Chapter 1

Fundamentals

1.1 Models of Computation

When designing algorithms, one has to agree on a model of computation according to which these algorithms can be executed. There are various such models, but when it comes to geometry some are more convenient to work with than others. Even using very elementary geometric operations—such as taking the center of a circle defined by three points or computing the length of a given circular arc—the realms of rational and even algebraic numbers are quickly left behind. Representing the resulting real numbers/coordinates would be a rather painful task in, for instance, a Turing machine type model of computation.

Therefore, other models of computation are more prominent in the area of geometric algorithms and data structures. In this course we will be mostly concerned with two models: the Real RAM and the algebraic computation/decision tree model. The former is rather convenient when designing algorithms, because it sort of abstracts from the aforementioned representation issues by simply assuming that it can be done. The latter model typically appears in the context of lower bounds, that is, proofs that certain problems cannot be solved more efficiently than some function depending on the problem size (and possibly some other parameters). So let us see what these models are in more detail.

Real RAM Model. A memory cell stores a real number (that is what the "Real" stands for)¹. Any single arithmetic operation (addition, subtraction, multiplication, division, and k-th root, for small constant k) or comparison can be computed in constant time.² This is a quite powerful (and somewhat unrealistic) model of computation, as a single real number in principle can encode an arbitrary amount of information. Therefore we

¹RAM stands for random access machine, meaning that every memory cell can be accessed in constant time. Not like, say, a list where one always has to start from the first element.

²In addition, sometimes also logarithms, other analytic functions, indirect addressing (integral), or floor and ceiling are used. As adding some of these operations makes the model more powerful, it is usually specified and emphasized explicitly when an algorithm uses them.

have to ensure that we do not abuse the power of this model. For instance, we may want to restrict the numbers that are manipulated by any single arithmetic operation to be bounded by some fixed polynomial in the numbers that appear in the input.

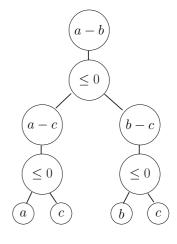
On the positive side, the real RAM model allows to abstract from the lowlands of numeric and algebraic computation and to concentrate on the algorithmic core from a combinatorial point of view.

But there are also downsides to using such a powerful model. First, it may be a challenge to efficiently implement a geometric algorithm designed for the real RAM on an actual computer. With bounded memory there is no way to represent general real numbers explicitly, and operations using a symbolic representation can hardly be considered constant time. Second, it is difficult if not impossible to derive reasonable lower bounds in the real RAM model.

Therefore, when interested in lower bounds, it is convenient to use a different, less powerful model of computation. One such model is the computation tree model, which encompasses and explicitly represents all possible execution paths of an algorithm.

Algebraic Computation Trees (Ben-Or [1]). A computation is regarded as a rooted binary tree, that is, each node has at most two children.

- The leaves contain the (possible) results of the computation.
- Every node ν with one child has an operation of the form $+,-,*,/,\sqrt{,\ldots}$ associated to it. The operands of this operation are constant input values, or among the ancestors of ν in the tree.
- Every node ν with two children has associated to it a branching of the form > 0, $\geqslant 0$, or = 0. The branch is with respect to the result of ν 's parent node. If the expression yields true, the computation continues with the left child of ν ; otherwise, it continues with the right child of ν .



The term decision tree is used if all of the final results (leaves) are either true or false. If every branch is based on a linear function in the input values, we face a linear decision tree. Analogously one can define, say, quadratic decision trees.

The complexity of a computation or decision tree is the maximum number of vertices along any root-to-leaf path. It is well known that $\Omega(n \log n)$ comparisons are required to sort n numbers. But also for some problems that appear easier than sorting at first glance, the same lower bound holds. Consider, for instance, the following problem.

Element Uniqueness

Input: $\{x_1, \ldots, x_n\} \subset \mathbb{R}, n \in \mathbb{N}$.

Output: Is $x_i = x_j$, for some $i, j \in \{1, ..., n\}$ with $i \neq j$?

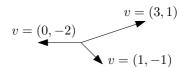
Ben-Or [1] has shown that any algebraic decision tree to solve Element Uniqueness for n elements has complexity $\Omega(n \log n)$.

1.2 **Basic Geometric Objects**

We will mostly be concerned with the d-dimensional Euclidean space \mathbb{R}^d , for small $d \in \mathbb{N}$; typically, d = 2 or d = 3. The basic objects of interest in \mathbb{R}^d are the following.

Points. A point
$$p \in \mathbb{R}^d$$
, typically described by its d Cartesian coordinates $p = (x_1, \dots, x_d)$.

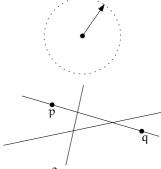
Vectors. A vector
$$v \in \mathbb{R}^d$$
, typically described by its d $v = (0, -2)$ Cartesian coordinates $v = (x_1, \dots, x_d)$.



What is the difference between a point and a vector? Mathematically, points and vectors are the same, they are both elements of Euclidean space \mathbb{R}^d . The different terms are still useful in order to indicate how we think of such an element in a given context. We think of a point as a location in space (a "dot"), while we think of a vector as a translation in space (an "arrow" starting from the origin). The point view is dominant in geometry, but every point is also a vector (the arrow from the origin to the dot), so we can seamlessly apply vector space operations (addition, scalar multiplication) to points. The resulting vector is also a point (the dot at the head of the arrow). There are a number of sources saying that points and vectors are not the same, and they are right when it comes to how we think about them. But when it comes to what they actually are, there is no need to make a difference.

Directions. A vector $v \in S^{d-1}$ (the (d-1)-dimensional unit sphere), typically described by its d Cartesian coordinates $\nu=(x_1,\ldots,x_d),$ with $\|\nu\|=\sqrt{\sum_{i=1}^d x_i{}^2}=1.$

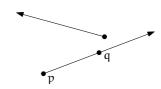
Lines. A line is a one-dimensional affine subspace. It can be described by two distinct points p and q as the set of all points r that satisfy $r = p + \lambda(q - p)$, for some $\lambda \in \mathbb{R}$.



While any pair of distinct points defines a unique line, a line in \mathbb{R}^2 contains infinitely many points and so it may happen that a collection of three or more points lie on a line. Such a collection of points is termed *collinear*³.

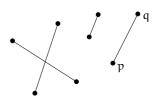
³Not colinear, which refers to a notion in the theory of coalgebras.

Rays. If we remove a single point from a line and take the closure of one of the connected components, then we obtain a ray. It can be described by two distinct points p and q as the set of all points r that satisfy $r = p + \lambda(q-p)$, for some $\lambda \geqslant 0$. The *orientation* of a ray is the direction $(q-p)/\|q-p\|$.



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Line segment. A line segment is a compact connected subset of a line. It can be described by two points p and q as the set of all points r that satisfy $r=p+\lambda(q-p)$, for some $\lambda\in[0,1]$. We will denote the line segment through p and q by \overline{pq} . Depending on the context we may allow or disallow degenerate line segments consisting of a single point only (p=q) in the above equation).



Hyperplanes. A hyperplane h is a (d-1)-dimensional affine subspace. It can be described algebraically by d+1 coefficients $h_1, \ldots, h_{d+1} \in \mathbb{R}$, as the set of all points (x_1, \ldots, x_d) that satisfy the linear equation $\sum_{i=1}^d h_i x_i = h_{d+1}$. Here, at least one of h_1, \ldots, h_d must be nonzero; otherwise, the equation is either satisfied by all points (if $h_{d+1} = 0$), or by no point (if $h_{d+1} \neq 0$). Such *degenerate* hyperplanes are useful in some contexts, and where we allow them, we explicitly say so.

If the above equation is converted into an inequality, we obtain the algebraic description of a *halfspace* (in \mathbb{R}^2 : halfplane).

Spheres and balls. A sphere is the set of all points that are equidistant to a fixed point. It can be described by a point c (center) and a number $\rho \in \mathbb{R}$ (radius) as the set of all points p that satisfy $||p-c|| = \rho$. The ball of radius ρ around c consists of all points p that satisfy $||p-c|| \leqslant \rho$.

1.3 Graphs

In this section we review some basic definitions and properties of graphs. For more details and proofs, refer to any standard textbook on graph theory [2, 3, 5].

A (simple undirected) graph G = (V, E) is defined on a set V of *vertices*. Unless explicitly stated otherwise, V is always finite. Vertices are associated to each other through *edges* which are collected in the set $E \subseteq \binom{V}{2}$. The two vertices defining an edge are *adjacent* to each other and *incident* to the edge. To avoid clutter we often omit brackets and write uv for an edge $\{u,v\}$.

For a vertex $v \in V$, denote by $N_G(v)$ the neighborhood of v in G, that is, the set of vertices from G that are adjacent to v. Similarly, for a set $W \subset V$ of vertices its neighborhood $N_G(W)$ is defined as $\bigcup_{w \in W} N_G(w)$. The degree $\deg_G(v)$ of a vertex $v \in V$

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is the size of its neighborhood, that is, the number of edges from E incident to ν . The subscript is often omitted if the graph under consideration is clear from the context.

Lemma 1.1 (Handshaking Lemma). In any graph G = (V, E) we have

$$\sum_{\nu \in V} \deg(\nu) = 2|E|.$$

Two graphs G = (V, E) and H = (U, W) are *isomorphic* if there is a bijection $\phi : V \to U$ such that $\{u, v\} \in E \iff \{\phi(u), \phi(v)\} \in W$. Such a bijection ϕ is called an *isomorphism* between G and H. The structure of isomorphic graphs is identical and often we do not distinguish between them when looking at them as graphs.

For a graph G denote by V(G) the set of vertices and by E(G) the set of edges. A graph H = (U, F) is a *subgraph* of G if $U \subseteq V$ and $F \subseteq E$. In case that U = V the graph H is a *spanning* subgraph of G. For a set $W \subseteq V$ of vertices denote by G[W] the *induced subgraph* of W in G, that is, the graph $(W, E \cap \binom{W}{2})$. For $F \subseteq E$ let $G \setminus F := (V, E \setminus F)$. Similarly, for $W \subseteq V$ let $G \setminus W := G[V \setminus W]$. In particular, for a vertex or edge $x \in V \cup E$ we write $G \setminus x$ for $G \setminus \{x\}$. The *union* of two graphs G = (V, E) and H = (W, F) is the graph $G \cup H := (V \cup W, E \cup F)$.

For an edge $e = \{u, v\} \in E$ the graph G/e is obtained from $G \setminus \{u, v\}$ by adding a new vertex w with $N_{G/e}(w) := (N_G(u) \cup N_G(v)) \setminus \{u, v\}$. This process is called *contraction* of e in G. Similarly, for a set $F \subseteq E$ of edges the graph G/F is obtained from G by contracting all edges from F (the order in which the edges from F are contracted does not matter).

Graph traversals. A walk in G is a sequence $W = (v_1, \ldots, v_k)$, $k \in \mathbb{N}$, of vertices such that v_i and v_{i+1} are adjacent in G, for all $1 \le i < k$. The vertices v_1 and v_k are referred to as the walk's endpoints, the other vertices are called interior. A walk with endpoints v_1 and v_k is sometimes referred to as a walk between v_1 and v_k . For a walk W denote by V(W) its set of vertices and by E(W) its set of edges (pairs of vertices adjacent along W). We say that W visits the vertices and edges in $V(W) \cup E(W)$. A walk for which both endpoints coincide, that is, $v_1 = v_k$, is called closed. Otherwise the walk is open.

If a walk uses each edge of G at most once, it is a *trail*. A closed walk that visits each edge and each vertex at least once is called a *tour* of G. An *Euler tour* is both a trail and a tour of G, that is, it visits each edge of G exactly once. A graph that contains an Euler tour is termed *Eulerian*.

If the vertices v_1, \ldots, v_k of a closed walk W are pairwise distinct except for $v_1 = v_k$, then W is a cycle of size k-1. If the vertices v_1, \ldots, v_k of a walk W are pairwise distinct, then W is a path of size k. A Hamilton cycle (path) is a cycle (path) that visits every vertex of G. A graph that contains a Hamilton cycle is Hamiltonian.

Two trails are *edge-disjoint* if they do not share any edge. Two paths are called (internally) vertex-disjoint if they do not share any vertices (except for possibly common endpoints). For two vertices $s, t \in V$ any path with endpoints s and t is called an (s, t)-path or a path between s and t.

Connectivity. Define an equivalence relation " \sim " on V by setting $\alpha \sim b$ if and only if there is a path between α and b in G. The equivalence classes with respect to " \sim " are called *components* of G and their number is denoted by $\omega(G)$. A graph G is *connected* if $\omega(G) = 1$ and *disconnected*, otherwise.

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A set $C \subset V$ of vertices in a connected graph G = (V, E) is a *cut-set* of G if $G \setminus C$ is disconnected. A graph is *k-connected*, for a positive integer k, if $|V| \geqslant k+1$ and there every cut-set has at least k vertices. Similarly a graph G = (V, E) is *k-edge-connected*, if $G \setminus F$ is connected, for any set $F \subseteq E$ of at most k-1 edges. Connectivity and cut-sets are related via the following well-known theorem.

Theorem 1.2 (Menger [4]). For any two nonadjacent vertices u, v of a graph G = (V, E), the size of a minimum cut that disconnects u and v is the same as the maximum number of pairwise internally vertex-disjoint paths between u and v.

Specific families of graphs. A graph with a maximum number of edges, that is, $(V, \binom{V}{2})$, is called a *clique*. Up to isomorphism there is only one clique on n vertices; it is referred to as the *complete graph* K_n , for $n \in \mathbb{N}$. At the other extreme, the *empty graph* $\overline{K_n}$ consists of n isolated vertices that are not connected by any edge. A set U of vertices in a graph G is *independent* if G[U] is an empty graph. A graph whose vertex set can be partitioned into at most two independent sets is *bipartite*. An equivalent characterization states that a graph is bipartite if and only if it does not contain any odd cycle. The bipartite graphs with a maximum number of edges (unique up to isomorphism) are the *complete bipartite graphs* $K_{m,n}$, for $m,n \in \mathbb{N}$. They consist of two disjoint independent sets of size m and n, respectively, and all mn edges in between.

A forest is a graph that is acyclic, that is, it does not contain any cycle. A connected forest is called *tree* and its *leaves* are the vertices of degree one. Every connected graph contains a spanning subgraph which is a tree, a so called *spanning tree*. Beyond the definition given above, there are several equivalent characterizations of trees.

Theorem 1.3. The following statements for a graph G are equivalent.

- (1) G is a tree (i.e., it is connected and acyclic).
- (2) G is a connected graph with n vertices and n-1 edges.
- (3) G is an acyclic graph with n vertices and n-1 edges.
- (4) Any two vertices in G are connected by a unique path.
- (5) G is minimally (edge-)connected, that is, G is connected but removal of any single edge yields a disconnected graph.
- (6) G is maximally acyclic, that is, G is acyclic but adding any single edge creates a cycle.

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Directed graphs. In a directed graph or, short, $digraph\ D=(V,E)$ the set E consists of ordered pairs of vertices, that is, $E\subseteq V^2$. The elements of E are referred to as arcs. To avoid clutter we often omit brackets and write uv for an arc (u,v). An arc $uv\in E$ is said to be directed from its $source\ u$ to its $target\ v$. For $uv\in E$ we also say "there is an arc from u to v in D". Usually, we consider loop-free graphs, that is, arcs of the type vv, for some $v\in V$, are not allowed.

The in-degree $\deg_D^-(v) := |\{(u,v)|uv \in E\}|$ of a vertex $v \in V$ is the number of incoming arcs at v. Similarly, the out-degree $\deg_D^+(v) := |\{(v,u) \mid vu \in E\}|$ of a vertex $v \in V$ is the number of outgoing arcs at v. Again the subscript is often omitted when the graph under consideration is clear from the context.

From any undirected graph G one can obtain a digraph on the same vertex set by specifying a direction for each edge of G. Each of these $2^{|E(G)|}$ different digraphs is called an *orientation* of G. Similarly every digraph D = (V, E) has an *underlying* undirected graph $G = (V, \{\{u, v\} \mid (u, v) \in E \text{ or } (v, u) \in E\})$. Hence most of the terminology for undirected graphs carries over to digraphs.

A directed walk in a digraph D is a sequence $W = (v_1, \ldots, v_k)$, for some $k \in \mathbb{N}$, of vertices such that there is an arc from v_i to v_{i+1} in D, for all $1 \le i < k$. In the same way we define directed trails, directed paths, directed cycles, and directed tours.

Multigraphs. Sometimes we also consider *multigraphs*, where edges have a multiplicity, that is, there may be several copies of an edge. Unless forbidden explictly, a multigraph may contain loops. Just as simple graphs, multigraphs may be undirected or directed, and also most of the other basic notions for graphs discussed above naturally generalize to multigraphs.

References

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