.emptv()) return {};
 vi prev(sz(S));
 typedef pair<I, int> p;
 vector res;
 rep(i, 0, sz(S)) {
   // change 0 -> i for longest non-decreasing subsequence
   auto it = lower_bound(all(res), p{S[i], 0});
   if (it == res.end()) res.emplace_back(), it = res.end()-1;
   *it = {S[i], i};
   prev[i] = it == res.begin() ? 0 : (it-1) -> second;
 int L = sz(res), cur = res.back().second;
 vi ans(L);
 while (L--) ans[L] = cur, cur = prev[cur];
 return ans:
```

FastKnapsack.h

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

Time: $O(N \max(w_i))$

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

rep(x,0,m) v[x+w[i]] = max(v[x+w[i]], u[x]);

10.3 Dynamic programming

KnuthDP.h

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

DivideAndConquerDP.h

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

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

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

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

void solve(int L, int R) { rec(L, R, INT_MIN, INT_MAX); }
};
```

10.4 Debugging tricks

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

10.5 Optimization tricks

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

10.5.1 Bit backs

b20ccc, 16 lines

- 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; $(((r^x) >> 2)/c) \mid r$ is the next number after x with the same number of bits set.
- rep(b,0,K) rep(i,0,(1 << K))
 if (i & 1 << b) D[i] += D[i^(1 << b)];
 computes all sums of subsets.</pre>

10.5.2 Pragmas

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

FastMod.h

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

```
typedef unsigned long long ull;
struct FastMod {
  ull b, m;
  FastMod(ull b) : b(b), m(-IULL / b) {}
  ull reduce(ull a) { // a % b + (0 or b)
    return a - (ull)((__uint128_t(m) * a) >> 64) * b;
  }
};
```

FastInput.h

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

Usage: ./a.out < input.txt</pre>

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

7b3c70, 17 lines

```
inline char gc() { // like getchar()
    static char buf[1 << 16];
    static size_t bc, be;
    if (bc >= be) {
        buf[0] = 0, bc = 0;
        be = fread(buf, 1, sizeof(buf), stdin);
    }
    return buf[bc++]; // returns 0 on EOF
}

int readInt() {
    int a, c;
    while ((a = gc()) < 40);
    if (a == '-') return -readInt();
    while ((c = gc()) >= 48) a = a * 10 + c - 480;
    return a - 48;
}
```

BumpAllocator.h

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

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

```
void operator delete(void*) {}
SmallPtr.h
Description: A 32-bit pointer that points into BumpAllocator memory.
template<class T> struct ptr {
 unsigned ind;
 ptr(T*p = 0) : ind(p ? unsigned((char*)p - buf) : 0) {
   assert (ind < sizeof buf);
 T& operator*() const { return *(T*)(buf + ind); }
 T* operator->() const { return &**this; }
 T& operator[](int a) const { return (&**this)[a]; }
 explicit operator bool() const { return ind; }
BumpAllocatorSTL.h
Description: BumpAllocator for STL containers.
Usage: vector<vector<int, small<int>>> ed(N);
                                                     bb66d4, 14 lines
char buf[450 << 20] alignas(16);</pre>
size t buf ind = sizeof buf;
template < class T > struct small {
 typedef T value type;
 small() {}
 template < class U > small(const U&) {}
 T* allocate(size t n) {
```

SIMD.h

};

buf ind -= n * sizeof(T);

buf_ind &= 0 - alignof(T);
return (T*) (buf + buf ind);

void deallocate(T*, size t) {}

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

```
551b82, 43 lines
#pragma GCC target ("avx2") // or sse4.1
#include "immintrin.h"
typedef __m256i mi;
#define L(x) _mm256_loadu_si256((mi*)&(x))
// High-level/specific methods:
// load(u)?_si256, store(u)?_si256, setzero_si256, _mm_malloc
// blendv_(epi8|ps|pd) (z?y:x), movemask_epi8 (hibits of bytes)
// i32gather_epi32(addr, x, 4): map addr[] over 32-b parts of x
// sad_epu8: sum of absolute differences of u8, outputs 4xi64
// maddubs_epi16: dot product of unsigned i7's, outputs 16xi15
// madd_epi16: dot product of signed i16's, outputs 8xi32
// extractf128_si256(, i) (256->128), cvtsi128_si32 (128->1032)
// permute2f128_si256(x,x,1) swaps 128-bit lanes
// shuffle_epi32(x, 3*64+2*16+1*4+0) == x for each lane
// shuffle_epi8(x, y) takes a vector instead of an imm
// Methods that work with most data types (append e.g. _epi32):
// set1, blend (i8?x:y), add, adds (sat.), mullo, sub, and/or,
// andnot, abs, min, max, sign(1,x), cmp(gt|eg), unpack(lo|hi)
```

```
int sumi32(mi m) { union {int v[8]; mi m;} u; u.m = m;
  int ret = 0; rep(i,0,8) ret += u.v[i]; return ret; }
mi zero() { return _mm256_setzero_si256(); }
mi one() { return _mm256_set1_epi32(-1); }
bool all_zero(mi m) { return _mm256_testz_si256(m, m); }
bool all_one(mi m) { return _mm256_testc_si256(m, one()); }
ll example_filteredDotProduct(int n, short* a, short* b) {
  int i = 0; 11 r = 0;
  mi zero = _mm256_setzero_si256(), acc = zero;
  while (i + 16 <= n) {
    mi \ va = L(a[i]), \ vb = L(b[i]); \ i += 16;
    va = _mm256_and_si256(_mm256_cmpgt_epi16(vb, va), va);
    mi vp = _mm256_madd_epi16(va, vb);
    acc = _mm256_add_epi64(_mm256_unpacklo_epi32(vp, zero),
      _mm256_add_epi64(acc, _mm256_unpackhi_epi32(vp, zero)));
  union {ll v[4]; mi m;} u; u.m = acc; rep(i,0,4) r += u.v[i];
  for (;i<n;++i) if (a[i] < b[i]) r += a[i]*b[i]; // <- equiv</pre>
  return r:
```

Techniques (A)

techniques.txt

Combinatorics

159 lines

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

Computation of binomial coefficients Pigeon-hole principle Inclusion/exclusion Catalan number Pick's theorem Number theory Integer parts Divisibility Euclidean algorithm Modular arithmetic * Modular multiplication * Modular inverses * Modular exponentiation by squaring Chinese remainder theorem Fermat's little theorem Euler's theorem Phi function Frobenius number 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 polynomial hashes Suffix array Suffix tree Aho-Corasick Manacher's algorithm Letter position lists Combinatorial search Meet in the middle Brute-force with pruning Best-first (A*) Bidirectional search Iterative deepening DFS / A* Data structures LCA (2^k-jumps in trees in general) Pull/push-technique on trees Heavy-light decomposition Centroid decomposition Lazy propagation Self-balancing trees Convex hull trick (wcipeg.com/wiki/Convex_hull_trick) Monotone queues / monotone stacks / sliding queues Sliding queue using 2 stacks Persistent segment tree

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