			3.7 k-max queue 11 3.8 Union-find 12
C	C++ contest library Miska Kananen August 12, 2019 ontents	4	Mathematics 13 4.1 Number theory 13 4.2 Combinatorics 13 4.3 Matrices 14 4.4 Summations and progressions 15 4.5 Miller-Rabin 15 4.6 Pollard-Rho 16 4.7 Extended Euclidean algorithm 17 4.8 Linear sieve 17
1	Environment and workflow	5 1	Geometry 18
2	1.2 Stress testing 2 General techniques 2 2.1 Bit tricks 2 2.2 Mo's algorithm 3 2.3 Arbitrary precision decimals 3 2.4 Arithmetic overflow checking 3 2.5 g++ pragmas 3 2.6 C++11 std::random 4 2.7 g++ vector extensions 4 2.8 XOR basis vectors 4	1 6 2 2 2 2 3 3 3 3 4 4 4 4 8	6.1 Kosaraju's algorithm 22 6.2 Bridges 22 6.3 Articulation points 23 6.4 Maximum flow (scaling algorithm) 24 6.5 Theorems on flows and cuts 24 6.6 Heavy-light decomposition 25 Tree algorithms 26 7.1 Smaller to larger 26 7.2 Subtree merging DP 27 String algorithms 28
3	3.1 Lazy segment tree	5 5 6 7	8.1 Polynomial hashing 28 8.2 Z-algorithm 29 8.3 Suffix array 29
	3.5 Sparse table 10 3.6 Policy-based data structures 10 3.6.1 Indexed set 10 3.6.2 Hashmap 11	0 0 1	Environment and workflow .1 Compilation script
		1	

```
#!/bin/bash
g++ $1 -o ${1%.*} -std=c++11 -Wall -Wextra -Wshadow -
    ftrapv -Wfloat-equal -Wconversion -Wlogical-op -
    Wshift-overflow=2 -fsanitize=address -fsanitize=
    undefined -fno-sanitize-recover
```

1.2 Stress testing

srand(time(NULL)); changes seed only once a second and is unsuitable for stress testing. RNG seed initialization (requires x86 and g++):

Shell script for stress testing with a brute force solution and a test generator:

```
for i in {1..1000}
do
        echo -n "Test_#$i:_"
        python gen.py > test_input
        ./corr < test_input > corr_output
        # time (seconds), memory (kilobytes)
        (ulimit -t 1 -v 128000; /usr/bin/time -f "%e, %M"
             -o exec_report ./hack < test_input >
            user_output)
        diff corr_output user_output > /dev/null
        res=$?
        if [ $res -ne 0 ]; then
                echo -e -n "\033[1;31mFailed_\033[0m"
                cat exec_report
                echo "Test input:"
                cat test_input
                cp test_input failed_test
```

```
echo ""
        echo "Correct output:"
        cat corr_output
        echo ""
        echo "User output:"
        cat user_output
fi
rm test_input
rm corr_output
rm user_output
if [ $res -ne 0 ]; then
        rm exec_report
        exit 1
echo -e -n "\033[1;32mAccepted_\033[0m"
cat exec_report
rm exec_report
```

2 General techniques

2.1 Bit tricks

done

g++ builtin functions:

- __builtin_clz(x): number of zeros in the beginning
- __builtin_ctz(x): number of zeros in the end
- __builtin_popcount(x): number of set bits
- __builtin_parity(x): parity of number of ones

There are separate functions of form __builtin_clzll(x) for 64-bit integers. For the compiler to utilize the native POPCNT instruction, #pragma GCC target("sse4.2") should be used. Iterate subsets of set s:

2.2 Mo's algorithm

Processes range queries on an array offline in $O(n\sqrt{n}\ f(n))$, where the array has n elements, there are n queries and addition/removal of an element to/from the active set takes O(f(n)) time.

The array is divided into \sqrt{n} blocks of $k = \sqrt{n}$ elements. Queries are sorted such that query $[a_i, b_i]$ goes before $[a_i, b_i]$ if:

1.
$$\lfloor \frac{a_i}{k} \rfloor < \lfloor \frac{a_j}{k} \rfloor$$
 or

2.
$$\lfloor \frac{a_i}{k} \rfloor = \lfloor \frac{a_j}{k} \rfloor$$
 and $b_i < b_j$

Active range is maintained between queries and the endpoints of the range are moved accordingly. Both endpoints move $O(n\sqrt{n})$ steps in total during the algorithm.

2.3 Arbitrary precision decimals

Python 3 implements arbitrary precision decimal arithmetic in module decimal. All decimal numbers are represented exactly and the precision is user-definable.

```
from decimal import *
a, b = [Decimal(x) for x in input().split("_")]
getcontext().prec = 50 # set precision
print(a/b)
```

2.4 Arithmetic overflow checking

g++ implements efficient builtin functions for checking for arithmetic overflow. Functions are of form bool __builtin_overflow(a, b, *res) and return true if operation overflows. The result of the operation is returned through res.

```
• __builtin_sadd_overflow(),
builtin saddll overflow: addition
```

```
• __builtin_ssub_overflow(),
__builtin_ssubll_overflow: subtraction
```

```
• __builtin_smul_overflow(),
__builtin_smulll_overflow: multiplication
```

There are separate functions for 32- and 64-bit integers. Unsigned versions are of form __builtin_uadd_overflow().

2.5 g++ pragmas

Pragmas optimize all functions defined afterwards. They should be located in the very beginning of the source code, even before includes in order to optimize imported standard library code.

```
#pragma GCC optimize("03")
#pragma GCC optimize("Ofast"), enables more opti-
mizations but isn't always faster.
```

```
#pragma GCC optimize("unroll-loops")
#pragma GCC target("arch=skylake")
#pragma GCC target("mmx,sse,sse2,sse3,
ssse3,sse4.2,popcnt,avx,tune=native") for ivybridge
if arch=ivybridge fails.
```

All possible target architectures are listed in compiler report if an invalid architecture is given to arch. Supported Intel Core generations in order: nehalem, sandybridge, ivybridge (for CF), haswell (first avx2), broadwell, skylake.

2.6 C++11 std::random

If different ranges are required on every iteration, just create a new distribution every time, it's quite fast.

2.7 g++ vector extensions

Requires AVX support from the grading CPU. If heap-allocating, memory must be aligned to a multiple of 32. Stack allocation works normally.

```
// elementwise minimum
inline float8_t min8(float8_t x, float8_t y) {
    return x < y ? x : y;
}</pre>
```

2.8 XOR basis vectors

XOR is equivalent to bitwise addition modulo 2. We can consider the bitwise representation of an integer a as a d-dimensional vector modulo 2, where d is the maximum needed bit count.

We can find the basis of a vector space of n d-element vectors in \mathbb{Z}_2^d in O(nd). We check for each vector whether it can be represented with current basis vectors, and if not, we add it to the basis.

Let f(v) be the position of the lowest 1-bit in the binary representation of v. We want all basis vectors to have a distinct value of f. Let b_1 be the vector in the current basis with the smallest f. If $f(v) < f(b_1)$ for a new vector, we just insert it, since no linear combination of current basis vectors has 1 at f(v). If $f(v) = f(b_1)$, we subtract b_1 from v and continue to b_2 . If in the end v is not a null vector, it can't be represented using current basis vectors and we have to add it as a new basis vector.

3 Data structures

3.1 Lazy segment tree

Implements range add and range sum query in $O(\log(n))$. 0-indexed

```
operator+logic
                if (1 != r) {
                        lz[2*s] += lz[s]; // change
                             operator
                        lz[2*s+1] += lz[s]; // change
                             operator
                        haslz[2*s] = true;
                        haslz[2*s+1] = true;
                lz[s] = 0; // set to identity
                haslz[s] = false;
ll kysy(int ql, int qr, int s = 1, int l = 0, int r = N
    -1) {
        push(s, l, r);
        if (l > qr || r < ql) {</pre>
                return 0; // set to identity
        if (ql <= l && r <= qr) {
                return st[s];
        int mid = (1+r)/2;
        11 res = 0; // set to identity
        res += kysy(ql, qr, 2*s, 1, mid); // change
            operator
        res += kysy(gl, gr, 2*s+1, mid+1, r); // change
            operator
        return res;
void muuta(int ql, int qr, ll x, int s = 1, int l = 0,
    int r = N-1) {
        push(s, l, r);
        if (1 > qr || r < ql) {</pre>
                return;
        if (ql <= l && r <= qr) {
                lz[s] += x; // change operator
```

st[s] += (r-l+1)*lz[s]; // change

if (haslz[s]) {

```
haslz[s] = true;
                return;
        int mid = (1+r)/2;
        muuta(ql, qr, x, 2*s, l, mid);
        muuta(ql, qr, x, 2*s+1, mid+1, r);
        st[s] = st[2*s] + st[2*s+1]; // change operator
        if (haslz[2*s]) {
                st[s] += (mid-l+1)*lz[2*s]; // change
                    operator+logic
        if (haslz[2*s+1]) {
                st[s] += (r-(mid+1)+1)*lz[2*s+1]; //
                    change operator+logic
void build(int s = 1, int l = 0, int r = N-1) {
        if (r-1 > 1) {
                int mid = (1+r)/2;
                build(2*s, 1, mid);
               build(2*s+1, mid+1, r);
        st[s] = st[2*s]+st[2*s+1]; // change operator
/*
        TESTED, correct
        Allowed indices 0..N-1
        2 types of queries: range add and range sum
int main() {
        for (int i = 1; i <= n; ++i) {</pre>
               cin >> st[i+N];
       build();
```

3.2 Sparse segment tree

Implements point update and range sum query in O(log(n)). 0-indexed.

```
#include <iostream>
using namespace std;
typedef long long 11;
const int N = 1 << 30; // max element index
struct node {
        11 s;
        node *1, *r;
        node (int cs) : s(cs) {
               1 = nullptr;
                r = nullptr;
};
node *st = new node(0); // segtree root node
void update(int k, ll val, int nl = 0, int nr = N-1,
    node *nd = st) {
        if (nl == nr) {
                nd->s += val; // change operator
        else {
                int mid = (nl + nr)/2;
                if (nl <= k && k <= mid) {
                        if (nd->1 == nullptr) nd->1 =
                            new node(0);
                        update(k, val, nl, mid, nd->1);
                else if (mid < k && k <= nr) {
                        if (nd->r == nullptr) nd->r =
                            new node(0);
                        update(k, val, mid+1, nr, nd->r)
                11 ns = 0; // set to identity
                if (nd->1 != nullptr) ns += (nd->1)->s;
                    // change operator
```

```
if (nd->r != nullptr) ns += (nd->r)->s;
                    // change operator
                nd->s = ns;
ll query(int ql, int qr, int nl = 0, int nr = N-1, node
    *nd = st) {
        if (ql <= nl && nr <= qr) return nd->s;
        if (nr < ql || nl > qr) return 0; // set to
            identity
        int mid = (nl + nr)/2;
        11 res = 0; // set to identity
        if (nd->1 != nullptr) res += query(q1, qr, n1,
            mid, nd->1); // change operator
        if (nd->r != nullptr) res += query(ql, qr, mid
            +1, nr, nd->r); // change operator
        return res;
}
```

3.3 2D segment tree

Implements point update and subgrid query in $O(log^2(n))$. Grid is 0-indexed

```
#include <iostream>
using namespace std;
typedef long long 11;

const int N = 1<<11;
int n, q;

11 st[2*N][2*N];

// calculate subgrid sum from {y1, x1} to {y2, x2}

// 0-indexed

11 summa(int y1, int x1, int y2, int x2) {
    y1 += N;
    x1 += N;
    y2 += N;
    x2 += N;</pre>
```

```
while (y1 <= y2) {
        if (y1%2 == 1) {
            int nx1 = x1;
            int nx2 = x2;
            while (nx1 \le nx2) {
                if (nx1\%2 == 1) sum += st[y1][nx1++];
                if (nx2\%2 == 0) sum += st[y1][nx2--];
                nx1 /= 2;
                nx2 /= 2;
            y1++;
        if (y2\%2 == 0) {
            int nx1 = x1;
            int nx2 = x2;
            while (nx1 \le nx2) {
                if (nx1\%2 == 1) sum += st[y2][nx1++];
                if (nx2\%2 == 0) sum += st[y2][nx2--];
                nx1 /= 2;
                nx2 /= 2;
            y2--;
        y1 /= 2;
        y2 /= 2;
    return sum;
// set {y, x} to u
// 0-indexed
void muuta(int y, int x, ll u) {
    y += N;
    x += N;
    st[y][x] = u;
    for (int nx = x/2; nx >= 1; nx /= 2) {
        st[y][nx] = st[y][2*nx]+st[y][2*nx+1];
    for (y /= 2; y >= 1; y /= 2) {
        for (int nx = x; nx >= 1; nx /= 2) {
            st[y][nx] = st[2*y][nx]+st[2*y+1][nx];
```

11 sum = 0;

}

3.4 Treap

Implements split, merge, kth element, range update and range reverse in O(log(n)). Range update adds a value to every element woid push (node *s) { if (s == nul

Note: Memory management tools warn of about 30 MB memory leak for $500\ 000$ elements. This is because nodes are not deleted when exiting program and is irrelevant in a competition. Deleting nodes would slow the treap down by a factor of 3.

```
#include <iostream>
#include <cstdlib>
#include <algorithm>
using namespace std;
typedef long long 11;
struct node {
        11 val; // change data type (char, integer...)
        int prio, size;
        bool lzinv;
        ll lzupd;
        bool haslz;
        node *left, *right;
        node(ll v) {
                val = v;
                prio = rand();
                size = 1;
                lzinv = false;
                lzupd = 0;
                haslz = false;
                left = nullptr;
                right = nullptr;
};
int gsize(node *s) {
        if (s == nullptr) return 0;
```

```
return s->size;
void upd(node *s) {
        if (s == nullptr) return;
        s->size = gsize(s->left) + 1 + gsize(s->right);
        if (s == nullptr) return;
        if (s->haslz) {
                 s->val += s->lzupd; // operator
        if (s->lzinv) {
                 swap(s->left, s->right);
        if (s->left != nullptr) {
                 if (s->haslz) {
                          s->left->lzupd += s->lzupd; //
                              operator
                          s->left->haslz = true;
                 if (s->lzinv) {
                          s \rightarrow left \rightarrow lzinv = !s \rightarrow left \rightarrow lzinv
        if (s->right != nullptr) {
                 if (s->haslz) {
                          s->right->lzupd += s->lzupd; //
                              operator
                          s->right->haslz = true;
                 if (s->lzinv) {
                          s->right->lzinv = !s->right->
                              lzinv:
        s->lzupd = 0; // operator identity value
        s->lzinv = false;
        s->haslz = false;
```

```
if (k < cval) return kthElem(t->left, k);
// split a treap into two treaps, size of left treap = k
                                                                   return kthElem(t->right, k-cval);
void split(node *t, node *&l, node *&r, int k) {
        push(t);
        if (t == nullptr) {
                                                           // do a lazy update on subarray [a..b]
                                                           void rangeUpd(node *&t, int a, int b, ll x) {
               l = nullptr;
                r = nullptr;
                                                                   node *cl, *cur, *cr;
                return;
                                                                   int tsz = gsize(t);
                                                                   bool lsplit = false;
        if (k \ge gsize(t->left)+1) {
                                                                   bool rsplit = false;
                split(t->right, t->right, r, k-(gsize(t
                                                                   cur = t;
                                                                   if (a > 1) {
                    ->left)+1));
               1 = t;
                                                                           split(cur, cl, cur, a-1);
                                                                           lsplit = true;
        else {
                split(t->left, l, t->left, k);
                                                                   if (b < tsz) {
                                                                           split(cur, cur, cr, b-a+1);
               r = t;
                                                                           rsplit = true;
        upd(t);
                                                                   cur->lzupd += x; // operator
                                                                   cur->haslz = true;
// merge two treaps
                                                                   if (lsplit) {
void merge(node *&t, node *1, node *r) {
                                                                           merge(cur, cl, cur);
        push(1);
                                                                   if (rsplit) {
        push(r);
        if (l == nullptr) t = r;
                                                                           merge(cur, cur, cr);
        else if (r == nullptr) t = 1;
                                                                   t = cur;
                if (l->prio >= r->prio) {
                        merge(l->right, l->right, r);
                                                           // reverse subarray [a..b]
                                                           void rangeInv(node *&t, int a, int b) {
                else {
                                                                   node *cl, *cur, *cr;
                                                                   int tsz = gsize(t);
                        merge(r->left, l, r->left);
                        t = r;
                                                                   bool lsplit = false;
                                                                   bool rsplit = false;
                                                                   cur = t;
        upd(t);
                                                                   if (a > 1) {
                                                                           split(cur, cl, cur, a-1);
                                                                           lsplit = true;
// get k:th element in array (1-indexed)
11 kthElem(node *t, int k) {
                                                                   if (b < tsz) {
        push(t);
                                                                           split (cur, cur, cr, b-a+1);
        int cval = qsize(t->left)+1;
                                                                           rsplit = true;
        if (k == cval) return t->val;
```

3.5 Sparse table

Implements range minimum/maximum query in O(1) with $O(n \ log(n))$ preprocessing. 0-indexed.

```
#include <iostream>
#include <cmath>

using namespace std;
typedef long long ll;

int n, q;
ll t[100005];
ll st[18][100005];

ll rmq(int a, int b) {
    int l = b-a+1;
    int k = (int)log2(l);
    return min(t[st[k][a]], t[st[k][a+(l-(1<<k))]]);
    // change function
}</pre>
```

3.6 Policy-based data structures

3.6.1 Indexed set

Works like std::set but adds support for indices. Set is 0-indexed. Requires g++. Has two additional functions:

- 1. find_by_order(x): return an iterator to element at index x
- 2. order_of_key(x): return the index that element x has or would have in the set, depending on if it exists

Both functions work in O(log(n)).

Changing less to less_equal makes the set work like multiset. In this case:

- find() will always return end
- lower_bound() works like upper_bound() (returns an iterator to first element > x)

- upper_bound() works like lower_bound() (returns an iterator to first element $\geq x$) using namespace std; using namespace __gn
- erase() works as expected with iterators obtained from lower_bound() and upper_bound(), deleting a single element
- order_of_key() returns the index of the first element, } if there are multiple ones
- find_by_order() works as expected

```
#include <iostream>
#include <ext/pb_ds/assoc_container.hpp>
using namespace std;
using namespace __gnu_pbds;
typedef tree<int, null_type, less<int>, rb_tree_tag,
    tree_order_statistics_node_update> indexed_set;
indexed_set s;
int main() {
        s.insert(2);
        s.insert(4);
        s.insert(5);
        auto x = s.find_by_order(1);
        cout << *x << "\n"; // prints 4
        cout << s.order_of_key(5) << "\n"; // prints 2</pre>
        cout << s.order_of_key(3) << "\n"; // prints 1</pre>
        return 0;
```

3.6.2 Hashmap

Works like std::unordered_map but is many times faster.

```
#include <iostream>
#include <ext/pb_ds/assoc_container.hpp>
```

```
using namespace __gnu_pbds;
// get a random number
uint32 t rd() {
        uint32_t ret;
        asm volatile("rdrand, %0" : "=a"(ret) :: "cc");
        return ret;
const uint32 t XR = rd();
// xor with a random number to avoid anti-hash tests
struct chash {
    int operator()(int x) const { return hash<int>{}(x^
        XR); }
};
int n;
gp_hash_table<11, int, chash> s;
int main() {
        ios_base::sync_with_stdio(false);
        cin.tie(0);
        cin >> n:
        for (int i = 0; i < n; ++i) {
                int x;
                cin >> x;
                s[x] = 1;
        cout << s.size() << "\n";
        return 0;
```

3.7 k-max queue

Works like std::queue, but implements O(1) max query for elements in queue. All operations are O(1), push_back(x) is amortized O(1). Can be used as a min queue if elements are inserted as negative.

lt's not possible to return popped element on pop_front().
#include <deque>

```
template <typename T>
struct kmax_queue {
private:
        std::deque<std::pair<T, int>> q;
        int q_size;
public:
        kmax_queue() {
                q_size = 0;
        void push_back(T x) {
                int unimp_before = 0;
                while ((!q.empty()) && (q.back().first
                    <= x)) {
                        unimp_before += q.back().second
                            + 1;
                        g.pop_back();
                q.push_back({x, unimp_before});
                q_size++;
        void pop_front() {
                if (empty()) {
                        throw ("The queue is empty");
                if (q.front().second > 0) {
                        q.front().second--;
                else {
                        q.pop_front();
                q_size--;
        T max() {
                if (empty()) {
                        throw ("The_queue_is_empty");
```

```
return q.front().first;
}
int size() {
    return q_size;
}
bool empty() {
    return size() == 0;
}
};
```

3.8 Union-find

Uses path compression, id(x) has amortized time complexity $O(a^{-1}(n))$ where a^{-1} is inverse Ackermann function.

```
#include <iostream>
#include <algorithm>
using namespace std;
int k[100005];
int s[100005];
int id(int x) {
        int tx = x;
        while (k[x] != x) x = k[x];
        return k[tx] = x;
bool equal(int a, int b) {
        return id(a) == id(b);
void join(int a, int b) {
        a = id(a);
        b = id(b);
        if (s[b] > s[a]) swap(a, b);
        s[a] += s[b];
        k[b] = a;
```

```
int n;
int main() {
    for (int i = 0; i < n; ++i) {
        k[i] = i;
        s[i] = 1;
}</pre>
```

4 Mathematics

4.1 Number theory

- Prime factorization of n: $p_1^{\alpha_1} p_2^{\alpha_2} \dots p_k^{\alpha_k}$
- Number of factors: $\tau(n) = \prod_{i=1}^k (\alpha_i + 1) \approx \sqrt[3]{n}$

-
$$max(\tau(1), \tau(2), \dots \tau(10^9)) = 1344$$

$$- \max(\tau(1), \tau(2), \dots, \tau(10^{18})) = 103680$$

- Sum of factors: $\sigma(n) = \prod_{i=1}^k \frac{p_i^{\alpha_i+1}-1}{p_i-1}$
- Product of factors: $\mu(n) = n^{\tau(n)/2}$

Euler's totient (phi) function $\varphi(n)$ $(1,1,2,2,4,2,6,4,6,4,\dots)$: counts numbers coprime with n in range $1\dots n$

$$\varphi(n) = \begin{cases} n-1 & \text{if } n \text{ is prime} \\ \prod_{i=1}^k p_i^{a_i-1}(p_i-1) & \text{otherwise} \end{cases}$$

The function can be precomputed for all natural numbers $\leq n$ in $O(n \ log(n))$ with a sieve:

```
for (int j = 2*i; j <= N; j += i) {
          phi[j] -= phi[i];
}</pre>
```

There are $\varphi(\frac{n}{d})$ numbers i $(1 \le i \le n)$ for which $\gcd(i,n) = d$ if $d \mid n$. If $d \nmid n$, there are none.

Fermat's theorem: $x^{m-1} \mod m = 1$ when m is prime and x and m are coprime. It follows that $x^k \mod m = x^{k \mod (m-1)} \mod m$.

Modular inverse $x^{-1} = x^{\varphi(m)-1}$. If m is prime, $x^{-1} = x^{m-2}$. Inverse exists if and only if x and m are coprime.

4.2 Combinatorics

Binomial coefficients:

$$\binom{n}{k} = \binom{n-1}{k-1} + \binom{n-1}{k}$$
$$\binom{n}{0} = \binom{n}{n} = 1$$

Catalan numbers (1, 1, 2, 5, 14, 42, 132, 429, 1430, 4862, 16796...):

$$C_n = \frac{1}{n+1} \binom{2n}{n}$$

Classic examples of Catalan numbers: number of balanced pairs of parentheses, number of mountain ranges (n upstrokes and n downstrokes all staying above the original line), number of paths from upper left corner to lower right corner staying above the main diagonal in a $n \times n$ square, ways to triangulate a n+2 sided regular polygon, ways to shake hands between 2n people in a circle such that no arms cross, number of rooted binary trees with n nodes that have 2 children, number of rooted trees with n edges, number of permutations of $1 \dots n$ that don't have an increasing subsequence of length 3.

Number of derangements (no element stays in original place) of $1, 2, \ldots, n$ $(1, 0, 1, 2, 9, 44, 265, 1854, 14833, 133496, 1334961, \ldots)$:

$$f(n) = \begin{cases} 0 & n = 1\\ 1 & n = 2\\ (n-1)(f(n-2) + f(n-1)) & n > 2 \end{cases}$$

Stirling numbers of the second kind $\binom{n}{k}$: number of ways to partition a set of n objects into k non-empty subsets.

$$\begin{array}{c}
1 \\
0, 1 \\
0, 1, 1 \\
0, 1, 3, 1 \\
0, 1, 7, 6, 1 \\
0, 1, 15, 25, 10, 1 \\
0, 1, 31, 90, 65, 15, 1
\end{array}$$

$$\begin{cases}
n+1 \\ k
\end{cases} = k \begin{Bmatrix} n \\ k \end{Bmatrix} + \begin{Bmatrix} n \\ k-1 \end{Bmatrix} \quad (k>0)$$

$$\begin{Bmatrix} 0 \\ 0 \end{Bmatrix} = 1, \begin{Bmatrix} n \\ 0 \end{Bmatrix} = \begin{Bmatrix} 0 \\ n \end{Bmatrix} = 0 \quad (n>0)$$

4.3 Matrices

Matrix $A = a \times n$, matrix $B = n \times b$. Matrix multiplication:

$$AB[i,j] = \sum_{k=1}^{n} A[i,k] \cdot B[k,j]$$

Let linear recurrence $f(n)=c_1f(n-1)+c_2f(n-2)+\cdots+c_kf(n-k)$ with initial values $f(0),f(1),\ldots,f(k-1).$ c_1,c_2,\ldots,c_n are constants.

Transition matrix X:

$$X = \begin{pmatrix} 0 & 1 & 0 & \dots & 0 \\ 0 & 0 & 1 & \dots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & 0 & \dots & 1 \\ c_k & c_{k-1} & c_{k-2} & \dots & c_1 \end{pmatrix}$$

Now f(n) can be calculated in $O(k^3 log(n))$:

$$\begin{pmatrix} f(n) \\ f(n+1) \\ \vdots \\ f(n+k-1) \end{pmatrix} = X^n \cdot \begin{pmatrix} f(0) \\ f(1) \\ \vdots \\ f(k-1) \end{pmatrix}$$

```
#include <iostream>
#include <cstring>
using namespace std;
typedef long long 11;
const int N = 2; // matrix size
const 11 M = 1000000007; // modulo
struct matrix {
    11 m[N][N];
    matrix() {
        memset(m, 0, sizeof m);
   matrix operator * (matrix b) {
        matrix c = matrix();
        for (int i = 0; i < N; ++i)
            for (int j = 0; j < N; ++j)
                for (int k = 0; k < N; ++k) {
                    c.m[i][j] = (c.m[i][j] + m[i][k] * b
                         .m[k][j])%M;
        return c;
   matrix unit() {
        matrix a = matrix();
```

```
for (int i = 0; i < N; ++i) a.m[i][i] = 1;</pre>
        return a;
};
matrix p(matrix a, ll e) {
    if (e == 0) return a.unit();
    if (e%2 == 0) {
        matrix h = p(a, e/2);
        return h*h;
    return (p(a, e-1)*a);
}
11 n;
// prints nth Fibonacci number mod M
int main() {
    cin >> n;
    matrix x = matrix();
    x.m[0][1] = 1;
    x.m[1][0] = 1;
    x.m[1][1] = 1;
    x = p(x, n);
    cout << x.m[0][1] << "\n";
    return 0;
```

4.4 Summations and progressions

- Sum of naturals: $\sum_{i=1}^{n} x = \frac{n(n+1)}{2}$
- Sum of squares: $\sum_{i=1}^{n} x^2 = \frac{n(n+1)(n+2)}{6}$
- Arithmetic progression: $a + \cdots + b = \frac{n(a+b)}{2}$, where n is the number of terms, a is the first term and b is the last term
- Geometric progression: $a+ar+ar^2+\cdots+ar^{n-1}=a\frac{1-r^n}{1-r}$, where n is the number of terms, a is the first term and $r(r\neq 1)$ is the ratio between two successive terms

```
- If r=1. sum is na
```

- Also $a + ar + ar^2 + \cdots + b = \frac{a - br}{1 - r}$, where a is the first term, b is the last term and r is the ratio between two successive terms

Terms of sum $S=\sum_{i=1}^n\lfloor\frac{n}{i}\rfloor$ get at most $O(\sqrt{n})$ distinct values. All terms and their counts can be found as follows in $O(\sqrt{n})$:

```
#include <iostream>
#include <vector>
using namespace std;
typedef long long 11;
11 n;
int main() {
        cin >> n;
        vector<ll> v;
        11 x = 0;
        for (ll i = 1; i <= n; i = x+1) {</pre>
                x = n/(n/i); // iterate all possible
                     values of floor(n/i) in increasing
                     order
                v.push_back(x);
        for (int i = 0; i < v.size(); ++i) {</pre>
                // current value of floor(n/i)
                ll cx = v[i];
                // smallest i for which floor(n/i) == cx
                ll imin = (i == v.size()-1 ? 1 : n/v[i
                     +1] + 1);
                // largest i for which floor(n/i) == cx
                11 imax = n/cx;
        return 0;
```

4.5 Miller-Rabin

Deterministic primality test for all 64-bit integers. Requires __int128 support to test over 32-bit integers.

```
#include <iostream>
using namespace std;
typedef long long 11;
typedef __int128 111;
// required bases to make test deterministic for 64-bit
    integers
11 \text{ mrb}[12] = \{2, 3, 5, 7, 11, 13, 17, 19, 23, 29, 31,
    371:
111 modpow(111 k, 111 e, 111 m) {
        if (e == 0) return 1;
        if (e == 1) return k;
        if (e%2 == 0) {
                lll h = modpow(k, e/2, m)%m;
                return (h*h)%m;
        return (k*modpow(k, e-1, m))%m;
bool witness(ll a, ll x, ll u, ll t) {
        lll cx = modpow(a, u, x);
        for (int i = 1; i <= t; ++i) {</pre>
                lll nx = (cx*cx)%x;
                if (nx == 1 && cx != 1 && cx != (x-1))
                     return true;
                cx = nx;
        return (cx != 1);
// TESTED, correct
// determines if x is prime
// deterministic for all 64-bit integers
bool miller_rabin(ll x) {
        if (x == 2) return true;
        if (x < 2 \mid | x \% 2 == 0) return false;
        11 u = x-1;
        11 t = 0;
        while (u%2 == 0) {
                u /= 2;
                t++;
```

```
for (int i = 0; i < 12; ++i) {
    if (mrb[i] >= x-1) break;
    if (witness(mrb[i], x, u, t)) return
        false;
}
return true;
```

4.6 Pollard-Rho

Finds a factor of x in $O(\sqrt[4]{x})$. Requires __int128 support to factor over 32-bit integers.

If x is prime or a perfect square, algorithm might not terminate or it might return 1. Primality must be checked separately.

```
#include <iostream>
#include <cstdlib>
#include <algorithm>
using namespace std;
typedef long long 11;
typedef __int128 111;
11 n:
ll f(lll x) {
    return (x*x+1)%n;
11 gcd(ll a, ll b) {
    if (b == 0) return a;
    return gcd(b, a%b);
// return a factor of a
// st is a starting seed for pseudorandom numbers, start
     with 2, if algorithm fails (returns -1), increment
ll pollardrho(ll a, ll st) {
    if (n%2 == 0) return 2;
```

```
11 x = st, y = st, d = 1;
    while (d == 1) {
        x = f(x);
        y = f(f(y));
        d = gcd(abs(x-y), a);
        if (d == a) return -1;
    return d;
ll is_square(ll x) {
        11 a = 1;
        for (11 b = (1LL<<30); b >= 1; b /= 2) {
                if ((a+b)*(a+b) \le x) a += b;
        if (a*a == x) return a;
        return -1;
}
/*
        TESTED, correct.
    Finds a factor of n in O(root_4(n))
    If n is prime, alg might not terminate or it might
        return 1. Check for primality.
int main() {
    cin >> n;
    // check if n is square, pollardrho might fail if
        the input is perfect square
    11 sq = is_square(n);
    if (sq !=-1) {
        cout << sq << "_" << sq << "\n";
        return 0;
    11 \text{ fa} = -1;
    11 \text{ st} = 2:
    while (fa == -1) {
        fa = pollardrho(n, st++);
    cout << min(fa, n/fa) << "." << max(fa, n/fa) << "\n</pre>
    return 0;
```

4.7 Extended Euclidean algorithm

4.8 Linear sieve

ios_base::sync_with_stdio(false);

```
cin.tie(0);
                                                             // point equality comparison
        cin >> n;
                                                             bool equal(point a, point b) {
        for (int i = 2; i <= n; ++i) {</pre>
                                                                     return (equal(a.X, b.X) && equal(a.Y, b.Y));
                if (mpf[i] == 0) {
                        mpf[i] = i;
                        pr.push_back(i);
                                                             // comparer for sorting points
                                                             // check if a < b
                for (int j = 0; j < pr.size(); ++j) {</pre>
                                                             bool point_comp(point a, point b) {
                        if (mpf[i] < pr[j]) break;</pre>
                                                                     if (equal(a.X, b.X)) {
                        int a = pr[j]*i;
                                                                             return a.Y < b.Y;</pre>
                        if (a > n) break;
                                                                     return a.X < b.X;</pre>
                        mpf[a] = pr[j];
                                                             struct line {
        for (int a : pr) cout << a << "..";</pre>
                                                                     point first, second;
        cout << "\n";
        return 0;
                                                                     line(point a, point b) {
                                                                             if (point_comp(b, a)) swap(a, b);
                                                                              first = a;
     Geometry
                                                                              second = b;
                                                                     // construct line from point and angle of
#include <iostream>
                                                                          elevation
#include <complex>
                                                                     line(point a, ct ang) : line(a, a+polar((ct)1.0,
#include <vector>
                                                                           ang)) {}
#include <algorithm>
#include <iomanip>
                                                                     // construct line from standard equation
                                                                          coefficients
using namespace std;
                                                                     // assume that a != 0 or b != 0
typedef long double ct; // coordinate type
                                                                     // TESTED
typedef complex<ct> point;
                                                                     line(ct a, ct b, ct c) {
#define X real()
                                                                             if (equal(b, 0.0)) {
#define Y imag()
                                                                                      // vertical line
#define F first
#define S second
                                                                                      ct cx = c/(-a);
                                                                                      first = \{cx, 0\};
                                                                                      second = \{cx, 1\};
const ct EPS = 0.000001; // 1e-6
const ct PI = 3.14159265359;
                                                                              else {
                                                                                      first = \{0, c/(-b)\};
// floating-point equality comparison
```

bool equal(ct a, ct b) {

return abs(a-b) < EPS;</pre>

second = $\{1, (a+c)/(-b)\};$

if (point_comp(second, first)) swap(

```
first, second);
                                                                   return (conj(a) *b) .Y;
};
                                                           // euclidean distance
struct line_segment {
                                                           // TESTED
        point first, second;
                                                           ct dist(point a, point b) {
                                                                    return abs(a-b);
        // implicit conversion
        operator line() {
                return line(first, second);
                                                           // squared distance
                                                            ct sq_dist(point a, point b) {
                                                                   return norm(a-b);
        line_segment(point a, point b) {
                if (point_comp(b, a)) swap(a, b);
                first = a;
                                                           // angle from a to b
                second = b;
                                                           // [0, 2*pi[
                                                           // TESTED
                                                           ct angle (point a, point b) {
        line_segment(point a, ct ang, ct len) :
                                                                   ct cres = arg(b-a);
            line_segment(a, a+polar(len, ang)) {};
                                                                   if (cres < 0) cres = 2*PI+cres;</pre>
};
                                                                    return cres;
// assume that the first and last vertices are the same
typedef vector<point> polygon;
                                                           // angle of elevation
                                                            // [-pi/2, pi/2]
// radians to degrees
                                                           ct elev_ang(point a, point b) {
ct rad_to_deg(ct arad) {
                                                                   if (point_comp(b, a)) swap(a, b);
        return (arad*((ct)180.0/PI));
                                                                   return arg(b-a);
// degrees to radians
                                                            // angle of elevation
ct deg_to_rad(ct adeg) {
                                                           ct elev_ang(line l) {
        return (adeg*(PI/(ct)180.0));
                                                                   return elev_ang(1.F, 1.S);
// dot product, > 0 if a, b point to same direction, 0
                                                            // slope of line
    if perpendicular, < 0 if pointing to opposite
                                                            ct slope(point a, point b) {
    directions
                                                                   return tan(elev_ang(a, b));
ct dot(point a, point b) {
       return (conj(a) *b) .X;
                                                           // slope of line
                                                            ct slope(line 1) {
// 2D cross product, > 0 if a+b turns left, 0 if
                                                                    return tan(elev_ang(1));
    collinear, < 0 if turns right
ct cross(point a, point b) {
```

```
// length of line segment
                                                                       S));
ct segment_len(line_segment ls) {
        return dist(ls.F, ls.S);
                                                           // get projection of a on 1
                                                           // TESTED
// rotate a around origin by ang
                                                           point point_line_proj(point a, line l) {
point rot_origin(point a, ct ang) {
                                                                   return (1.F+(1.S-1.F) *dot(a-1.F, 1.S-1.F) /norm(1
        return (a*polar((ct)1.0, ang));
                                                                       .S-1.F));
// rotate a around ps by ang
                                                           // reflect a across l
point rot_pivot(point a, point ps, ct ang) {
                                                           point point_line_refl(point a, line l) {
        return ((a-ps)*polar((ct)1.0, ang)+ps);
                                                                   return (1.F+conj((a-1.F)/(1.S-1.F)) * (1.S-1.F));
// translate a by dist to the direction of ang
                                                           // angle a-b-c
point translate(point a, ct dist, ct ang) {
                                                           // [O, PI]
        return a+polar(dist, ang);
                                                           // TESTED
                                                           ct ang_abc(point a, point b, point c) {
                                                                   return abs(remainder(arg(a-b)-arg(c-b), (ct)2.0*
// check if a -> b -> c turns counterclockwise
                                                                       PI));
bool ccw(point a, point b, point c) {
        return cross({b.X-a.X, b.Y-a.Y}, {c.X-a.X, c.Y-a
            .Y}) > 0;
                                                           // shortest distance between point a and line 1
                                                           ct point_line_dist(point a, line l) {
// < 0 if point is left, ~0 if on line, > 0 if right
                                                                   point proj = point_line_proj(a, 1);
                                                                   return dist(a, proj);
ct point_line_side(point a, line l) {
        return cross(a-1.F, a-1.S);
                                                           // shortest distance between point a and line segment ls
                                                           // TESTED
// check if point is on line
                                                           ct point_segment_dist(point a, line_segment ls) {
// TESTED
                                                                   point proj = point_line_proj(a, ls);
bool point on line(point a, line 1) {
                                                                   if (point_on_seg(proj, ls)) {
                                                                           return dist(a, proj);
        return equal(point_line_side(a, 1), (ct)0.0);
                                                                   return min(dist(a, ls.F), dist(a, ls.S));
// check if point is on line segment
// TESTED
bool point_on_seg(point a, line_segment ls) {
                                                           // get intersection point of two lines
                                                           // first return val 0 = no intersection, 1 = single
        if (!point on line(a, ls)) return false;
        if (equal(a, ls.F) || equal(a, ls.S)) return
                                                               point, 2 = infinitely many
                                                           // second return val = intersection point if first
        return (point_comp(ls.F, a) && point_comp(a, ls.
                                                               return val = 1, otherwise undefined
```

```
// TESTED (only non-degenerate cases, single
                                                                   auto tres = intersect(a, b);
    intersection point)
                                                                   if (tres.F == 0) {
pair<int, point> intersect(line a, line b) {
                                                                            return tres;
        ct c1 = cross(b.F-a.F, a.S-a.F);
        ct c2 = cross(b.S-a.F, a.S-a.F);
                                                                   else if (tres.F == 2) {
                                                                           vector<pair<point, int>> v = {{a.F, 1},
        if (equal(c1, c2)) {
                if (point_on_line(b.F, a)) {
                                                                                {a.S, 1}, {b.F, 2}, {b.S, 2}};
                        return {2, a.F};
                                                                            sort(v.begin(), v.end(), pi_comp);
                                                                           if (v[0].S != v[1].S) return {2, a.F};
                                                                                // overlapping segments
                return {0, a.F};
        return {1, (c1*b.S-c2*b.F)/(c1-c2)};
                                                                            // common vertex
}
                                                                           if (equal(a.S, b.F)) return {1, a.S};
                                                                            if (equal(a.F, b.S)) return {1, a.F};
// sort comparer for seg_intersect
bool pi_comp(pair<point, int> p1, pair<point, int> p2) {
                                                                           // not intersecting but on the same line
        if (equal(p1.F, p2.F)) return p1.S < p2.S;</pre>
                                                                            return {0, a.F};
        return point_comp(p1.F, p2.F);
}
                                                                   if (point_on_seg(tres.S, a) && point_on_seg(tres
                                                                        .S, b)) {
// get intersection point of two line segments
                                                                           return tres;
// first return val 0 = no intersection, 1 = single
    point, 2 = infinitely many
                                                                   return {0, a.F};
// second return val = intersection point if first
    return val = 1, otherwise undefined
// might miss an intersection due to precision issues if // get polygon area
     coordinates are too large, increasing epsilon works // O(n)
pair<int, point> seq_intersect(line_segment a,
                                                           // TESTED
    line segment b) {
                                                           ct pgon_area(polygon pg) {
        ct alen = segment_len(a);
                                                                   ct cres = 0;
        ct blen = segment len(b);
                                                                   for (int i = 0; i < pq.size()-1; ++i) {
                                                                           cres += cross(pg[i], pg[i+1]);
        if (equal(alen, (ct)0) && equal(blen, (ct)0)) {
                return (equal(a.F, b.F) ? make_pair(1, a
                                                                   return (abs(cres)/(ct)2.0);
                    .F) : make_pair(0, a.F));
        else if (equal(alen, (ct)0)) {
                                                            // check if point is inside polygon
                                                           // 0 = outside, 1 = inside, 2 = on polygon edge
                return (point_on_seg(a.F, b) ? make_pair
                                                           // O(n)
                    (1, a.F) : make_pair(0, a.F));
                                                           // TESTED
        else if (equal(blen, (ct)0)) {
                                                           int point_in_pgon(point a, polygon pg) {
                return (point_on_seg(b.F, a) ? make_pair
                                                                   for (int i = 0; i < pq.size()-1; ++i) {</pre>
                    (1, b.F) : make_pair(0, b.F));
                                                                           if (point_on_seq(a, line_segment(pg[i],
                                                                               pg[i+1]))) {
                                                                                    return 2;
```

```
// arbitrary angle, try to avoid polygon
            vertices (likely lattice points)
        line_segment tl = line_segment(a, {(ct)1092854,
            (ct)1085417});
        int icnt = 0;
        for (int i = 0; i < pg.size()-1; ++i) {</pre>
                auto cur = seg_intersect(t1,
                    line_segment(pg[i], pg[i+1]));
                if (cur.F == 1) {
                        icnt++;
        return (icnt%2 == 1);
// return the points that form given point set's convex
// O(n log n)
vector<point> convex_hull(vector<point> ps) {
        vector<point> ch;
        sort(ps.begin(), ps.end(), point_comp);
    for (int cv = 0; cv < 2; ++cv) {</pre>
        for (int i = 0; i < ps.size(); ++i) {</pre>
            int cs = ch.size();
            while (cs \ge 2 \&\& ccw(ch[cs-2], ch[cs-1], ps
                [i])) {
                ch.pop_back();
                --cs;
            ch.push_back(ps[i]);
        ch.pop_back();
        reverse(ps.begin(), ps.end());
    return ch;
```

6 Graph algorithms

6.1 Kosaraju's algorithm

Finds strongly connected components in a directed graph in O(n+m).

- 1. Create an inverse graph where all edges are reversed.
- 2. Do a DFS traversal on original graph and add all nodes in post-order to a vector.
- 3. Reverse the obtained vector.
- 4. Iterate the vector. If a node doesn't belong to a component, create new component and assign current node to it, and do a DFS in inverse graph from current node and add all reachable nodes to the component that was just created.

6.2 Bridges

An edge u-v is a bridge if there is no edge from the subtree of v to any node with lower depth than u in DFS tree. O(n+m).

```
#include <iostream>
#include <vector>
#include <algorithm>
using namespace std;
int n, m;
vector<int> g[200010];
int v[200010];
int d[200010];
// found bridges
vector<pair<int, int>> res;
// find bridges
int bdfs(int s, int cd, int p) {
    if (v[s]) return d[s];
```

```
v[s] = 1;
d[s] = cd;

int minh = cd;

for (int a : g[s]) {
    if (a == p) continue;
      minh = min(minh, bdfs(a, cd+1, s));
}

if (p != -1) {
    if (minh == cd) {
      res.push_back({s, p});
    }
}

return minh;
}

int main() {
    for (int i = 1; i <= n; ++i) {
      if (!v[i]) bdfs(i, 1, -1);
    }
}</pre>
```

6.3 Articulation points

A vertex u is an articulation point if there is no edge from the subtree of u to any parent of u in DFS tree, or if u is the root of DFS tree and has at least 2 children. O(n+m) if removing duplicates doesn't count.

Set res can be replaced with a vector if duplicates are removed afterwards.

```
#include <iostream>
#include <vector>
#include <algorithm>
#include <set>

using namespace std;
int n, m;
vector<int> g[200010];
```

```
int dt[200010];
int low[200010];
// found articulation points
// can be replaced with vector, but duplicates must be
    removed
set<int> res;
int curt = 1;
void adfs(int s, int p) {
    if (v[s]) return;
    v[s] = 1;
    dt[s] = curt++;
    low[s] = dt[s];
    int ccount = 0;
    for (int a : q[s]) {
        if (!v[a]) {
            ++ccount;
            adfs(a, s);
            low[s] = min(low[s], low[a]);
            if (low[a] >= dt[s] && p != -1) res.insert(s
                );
        else if (a != p) {
            low[s] = min(low[s], dt[a]);
        if (p == -1 && ccount > 1) {
            res.insert(s);
int main() {
    for (int i = 1; i <= n; ++i) {</pre>
        if (!v[i]) adfs(i, -1);
```

int v[200010];

6.4 Maximum flow (scaling algorithm)

Scaling algorithm, uses DFS to find an augmenting path where each edge weight is larger than or equal to a certain threshold. Time complexity $O(m^2 \; log(c))$, where c is the starting threshold (sum of all edge weights in the graph).

```
#include <iostream>
#include <vector>
#include <algorithm>
using namespace std;
typedef long long 11;
const int N = 105; // vertex count
const 11 LINF = 1000000000000000005;
int n, m;
vector<int> q[N];
ll d[N][N]; // edge weights
int v[N];
vector<int> cp; // current augmenting path
11 \text{ res} = 0;
// find augmenting path using scaling
// prerequisities: clear current path, divide threshold
    by 2, increment cvis
ll dfs(int s, int t, ll thresh, int cvis, ll cmin) {
    if (v[s] == cvis) return -1;
    v[s] = cvis;
    cp.push_back(s);
    if (s == t) return cmin;
    for (int a : q[s]) {
        if (d[s][a] < thresh) continue; // scaling</pre>
        ll cres = dfs(a, t, thresh, cvis, min(cmin, d[s
            ][a]));
        if (cres != -1) return cres;
    cp.pop_back();
    return -1;
```

```
int main()
    ios_base::sync_with_stdio(false);
    cin.tie(0);
    cin >> n >> m;
    11 \text{ cthresh} = 0;
    for (int i = 0; i < m; ++i) {</pre>
        int a, b;
        11 c;
        cin >> a >> b >> c;
        g[a].push_back(b);
        q[b].push_back(a);
        d[a][b] += c;
        d[b][a] = 0;
        cthresh += c;
    int cvis = 0;
    while (true) {
        cvis++;
        cp.clear();
        11 minw = dfs(1, n, cthresh, cvis, LINF);
        if (minw != -1) {
            res += minw;
            for (int i = 0; i < cp.size()-1; ++i) {</pre>
                d[cp[i]][cp[i+1]] -= minw;
                d[cp[i+1]][cp[i]] += minw;
        else {
            if (cthresh == 1) break;
            cthresh /= 2:
    cout << res << "\n";
    return 0;
```

6.5 Theorems on flows and cuts

Maximum flow is always equal to minimum cut. Minimum cut can be found by running a maximum flow algorithm and dividing the resulting flow graph into two sets of vertices. Set A contains all vertices that can be reached from source using positive-weight

edges. Set B contains all other vertices. Minimum cut consists of the edges between these two sets.

Number of edge-disjoint (= each edge can be used at most once) paths in a graph is equal to maximum flow on graph where capacity of each edge is 1.

Number of vertex-disjoint paths can be found the same way as edge-disjoint paths, but each vertex is duplicated and an edge is added between the two vertices. All incoming edges go to the first vertex and all outgoing edges start from the second vertex.

Maximum matching of a bipartite graph can be found by adding a source and a sink to the graph and connecting source to all left vertices and sink to all right vertices. Maximum matching equals maximum flow on this graph.

König's theorem: sizes of a minimum vertex cover (= minimum set of vertices such that each edge has at least one endpoint in the set) and a maximum matching are always equal in a bipartite graph. Maximum independent set (= maximum set of vertices such that no two vertices in the set are connected with an edge) consists of the vertices not in a minimum vertex cover.

6.6 Heavy-light decomposition

Supports updates and queries on path between two vertices a and b in $O(log^2(n))$.

Doesn't explicitly look for LCA, instead climbs upwards from the lower chain until both vertices are in the same chain.

Requires a segment tree implementation that corresponds to the queries. Lazy segtree, for example, can be pasted directly in.

```
#include <iostream>
#include <vector>
#include <algorithm>

using namespace std;
typedef long long ll;

const int S = 100005; // vertex count
const int N = (1<<18); // seqtree size, must be >= S
```

```
vector<int> q[S];
int sz[S], de[S], pa[S];
int cind[S], chead[S], cpos[S];
int cchain, cstind, stind[S];
// IMPLEMENT SEGMENT TREE HERE
// st_update() and st_query() should call segtree
    functions
ll st[2*N];
void hdfs(int s, int p, int cd) {
    de[s] = cd;
    pa[s] = p;
    sz[s] = 1;
    for (int a : q[s]) {
        if (a == p) continue;
        hdfs(a, s, cd+1);
        sz[s] += sz[a];
void hld(int s) {
    if (chead[cchain] == 0) {
        chead[cchain] = s;
        cpos[s] = 0;
    else {
        cpos[s] = cpos[pa[s]]+1;
    cind[s] = cchain;
    stind[s] = cstind;
    cstind++;
    int cmx = 0, cmi = -1;
    for (int i = 0; i < g[s].size(); ++i) {</pre>
        if (g[s][i] == pa[s]) continue;
        if (sz[g[s][i]] > cmx) {
            sz[g[s][i]] = cmx;
            cmi = i;
```

```
if (cmi != -1) {
        hld(q[s][cmi]);
    for (int i = 0; i < g[s].size(); ++i) {</pre>
        if (i == cmi) continue;
        if (g[s][i] == pa[s]) continue;
        cchain++;
        cstind++;
        hld(g[s][i]);
// do a range update on underlying segtree
// sa and sb are segtree indices
void st_update(int sa, int sb, ll x) {
// do a range query on underlying segtree
// sa and sb are segtree indices
11 st_query(int sa, int sb) {
}
// update all vertices on path from vertex a to b
// a and b are vertex numbers
void path_update(int a, int b, ll x) {
    while (cind[a] != cind[b]) {
        if (de[chead[cind[b]]] > de[chead[cind[a]]])
            swap(a, b);
        st_update(stind[chead[cind[a]]], stind[a], x);
        a = pa[chead[cind[a]]];
    if (stind[b] < stind[a]) swap(a, b);</pre>
    st_update(stind[a], stind[b], x);
// query all vertices on path from vertex a to b
// a and b are vertex numbers
11 path_query(int a, int b) {
        11 cres = 0; // set to identity
        while (cind[a] != cind[b]) {
        if (de[chead[cind[b]]] > de[chead[cind[a]]])
            swap(a, b);
```

7 Tree algorithms

7.1 Smaller to larger

Answers queries offline on entire subtrees or specifically on vertices with depth d in a subtree. Normally $O(n \ log \ n)$ for all queries, the complexity may worsen depending on what is stored for each node. If the depth is queried on, merge to the deepest subtree, otherwise to the largest one. When storing data for each depth, store the highest vertex last so it's efficient to append higher vertices.

```
int n, q;

vector<int> g[N];
vector<int> nd[N]; // subtree root -> depth -> data,
    highest vertex is the last one

vector<int> nq[N]; // queries for each vertex
vector<pair<int, int>> rq; // raw queries in original
    order
```

```
map<int, int> res[N];
void dfs(int s, int p) {
        // find deepest subtree
        int mxs = 0, mxi = -1;
        for (int i = 0; i < g[s].size(); ++i) {</pre>
                int a = q[s][i];
                if (a == p) continue;
                dfs(a, s);
                if (nd[a].size() > mxs) {
                        mxs = nd[a].size();
                        mxi = i;
        // swap deepest subtree with current one
        if (mxi != -1) {
                swap(nd[s], nd[g[s][mxi]]);
        // merge shallower subtrees to the largest one
        for (int i = 0; i < g[s].size(); ++i) {</pre>
                int a = g[s][i];
                if (a == p || i == mxi) continue;
                for (int j = 0; j < nd[a].size(); ++j) {</pre>
                        int sr = nd[a].size()-(j+1); //
                             source
                        int de = nd[s].size()-(j+1); //
                             destination
                         // merge vertices with same
                             depth
                        nd[s][de] += nd[a][sr];
        // add current vertex
        nd[s].push back(1);
        // nd[s] represents now the subtree of s
        // answer all queries on this subtree offline
            and store the answers
        for (int de : nq[s]) {
                int di = nd[s].size()-(de+1);
                if (di < 0) res[s][de] = 0;
                else res[s][de] = nd[s][di]-1;
```

7.2 Subtree merging DP

For each subtree of a tree, some DP is calculated for each vertex by merging all child subtrees of the vertex together one by one. Basically we take a elements from current subtree root and the already merged child subtrees and b elements from the child subtree being merged. This is the technique used in Looking for a Challenge - Barricades.

The algorithm looks like $O(n^3)$, but actually runs in $O(n^2)$.

```
#include <iostream>
#include <vector>
using namespace std;

const int N = 3005;
const int INF = 1000000005;
int n, m;
vector<int> g[N];
int sz[N];
```

```
// dp[i][j] = min number of blocked edges to get a
                                                                                     // In Barricades, innermost loop
    security zone of
// size j in the subtree of vertex i such that i is in
                                                                                     // dp[s][a]++;
    the zone
int dp[N][N];
                                                                             // now v is completely merged, count its
                                                                                  size
// Looking for a challenge: Barricades style
                                                                             sz[s] += sz[v];
// Merge child subtrees of s to s one-by-one
// Runs in O(n^2) even though it looks like O(n^3)
void solve(int s, int p) {
        // maintain the combined size of already merged
                                                            int main() {
            child subtrees
                                                                     for (int i = 0; i <= n; ++i) {</pre>
                                                                             for (int j = 0; j <= n; ++j) {
        sz[s] = 1;
                                                                                     dp[i][j] = INF;
        // initial dp conditions (how to solve if s is a
             leaf node)
                                                                     solve(1, -1);
        dp[s][1] = 0;
                                                                     return 0;
        // merge the subtree of v to (s + previous v:s)
        // first v requires no special case, since we
            just merge to s
        for (int v : q[s]) {
                                                                 String algorithms
                if (v == p) continue;
                solve(v, s);
                                                                  Polynomial hashing
                // take a elements from already merged
                     ones and b from the subtree of v
                                                            If hash collisions are likely, compute two hashes with two distinct
                // we don't need an auxiliary dp array
                                                            pairs of constants of magnitude 10^9 and use their product as the
                    since we write to larger indices
                                                            actual hash.
                // from where we read during current
                                                            #include <iostream>
                    subtree merge operation ((a+b) > a)
                for (int a = sz[s]; a >= 0; --a) {
                        for (int b = 0; b <= sz[v]; ++b)</pre>
                                                            using namespace std;
                                 // do dp transition here
                                                            const 11 A = 957262683;
                                                            const 11 B = 998735246;
                                 dp[s][a+b] = min(dp[s][a
                                     +b], dp[s][a] + dp[v]
                                                            string s;
                                     ][b]);
                                                            ll h[1000005];
                                                            ll p[1000005];
                        // Barricades specific: if we
                                                            11 ghash(int a, int b) {
                             take 0 nodes from v, we have
                                                                    if (a == 0) return h[b];
                              to
                                                                    ll cres = (h[b]-h[a-1]*p[b-a+1])%B;
                        // block the edge to v
```

should start from b=1

```
if (cres < 0) cres += B;
    return cres;
}

int main() {
    cin >> s;

    h[0] = s[0];
    p[0] = 1;

for (int i = 1; i < s.length(); ++i) {
        h[i] = (h[i-1]*A+s[i])%B;
        p[i] = (p[i-1]*A)%B;
    }
    return 0;
}</pre>
```

8.2 Z-algorithm

Constructs the Z-array for string s. Z-array tells for each i the length of the longest substring that begins at i and is a prefix of s. O(n).

```
vector<int> z_alg(string s) {
   int cn = s.size();
   vector<int> z(cn);
   int x = 0;
   int y = 0;
   for (int i = 1; i < cn; ++i) {
        z[i] = max(0, min(z[i-x], y-i+1));
        while (i+z[i] < cn && s[z[i]] == s[i+z[i]])) {
            x = i;
            y = i+z[i];
            z[i]++;
        }
   }
   return z;
}</pre>
```

8.3 Suffix array

Constructs the suffix array for string s. By default, the array is a cyclic suffix array which has all the cyclic rotations of the string in lexicographic order. Creates a normal suffix array if \$ is appended to the string. In that case the first element in the suffix array must be discarded

```
// creates a circular suffix array (sorted array of
    cyclic rotations)
// to get a normal suffix array, add $ to the end of the
     string
// and discard the first element of returned suffix
// n = 7*10^5 takes around 1 second
vector<int> suffix_array(string cs) {
        int cn = (int)cs.length();
        int MXN = cn+256; // size of alphabet
        vector<int> sa(cn), ra(cn);
        for (int i = 0; i < cn; ++i) {</pre>
                sa[i] = i;
                ra[i] = (int)cs[i];
        for (int k = 0; k < cn; k ? k *= 2 : ++k) {
                vector<int> nsa(sa), nra(cn), ccnt(MXN);
                for (int i = 0; i < cn; ++i) {</pre>
                        nsa[i] = (nsa[i]-k+cn)%cn;
                        ccnt[ra[i]]++;
                for (int i = 1; i < MXN; ++i) {</pre>
                        ccnt[i] += ccnt[i-1];
                for (int i = cn-1; i >= 0; --i) {
                        sa[--ccnt[ra[nsa[i]]]] = nsa[i];
                int r = 0;
                for (int i = 1; i < cn; ++i) {</pre>
                        if (ra[sa[i]] != ra[sa[i-1]]) {
                                 r++;
                         else if (ra[(sa[i] + k)%cn] !=
                             ra[(sa[i-1] + k)%cn]) {
```

```
r++;
}
nra[sa[i]] = r;
}
ra = nra;
}
return sa;
```