

Universidade de Brasilia

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1 Contest	1 .bashrc
2 Mathematics	alias comp='g++ -std=c++17 -g3 -ggdb3 -03 -Wall -Wextra - fsanitize=address,undefined -Wshadow -Wconversion - D_GLIBCXX_ASSERTIONS -o test'
3 Data structures	3
	hash.sh 3 lines
4 Numerical	# Hashes a file, ignoring all whitespace and comments. Use for
5 Number theory	# verifying that code was correctly typed. CTRL+D to send EOF cpp -dD -P -fpreprocessed tr -d '[:space:]' md5sum cut -c -6
6 Combinatorial	3
	troubleshoot.txt 52 lines
7 Graph	Pre-submit:
8 Geometry	Write a few simple test cases if sample is not enough. Are time limits close? If so, generate max cases. Is the memory usage fine?
9 Strings	Could anything overflow? Make sure to submit the right file.
40351	Wrong answer:
10 Miscellaneous	Print your solution! Print debug output, as well.
Contest (1)	Are you clearing all data structures between test cases? Can your algorithm handle the whole range of input? Read the full problem statement again. Do you handle all corner cases correctly? Have you understood the problem correctly?
template.cpp 33 line	Any uninitialized variables?
#include <bits stdc++.h=""></bits>	<pre>Many overflows? Confusing N and M, i and j, etc.?</pre>
<pre>using namespace std; #define sws cin.tie(0)->sync_with_stdio(0)</pre>	Are you sure your algorithm works? What special cases have you not thought of?
	Are you sure the STL functions you use work as you think?
#define endl '\n' #define ll long long	Add some assertions, maybe resubmit. Create some testcases to run your algorithm on.
#define ld long double	Go through the algorithm for a simple case.
#define pb push_back	Go through this list again.
<pre>#define ff first #define ss second</pre>	Explain your algorithm to a teammate. Ask the teammate to look at your code.
#define pll pair <ll, 11=""></ll,>	Go for a small walk, e.g. to the toilet.
<pre>#define vll vector<11></pre>	Is your output format correct? (including whitespace) Rewrite your solution from the start or let a teammate do it.
#define teto(a, b) ((a+b-1)/(b))	Runtime error:
#define LSB(i) ((i) & -(i)) #define MSB(i) (32builtin_clz(i)) //64 - clzll	Have you tested all corner cases locally?
<pre>#define BITS(i)builtin_popcountll(i) //count set bits</pre>	Any uninitialized variables?
<pre>mt19937 rng(chrono::steady_clock::now().time_since_epoch().</pre>	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?
<pre>#define debug(a) cerr<<#a<<": ";for(auto b:a)cerr<<b<<" ";="" cerr<<endl;<="" pre=""></b<<"></pre>	Invalidated pointers or iterators? Are you using too much memory?
<pre>template<typename a=""> void dbg(A const& a) { ((cerr<<"{"<<a "),);="" <"}="" cerr<<endl;="" pre="" }<=""></typename></pre>	Debug with resubmits (e.g. remapped signals, see Various).
	Time limit exceeded:
<pre>const int MAX = 3e5+10; const int INF = INT32_MAX;</pre>	Do you have any possible infinite loops? What is the complexity of your algorithm?
const long long MOD = 1e9+7;	Are you copying a lot of unnecessary data? (References)
const long long LLINF = INT64_MAX;	How big is the input and output? (consider scanf)
const long double EPS = 1e-7;	Avoid vector, map. (use arrays/unordered_map)
<pre>const long double PI = acos(-1);</pre>	What do your teammates think about your algorithm?
<pre>int32_t main(){ sws;</pre>	Memory limit exceeded: What is the max amount of memory your algorithm should need?
}	Are you clearing all data structures between test cases?

Mathematics (2)

2.1 Equations

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

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

$$ax + by = e$$

$$cx + dy = f$$

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

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

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

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

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

2.2 Recurrences

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

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

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

2.3 Trigonometry

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

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

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

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

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

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

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

Geometry

2.4.1Triangles

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

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):

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

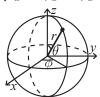
Law of sines: $\frac{\sin \alpha}{a} = \frac{\sin \beta}{b} = \frac{\sin \gamma}{c} = \frac{1}{2R}$ Law of cosines: $a^2 = b^2 + c^2 - 2bc \cos \alpha$

2.4.2 Quadrilaterals $\tan \frac{\alpha + \beta}{2}$ with of the nearths a, b, c, \overline{a} , diagonals e, f, diagonals angle θ , area A and magic flux $F = b^2 + \frac{1}{2}a^2 - \frac{1}{2}c^2 - c^2$:

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

2.4.3 Spherical coordinates

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



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

2.5 Derivatives/Integrals

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

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

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

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

Integration by parts:

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

2.6 Sums

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

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

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

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

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

2.7Series

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

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

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

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

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

2.8 Probability theory

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

Expectation is linear:

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

For independent X and Y,

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

2.8.1 Discrete distributions Binomial distribution

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

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

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

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

First success distribution

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

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

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

Poisson distribution

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

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

$$\mu = \lambda, \, \sigma^2 = \lambda$$

2.8.2 Continuous distributions Uniform distribution

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

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

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

Exponential distribution

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

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

Normal distribution

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

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

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

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

2.9 Markov chains

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

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

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

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

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

Data structures (3)

3.1 Disjoint Set Union

There are two optional improvements: -Tree Balancing -Path Compression $\,$

If one improvement is used, the time complexity will become $O(\log N)$

If both are used, $O(\alpha) \approx O(5)$

dsu.cpp

Description: Disjoint Set Union with path compression and tree balancing Time: $\mathcal{O}\left(\alpha\right)$ 0479c4. 22 lines

```
struct DSU{
    vll group, card;
    DSU (11 n) {
        n += 1; // 0 - i dx \rightarrow 1 - i dx
        group = vll(n);
        iota(group.begin(), group.end(), 0);
        card = vll(n, 1);
    ll find(ll i){
        return (i == group[i]) ? i : (group[i] = find(group[i])
    // returns false if a and b are already in the same
         component
    bool join(ll a , ll b) {
        a = find(a);
        b = find(b);
        if (a == b) return false;
        if (card[a] < card[b]) swap(a, b);</pre>
        card[a] += card[b];
        group[b] = a;
        return true;
};
```

Numerical (4)

Number theory (5)

Combinatorial (6)

6.1 Permutations

6.1.1 Factorial

	_		-		8	-	10	
n!	1 2 6	24 1	20 72	0 5040	40320	362880	3628800	
n	11	12	13	14	15	16	17	
							13 3.6e14	
							0 171	
n!	2e18	2e25	3e32	8e47 3	664 9e	157 6e2	$62 > DBL_MA$	X

Graph (7)

- 7.1 Fundamentals
- 7.2 Network flow
- 7.3 Matching
- 7.4 DFS algorithms
- 7.5 Coloring
- 7.6 Undirected Graph

Bridges and Articulation Points are concepts for undirected graphs!

7.6.1 Bridges (Cut Edges)

Also called **isthmus** or **cut arc**.

A back-edge is never a bridge!

A **lowlink** for a vertice U is the closest vertice to the root reachable using only span edges and a single back-edge, starting in the subtree of U.

After constructing a DFS Tree, an edge (u, v) is a bridge \iff there is no back-edge from v (or a descendent of v) to u (or an ancestor of u)

To do this efficiently, it's used tin[i] (entry time of node i) and low[i] (minimum entry time considering all nodes that can be reached from node i).

In another words, a edge (u, v) is a bridge \iff the low[v] ξ tin[u].

bridges.cpp

Description: Using the concepts of entry time (tin) and lowlink (low), an edge is a bridge if, and only if, low[v] > tin[u]

87e0d3, 25 lines

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

```
vector<vll> g(MAX);
ll timer = 1;
```

```
11 tin[MAX], low[MAX];
vector<pll> bridges;
void dfs(ll u, ll p = -1){
    tin[u] = low[u] = timer++;
    for (auto v : g[u]) if (v != p) {
        if (tin[v]) // v was visited (\{u,v\} \text{ is a back-edge})
            // considering a single back-edge:
            low[u] = min(low[u], tin[v]);
       else { // v wasn't visited ({u, v} is a span-edge)
            dfs(v, u);
            // after low[v] was computed by dfs(v, u):
            low[u] = min(low[u], low[v]);
            if (low[v] > tin[u])
                bridges.pb({u, v});
void findBridges(ll n) {
    for(ll i=1; i<=n; i++) if (!tin[i])
       dfs(i);
```

7.6.2 Bridge Tree

After merging *vertices* of a **2-edge connected component** into single vertices, and leaving only bridges, one can generate a Bridge Tree.

Every **2-edge connected component** has following properties:

• For each pair of vertices A, B inside the same component, there are at least 2 distinct paths from A to B (which may repeat vertices).

bridgeTree.cpp

Description: After finding bridges, set an component id for each vertice, then merge vertices that are in the same 2-edge connected component

```
Time: \mathcal{O}(V+E)
                                                       6d00bd, 47 lines
// g: u \rightarrow \{v, edge id\}
vector<vector<pll>> g(MAX);
vector<vll> gc(MAX);
11 \text{ timer} = 1;
11 tin[MAX], low[MAX], comp[MAX];
bool isBridge[MAX];
void dfs(ll u, ll p = -1) {
    tin[u] = low[u] = timer++;
    for (auto [v, id] : g[u]) if (v != p) {
        if (tin[v])
             low[u] = min(low[u], tin[v]);
        else {
            dfs(v, u);
            low[u] = min(low[u], low[v]);
            if (low[v] > tin[u])
                 isBridge[id] = 1;
void dfs2(11 u, 11 c, 11 p = -1) {
    comp[u] = c;
    for(auto [v, id] : g[u]) if (v != p) {
        if (isBridge[id]) continue;
        if (!comp[v]) dfs2(v, c, u);
```

```
void bridgeTree(ll n) {
    // find bridges
    for(ll i=1; i<=n; i++) if (!tin[i])
        dfs(i);
    // find components
    for(ll i=1; i<=n; i++) if (!comp[i])
        dfs2(i, i);
    // condensate into a TREE (or TREES if disconnected)
    for(11 u=1; u<=n; u++) {
        for(auto [v, id] : g[u]) {
            if (comp[u] != comp[v]) {
                gc[comp[u]].pb(comp[v]);
```

7.6.3 Articulation Points

One Vertice in a graph is considered a Articulation Points or Cut Vertice if its removal in the graph will generate more disconnected components

articulation.cpp

Description: if low[v] >= tin[u], u is an articulation points The root is a corner case Time: $\mathcal{O}(V+E)$

8707a0, 29 lines

```
vector<vll> q(MAX);
11 timer = 1;
11 low[MAX], tin[MAX], isAP[MAX];
// when vertex i is removed from graph
// isAP[i] is the quantity of new disjoint components created
// is AP[i] >= 1 {i is a Articulation Point}
void dfs(ll u, ll p = -1) {
    low[u] = tin[u] = timer++;
    for (auto v : g[u]) if (v != p) {
        if (tin[v]) // visited
            low[u] = min(low[u], tin[v]);
        else { // not visited
            dfs(v, u);
            low[u] = min(low[u], low[v]);
            if (low[v] >= tin[u])
                isAP[u]++;
    // corner case: root
    if (p == -1 \text{ and } isAP[u]) isAP[u]--;
void findAP(ll n) {
    for(ll i=1; i<=n; i++) if (!tin[i])</pre>
        dfs(i);
```

7.6.4 Block Cut Tree

After merging edges of a 2-vertex connected component into single vertices, one can obtain a block cut tree.

2-vertex connected components are also called as biconnected component

Every bridge by itself is a biconnected component

Each edge in the block-cut tree connects exactly an Articulation Point and a biconnected component (bipartite graph)

Each biconnected component has the following properties:

- For each pair of edges, there is a cycle that contains both edges
- For each pair of vertices A, B inside the same connected component, there are at least 2 distinct paths from A to B (which do not repeat vertices).

blockCutTree.cpp

Description: After Merging 2-Vertex Connected Components, one can generate a block cut tree

Time: $\mathcal{O}(V+E)$ f752d5, 100 lines

```
// Block-Cut Tree (bruno monteiro)
// Cria a block-cut tree, uma arvore com os blocos
// e os pontos de articulação
// Blocos sao as componentes 2-vertice-conexos maximais
// Uma 2-coloração da arvore eh tal que uma cor são
// os componentes, e a outra cor sao os pontos de articulação
// Funciona para grafo nao conexo
// isAP[i] responde o numero de novas componentes conexas
// criadas apos a remocao de i do grafo g
// Se isAP[i] >= 1, i eh ponto de articulação
// Para todo i < blocks.size()
// blocks[i] eh uma componente 2-vertce-conexa maximal
// blockEdges[i] sao as arestas do bloco i
// tree eh a arvore block-cut-tree
// tree[i] eh um vertice da arvore que corresponde ao bloco i
// comp[i] responde a qual vertice da arvore vertice i pertence
// Arvore tem no maximo 2n vertices
// O(n+m)
// 0-idx graph!!!
vector<vll> g(MAX), tree, blocks; // 2-vertex-connected-
    COMPonentS
vector<vector<pll>>> blockEdges;
stack<11> st; // st for vertices,
stack<pl1> st2; // st2 for edges
vector<11> low, tin, comp, isAP;
11 \text{ timer} = 1;
void dfs(ll u, ll p = -1) {
    low[u] = tin[u] = timer++;
    st.push(u);
    // add only back-edges to stack
    if (p != -1) st2.push({u, p});
    for (auto v : q[u]) if (v != p)
```

if (tin[v] != -1) // visited

```
st2.push({u, v});
    for (auto v : g[u]) if (v != p) {
        if (tin[v] != -1) // visited
            low[u] = min(low[u], tin[v]);
       else { // not visited
            dfs(v, u);
            low[u] = min(low[u], low[v]);
            if (low[v] >= tin[u]) {
                isAP[u] += 1;
                blocks.pb(vll(1, u));
                while(blocks.back().back() != v)
                    blocks.back().pb(st.top()), st.pop();
                blockEdges.pb(vector<pl1>(1, st2.top())), st2.
                while(blockEdges.back().back() != pair<11, 11>(
                    blockEdges.back().pb(st2.top()), st2.pop();
    // corner case: root
    if (p == -1 \text{ and } isAP[u]) isAP[u]--;
void blockCutTree(ll n) {
    // initialize vectors and reset
    tree.clear(), blocks.clear(), blockEdges.clear();
    st = stack<11>(), st2 = stack<pl1>();
    tin.assign(n, -1);
    low.assign(n, 0), comp.assign(n, 0), isAP.assign(n, 0);
    // find Articulation Points
    for(ll i=0; i<n; i++) if (tin[i] == -1)
        dfs(i);
    // set component id for APs
    tree.assign(blocks.size(), vll());
    for(ll i=0; i<n; i++) if (isAP[i])</pre>
        comp[i] = tree.size(), tree.pb(vll());
    // set component id for non-APs and construct tree
    for(11 u=0; u<(11)blocks.size(); u++) {</pre>
        for(auto v : blocks[u]) {
            if (!isAP[v])
                comp[v] = u;
                tree[u].pb(comp[v]), tree[comp[v]].pb(u);
```

7.6.5 Minimum Spanning Tree

A minimum spanning tree (MST) or minimum weight spanning tree is a subset of the edges of a connected, edge-weighted undirected graph that connects all the vertices together, without any cycles and with the minimum possible total edge weight. That is, it is a spanning tree whose sum of edge weights is as small as possible.

kruskal.cpp

Description: Sort all edges in crescent order by weight, include all edges which joins two disconnected trees. In case of tie, choose whichever. Dont include edges that will join a already connected part of the tree.

Time: $\mathcal{O}\left(E\log E\alpha\right)$ 206ba3, 21 line

```
// use DSU struct
struct DSU{};
set<array<11, 3>> edges;
int32 t main() { sws;
   11 n, m; cin >> n >> m;
   DSU dsu(n+1);
    for(ll i=0; i<m; i++) {</pre>
        11 u, v, w; cin >> u >> v >> w;
        edges.insert({w, u, v});
    11 \min Cost = 0;
    for(auto [w, u, v] : edges) {
        if (dsu.find(u) != dsu.find(v)) {
            dsu.join(u, v);
            minCost += w;
   }
    cout << minCost << endl;
```

7.7 Directed Graph

7.7.1 Topological Sort

Sort a directed graph with no cycles (DAG) in an order which each source of an edge is visited before the sink of this edge.

Cannot have cycles, because it would create a contradition of which vertices whould come before.

It can be done with a DFS, appending in the reverse order of transversal. Also a stack can be used to reverse order

toposort.cpp

Description: Using DFS pos order transversal and inverting the order, one can obtain the topological order

7.7.2 Kosaraju

A Strongly Connected Component is a maximal subgraph in which every vertex is reachable from any vertex inside this same subgraph.

5

A important *property* is that the inverted graph or transposed graph has the same SCCs as the original graph.

kosaraju.cpp

Description: By using the fact that the inverted graph has the same SCCs, just do a DFS twice to find all SCCs. A condensated graph can be created if wished. The condensated graph is a DAG!!

```
Time: \mathcal{O}(V+E)
                                                     381904, 45 lines
struct Kosaraju {
    11 n;
    vector<vll> g, gi, gc;
    vector<bool> vis;
    vector<11> comp;
    stack<11, v11> st;
    void dfs(ll u) { // g
       vis[u] = 1;
        for(auto v : g[u]) if (!vis[v]) dfs(v);
        st.push(u);
    void dfs2(11 u, 11 c) { // gi
        comp[u] = c;
        for (auto v : gi[u]) if (comp[v] == -1) dfs2(v, c);
    Kosaraju (vector<vll> &g )
      : g(g_), n(g_size()-1) { // 1-idx
        gi.assign(n+1, vll());
        for(ll i=1; i<=n; i++) {
            for(auto j : g[i])
                gi[j].pb(i);
        gc.assign(n+1, vll());
        vis.assign(n+1, 0);
        comp.assign(n+1, -1);
        st = stack<11, v11>();
        for(ll i=1; i<=n; i++) if (!vis[i]) dfs(i);
        while(!st.empty()) {
            auto u = st.top(); st.pop();
            if (comp[u] == -1) dfs2(u, u);
        for(11 u=1; u<=n; u++)
            for(auto v : q[u])
                if (comp[u] != comp[v])
```

$\begin{array}{ll} \textbf{7.7.3} & \textbf{2-SAT} & \texttt{gc[comp[u]].pb(comp[v]);} \\ \textbf{SAT}^{\text{l}}(Boolean \ satisfiability \ problem) \ is \ NP-Complete. \end{array}$

2-SAT is a restriction of the SAT problem, in 2-SAT every clause has exactly two variables: $(X_1 \lor X_2) \land (X_2 \lor X_3)$

Every restriction or implication are represented in the graph as directed edges.

The algorithm uses kosaraju to check if any $(X \text{ and } \neg X)$ are in the same Strongly Connected Component (which implies that the problem is impossible).

If it doesn't, there is at least one solution, which can be generated using the topological sort of the same kosaraju (opting for the variables that appers latter in the sorted order)

2sat.cpp

Description: Kosaraju to find if there are SCCs. If there are not cycles, use toposort to choose states

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

87417c, 83 lines

```
// 0-idx graph !!!!
struct TwoSat {
    11 N; // needs to be the twice of the number of variables
    // node with idx 2x \Rightarrow variable x
    // node with idx \ 2x+1 \Rightarrow variable \ !x
    vector<vll> q, qi;
    // g = graph; gi = transposed graph (all edges are inverted)
    TwoSat(ll n) { // number of variables (add +1 faor 1-idx)
       N = 2 * n;
        g.assign(N, vll());
       qi.assiqn(N, vll());
   11 idx; // component idx
    vector<11> comp, order; // topological order (reversed)
    vector<bool> vis, chosen;
    // chosen[x] = 0 \Rightarrow x was assigned
    // chosen[x] = 1 \rightarrow !x was assigned
    // dfs and dfs2 are part of kosaraju algorithm
   void dfs(ll u) {
       vis[u] = 1;
        for (ll v : g[u]) if (!vis[v]) dfs(v);
        order.pb(u);
   void dfs2(ll u, ll c) {
        comp[u] = c;
        for (ll v : qi[u]) if (comp[v] == -1) dfs2(v, c);
   bool solve() {
       vis.assign(N, 0);
        order = vector<11>();
        for (ll i = 0; i < N; i++) if (!vis[i]) dfs(i);
        comp.assign(N, -1); // comp = 0 \ can \ exist
        idx = 1:
        for(ll i=(ll)order.size()-1; i>=0; i--) {
            11 u = order[i];
            if (comp[u] == -1) dfs2(u, idx++);
        chosen.assign(N/2, 0);
        for (11 i = 0; i < N; i += 2) {
            // x and !x in the same component \Rightarrow contradiction
            if (comp[i] == comp[i+1]) return false;
```

```
chosen[i/2] = comp[i] < comp[i+1]; // choose latter</pre>
        return true;
    // a (with flagA) implies \Rightarrow b (with flagB)
    void add(ll a, bool fa, ll b, bool fb) {
        // \{fa == 0\} \Rightarrow a
        // \{fa == 1\} \Rightarrow !a
        a = 2*a + fa;
        b = 2*b + fb;
        g[a].pb(b);
        gi[b].pb(a);
    // force a state for a certain variable (must be true)
    void force(ll a, bool fa) {
        add(a, fa^1, a, fa);
    // xor operation: one must exist, and only one can exist
    void exclusive(ll a, bool fa, ll b, bool fb) {
        add(a, fa^0, b, fb^1);
        add(a, fa^1, b, fb^0);
        add(b, fb^0, a, fa^1);
        add(b, fb^1, a, fa^0);
    // nand operation: no more than one can exist
    void nand(ll a, bool fa, ll b, bool fb) {
        add(a, fa^0, b, fb^1);
        add(b, fb^0, a, fa^1);
};
```

7.8 Trees

lca.cpp

Description: Solves LCA for trees

```
Time: \mathcal{O}(N \log(N)) to build, \mathcal{O}(\log(N)) per query
                                                        7afc1a, 54 lines
struct BinaryLifting {
    11 n, logN = 20; // \sim 1e6
    vector<vll> q;
    vector<ll> depth;
    vector<vll> up;
    BinaryLifting(vector<vll> &g_)
    : g(g_), n(g_size() + 1) { // 1-idx}
        depth.assign(n, 0);
        while ((1 << logN) < n) logN++;
        up.assign(n, vll(logN, 0));
        build();
    }
    void build(ll u = 1, ll p = -1) {
        for(ll i=1; i<logN; i++) {</pre>
             up[u][i] = up[up[u][i-1]][i-1];
        for (auto v : q[u]) if (v != p) {
            up[v][0] = u;
            depth[v] = depth[u] + 1;
            build(v, u);
    ll go(ll u, ll dist) { // O(log(n))
```

```
for(11 i=logN-1; i>=0; i--) { // bigger jumps first
            if (dist & (1LL << i)) {
                u = up[u][i];
        return u;
    ll lca(ll a, ll b) { // O(log(n))
        if (depth[a] < depth[b]) swap(a, b);</pre>
        a = go(a, depth[a] - depth[b]);
        if (a == b) return a;
        for(ll i=logN-1; i>=0; i--) {
            if (up[a][i] != up[b][i]) {
                a = up[a][i];
                b = up[b][i];
        return up[a][0];
    ll lca(ll a, ll b, ll root) { // lca(a, b) when tree is
         rooted at 'root'
        return lca(a, b) ^lca(b, root) ^lca(a, root); //magic
};
```

queryTree.cpp

Description: Binary Lifting for min, max weight present in a simple path **Time:** $\mathcal{O}(N \log(N))$ to build; $\mathcal{O}(\log(N))$ per query 75ba37, 67 lines

```
struct BinaryLifting {
    11 n, logN = 20; // \sim 1e6
    vector<vpll> q;
    vector<11> depth;
    vector<vll> up, mx, mn;
    BinaryLifting(vector<vpll> &g )
    : g(g_), n(g_size() + 1) { // 1-idx}
        depth.assign(n, 0);
        while ((1 << logN) < n) logN++;
        up.assign(n, vll(logN, 0));
        mx.assign(n, vll(logN, -INF));
        mn.assign(n, vll(logN, INF));
        build();
    void build(ll u = 1, ll p = -1) {
        for(ll i=1; i<logN; i++) {</pre>
            mx[u][i] = max(mx[u][i-1], mx[up[u][i-1]][i-1]);
            mn[u][i] = min(mn[u][i-1], mn[up[u][i-1]][i-1]);
            up[u][i] = up[up[u][i-1]][i-1];
        for (auto [v, w] : g[u]) if (v != p) {
            mx[v][0] = mn[v][0] = w;
            up[v][0] = u;
            depth[v] = depth[u] + 1;
            build(v, u);
    array<11, 3> go(11 u, 11 dist) { // O(log(n))
        11 mxval = -INF, mnval = INF;
        for(11 i=logN-1; i>=0; i--) { // bigger jumps first
```

if (dist & (1LL << i)) {

UnB

```
mxval = max(mxval, mx[u][i]);
           mnval = min(mnval, mn[u][i]);
           u = up[u][i];
   return {u, mxval, mnval};
array<11, 3> query(11 u, 11 v) { // O(log(n))
   if (depth[u] < depth[v]) swap(u, v);</pre>
   auto [a, mxval, mnval] = go(u, depth[u] - depth[v]);
   11 b = v;
   if (a == b) return {a, mxval, mnval};
    for(ll i=logN-1; i>=0; i--) {
       if (up[a][i] != up[b][i]) {
           mxval = max({mxval, mx[a][i], mx[b][i]});
           mnval = min({mnval, mn[a][i], mn[b][i]});
           a = up[a][i];
           b = up[b][i];
   mxval = max(\{mxval, mx[a][0], mx[b][0]\});
   mnval = min(\{mnval, mn[a][0], mn[b][0]\});
   return {up[a][0], mxval, mnval};
```

7.9 Math

};

Geometry (8)

Strings (9)

Miscellaneous (10)

Techniques (A)

techniques.txt

Combinatorics

159 lines

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

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

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