

# Exhaustive Search, Combinatorial Optimization and Enumeration: Exploring the Potential of Raw Computing Power

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**Abstract.** For half a century since computers came into existence, the goal of finding elegant and efficient algorithms to solve “simple” (well-defined and well-structured) problems has dominated algorithm design. Over the same time period, both processing and storage capacity of computers have increased by roughly a factor of a million. The next few decades may well give us a similar rate of growth in raw computing power, due to various factors such as continuing miniaturization, parallel and distributed computing. If a quantitative change of orders of magnitude leads to qualitative advances, where will the latter take place? Only empirical research can answer this question.

Asymptotic complexity theory has emerged as a surprisingly effective tool for predicting run times of polynomial-time algorithms. For NP-hard problems, on the other hand, it yields overly pessimistic bounds. It asserts the non-existence of algorithms that are efficient across an entire problem class, but ignores the fact that many instances, perhaps including those of interest, can be solved efficiently. For such cases we need a complexity measure that applies to problem instances, rather than to over-sized problem classes.

Combinatorial optimization and enumeration problems are modeled by state spaces that usually lack any regular structure. Exhaustive search is often the only way to handle such “combinatorial chaos”. Several general purpose search algorithms are used under different circumstances. We describe reverse search and illustrate this technique on a case study of enumerative optimization: enumerating the  $k$  shortest Euclidean spanning trees.

## 1 Catching Up with Technology

Computer science is technology-driven: it has been so for the past 50 years, and will remain this way for the foreseeable future, for at least a decade. That is about as far as specialists can extrapolate current semiconductor technology and foresee that advances based on refined processes, without any need for fundamental innovations, will keep improving the performance of computing devices. Moreover, performance can be expected to advance at the rate of “Moore’s law”,

the same rate observed over the past 3 decades, of doubling in any period of 1 to 2 years. An up-to-date summary of possibilities and limitations of technology can be found in [12].

What does it mean for a discipline to be *technology-driven*? What are the implications?

Consider the converse: disciplines that are demand-driven rather than technology-driven. In the 60s the US stated a public goal “to put a man on the moon by the end of the decade”. This well-defined goal called for a technology that did not as yet exist. With a great national effort and some luck, the technology was developed just in time to meet the announced goal — a memorable technical achievement. More often than not, when an ambitious goal calls for the invention of new technology, the goal remains wishful thinking. In such situations, it is prudent to announce fuzzy goals, where one can claim progress without being able to measure it. The computing field has on occasion been tempted to use this tactic, for example in predicting “machine intelligence”. Such an elastic concept can be re-defined periodically to mirror the current state-of-the-art.

Apart from some exceptions, the dominant influence on computing has been a technology push, rather than a demand pull. In other words, computer architects, systems and application designers have always known that clock rates, flop/s, data rates and memory sizes will go up at a predictable, breath-taking speed. The question was less “what do we need to meet the demands of a new application?” as “what shall we do with the newly emerging resources?”. Faced with an embarrassment of riches, it is understandable that the lion’s share of development effort, both in industry and in academia, has gone into developing bigger, more powerful, hopefully a little better versions of the same applications that have been around for decades. What we experience as revolutionary in the break-neck speed with which computing is affecting society is not technical novelty, but rather, an unprecedented penetration of computing technology in all aspects of the technical infrastructure on which our civilization has come to rely. In recent years, computing technology’s outstanding achievement has been the *breadth* of its impact rather than the originality and depth of its scientific/technical innovations. The explosive spread of the Internet in recent years, based on technology developed a quarter-century ago, is a prominent example.

This observation, that computing technology in recent years has been “spreading inside known territory” rather than “growing into new areas”, does not imply that computing has run out of important open problems or new ideas. On the contrary, tantalizing open questions and new ideas call for investigation, as the following two examples illustrate:

1. A challenging, fundamental open problem in our “information age”, is a scientific definition of information. Shannon’s pioneering information theory is of unquestioned importance, but it does *not* capture the notion of “information” relevant in daily life (“what is your telephone number?”) or in business transactions (“what is today’s exchange rate”). The fact that we process information at all times without having a scientific definition of what we are processing is akin to the state of physics before Newton: humanity

has always been processing energy, but a scientific notion of energy emerged only three centuries ago. This discovery was a prerequisite of the industrial age. Will we ever discover a scientific notion of “information” of equal rigor and impact?

2. Emerging new ideas touch the very core of what “computing” means. Half a century of computing has been based on electronic devices to realize a von Neumann architecture. But computation can be modeled by many other physical phenomena. In particular, the recently emerged concepts of quantum computing or DNA computing are completely different models of computation, with strikingly different strengths and weaknesses.

The reason fundamentally new ideas are not receiving very much attention at the moment is that there is so much to do, and so much to gain, by merely riding the wave of technological progress, e. g. by bringing to market the next upgrade of the same old operating system, which will surely be bigger, if not better. Whereas business naturally looks for short-term business opportunities, let this be a call to academia to focus on long-term issues, some of which will surely revolutionize computing in decades to come.

## 2 “Ever-Growing” Computing Power: What Is It Good for?

Computer progress over the past several decades has measured several orders of magnitude with respect to various physical parameters such as computing power, memory size at all hierarchy levels from caches to disk, power consumption, physical size and cost. Both computing power and memory size have easily grown by a factor of a million. There are good reasons to expect both computing power and memory size to grow by the same two factors of a million over the next couple of decades. A factor of 1000 can be expected from the extrapolation of Moore’s law for yet another dozen years. Another factor of 1000, in both computing power and memory size, can be expected from increased use of parallel and distributed systems—a resource whose potential will be fully exploited only when microprocessor technology improvement slows down.

With hindsight we know what these two factors of a million have contributed to the state of the art—the majority of today’s computer features and applications would be impossible without them. The list includes:

- graphical user interfaces and multimedia in general,
- end-user application packages, such as spreadsheets,
- data bases for interactive, distributed information and transaction systems,
- embedded systems, for example in communications technology.

Back in the sixties it was predictable that computing power would grow, though the rapidity and longevity of this growth was not foreseen. People speculated what more powerful computers might achieve. But this discussion was generally limited to applications that were already prominent, such as scientific computing.

The new applications that emerged were generally not foreseen. Evidence for this is provided by quotes from famous pioneers, such as DEC founder Ken Olsen’s dictum “there is no reason why anyone would want a computer in his home” ([9] is an amusing collection of predictions).

If past predictions fell short of reality, we cannot assume that our gaze into the crystal ball will be any clearer today. We do not know what problems can be attacked with computing power a million times greater than available today. Thus, to promote progress over a time horizon of a decade or more, experimenting is a more promising approach than planning.

### 3 Uneven Progress in Algorithmics

After this philosophical excursion into the world of computing, let us now consider the small but important discipline of algorithmics, the craft of algorithm design, analysis and implementation. In the early days of computing, algorithmics was a relatively bigger part of computer science than it is now—all computer users had to know and program algorithms to solve their problem. Until the seventies, a sizable fraction of computer science research was dedicated to expand and improve our knowledge of algorithms.

The computer user community at large may be unaware that the progress of algorithmics can be considered spectacular. Whereas half a century ago an algorithm was just a prescription to be followed, nowadays an algorithm is a mathematical object whose properties are stated in terms of theorems and proofs. A large number of well-understood algorithms have proven their effectiveness, embedded in program libraries and application packages. They empower users to work with techniques that they could not possibly program themselves. Who could claim, for example, to know and be able to program all the algorithms in a symbolic computation system, or in a computational geometry library? Asymptotic complexity analysis has turned out to be a surprisingly effective technique to predict the performance of algorithms. When we classify an algorithm as running in time  $O(\log n)$  or  $O(n \log n)$ , generously ignoring constant factors, how could we expect to convert this rough measure into seconds? It’s easy: you time the program for a few small data sets, and extrapolate to obtain an accurate prediction of running times for much larger data sets, as the measurements in Table 1 illustrate.

By and large, the above claim of spectacular success is limited to algorithms that run in polynomial time, said to be in the class  $P$ . We have become so used to asymptotic complexity analysis that we forget to marvel how such a simple formula such as “ $\log n$ ” yields such accurate timing information. But we may marvel again when we consider the class of algorithms called NP-hard, which presumably require time exponential in the size  $n$  of the data set.

For these hard problems, occasionally called *intractable*, we have a theory that is not nearly as practical as the theory for  $P$ , because it yields overly pessimistic bounds. Our current theory of NP-hard problems is modeled on the same approach that worked so successfully for problems in  $P$ : prove theorems

**Table 1.** Running times of Binary Search

$n$	$\log n$	$t$ bin search	$t/\log n$
1	0	0.60	
2	1	0.81	0.81
4	2	0.91	0.45
8	3	1.08	0.36
16	4	1.26	0.32
32	5	1.46	0.29
64	6	1.66	0.27
128	7	1.88	0.27
256	8	2.08	0.26
512	9	2.30	0.26
1024	10	2.51	0.25
2048	11	2.71	0.25
4096	12	2.96	0.25
8192	13	3.23	0.25
16384	14	3.46	0.25

that hold for an entire class  $C(n)$  of problems, parametrized by the size  $n$  of the data. What is surprising, with hindsight, is that this ambitious sledge-hammer approach almost always works for problems in  $P$ ! We generally find a *single* algorithm that works well for all problems in  $C$ , from small to large, whose complexity is accurately described by a simple formula for all values of  $n$  of practical interest.

If we aim at a result of equal generality for some NP-hard problem class  $C(n)$ , the outcome is disappointing from a practical point of view, for two reasons:

1. Since the class  $C(n)$  undoubtedly contains many hard problem instances, a bound that covers all instances will necessarily be too high for the relatively harmless instances. And even though the harmless instances might be a minority within the class  $C(n)$ , they may be more representative of the actual problems a user may want to solve.
2. It is convenient, but not mandatory, to be able to use a single algorithm for an entire class  $C(n)$ . But when this approach fails in practice we have another option which is more thought-intensive but hopefully less compute-intensive: to analyze the specific problem instance we want to solve, and to tailor the algorithm to take advantage of the characteristics of this instance.

In summary, the standard complexity theory for NP-hard problems asserts the non-existence of algorithms that are efficient across an entire problem class, but ignores the possibility that many instances, perhaps including those of interest, can be solved efficiently.

An interesting and valuable approach to bypass the problem above is to design algorithms that efficiently compute approximate solutions to NP-hard problems. The practical justification for this approach is that an exact or optimal

solution is often not required, provided an error bound is known. The limitation of approximation algorithms is due to the same cause as for exact algorithms for NP-hard problems: that they aim to apply to an entire class  $C(n)$ , and thus cannot take advantage of the features of specific instances. As a consequence, if we insist on a given error bound, say 20%, the approximation problem often remains NP-hard.

There is a different approach to NP-hard problems that still insists on finding exact or optimal solutions. We do not tackle an entire problem class  $C(n)$ ; instead, we attack a challenging problem instance, taking advantage of all the specific features of this individual instance. If later we become interested in another instance of the same class  $C$ , the approach that worked for the first instance will have to be reappraised and perhaps modified. This approach changes the rules of algorithm design and analysis drastically: we still have to devise and implement clever algorithms, but complexity is not measured asymptotically in terms of  $n$ : it is measured by actually counting operations, disk accesses, and seconds.

## 4 Exhaustive Search and Enumeration: Concepts and Terminology

One of the oldest approaches to problem solving with the help of computers is brute-force enumeration and search: generate and inspect all data configurations in a large state space that is guaranteed to contain the desired solutions, and you are bound to succeed — if you can wait long enough. Although exhaustive search is conceptually simple and often effective, such an approach to problem solving is sometimes considered inelegant. This may be a legacy of the fact that computer science concentrated for decades on fine-tuning highly efficient polynomial-time algorithms. The latter solve problems that, by definition, are said to be in  $P$ . The well known class of NP-hard problems is widely believed to be intractable, and conventional wisdom holds that one should not search for exact solutions, but rather for good approximations. But the identification of NP-hard problems as *intractable* is being undermined by recent empirical investigation of extremely compute-intensive problems. The venerable *traveling salesman problem* is just one example of a probably worst-case hard problem class where many sizable instances turn out to be surprisingly tractable.

The continuing increase in computing power and memory sizes has revived interest in brute-force techniques for a good reason. The universe of problems that can be solved by computation is messy, not orderly, and does not yield, by and large, to the elegant, highly efficient type of algorithms that have received the lion's share of attention of the algorithms research community. The paradigm shift that may be changing the focus of computational research is that *combinatorial chaos* is just as interesting and rewarding as well-structured problems. Many “real” problems exhibit no regular structures to be exploited, and that leaves exhaustive enumeration as the only approach in sight. And even though we look in vain for order of magnitude asymptotic improvements due to “optimal”

algorithms, there are order of magnitude improvements waiting to be discovered due to program optimization, and to the clever use of limited computational resources. It is a game of algorithm design and analysis played according to a new set of rules: forget asymptotics and see what can be done for some specific problem instance, characterized by some large value of  $n$ . The main weapon in attacking instances of NP-hard problems, with an invariably irregular structure, is always *search and enumeration*.

The basic concepts of search algorithms are well known, but the terminology used varies. The following recapitulation of important concepts serves to introduce our terminology.

**State space  $S$ .** Discrete, often (but not necessarily) finite. Modeled as a graph,  $S = (V, E)$  where  $V$  is the set of vertices or nodes,  $E$  the set of edges (or perhaps arcs, i. e. directed edges). Nodes represent states, arcs represent relationships defined by given operators.

One or more **operators**,  $o: S \rightarrow 2^S$ , the powerset of  $S$ . An operator transforms a state  $s$  into a number of neighboring states that can easily be computed given  $s$ . In the frequently occurring *symmetric case*, an operator  $o$  is *its own inverse* in the following sense:  $s' \in o(s)$  implies  $s \in o(s')$ .

**Distinguished states**, e. g. starting state(s), goal or target state(s). The latter are usually defined by some target predicate  $t: S \rightarrow \{true, false\}$ .

**Objective function**, cost function  $f: S \rightarrow Reals$ . Serves to define an optimization problem, where one asks for any or all states  $s$  that optimize (minimize or maximize)  $f(s)$ .

**Search space.** Often used as a synonym for state space. Occasionally it is useful to make a distinction: a state space defines the problem to be solved, whereas different search algorithms may traverse different subsets of  $S$  as their search space.

**Traversal.** Sequentialization of the states of  $S$ . Common traversals are based on imposing tree structures over  $S$ , called search tree(s) or search forest.

**Search tree.** A rooted, ordered tree superimposed on  $S$ . The children  $s_1 \dots s_f$  of a node  $s$  are obtained by applying to  $s$  one of the operators defined on  $S$ , and they are ordered (*left-to-right order* when drawn). The number  $f$  of children of a node is called its *fan-out*. Nodes without children are called *leaves*.

**Search DAG.** It is commonly the case that the same state  $s$  will be encountered along many different paths from the root of a search tree towards its leaves. Thus, the same state  $s$  may generate many distinct nodes of the same search tree. By identifying (merging) all the tree nodes corresponding to the same state  $s$  we obtain a directed acyclic graph, the search DAG.

**DFS, BFS, etc.** The most common traversal of  $S$  w. r. t. a given search tree over  $S$  is *depth-first search* (DFS) or *backtrack*. DFS maintains at all times a single path from the root to the current node, and extends this path whenever possible. *Breadth-first search* (BFS) is also common, and can be likened to wave-propagation outwards from a starting state. BFS maintains at all times a frontier, or wave-front, that separates the states already visited from

those yet to be encountered. Several other traversals are useful in specific instances, such as best-first search, or iterative deepening.

**Search structures.** Every traversal requires data structures that record the current state of the search, i.e. the subspace of  $S$  already visited. DFS requires a stack, BFS a queue. The entire state  $S$  may require a mark (e.g. a bit) for each state to distinguish states already visited from those not yet encountered. The size of these data structures is often a limiting factor that determines whether or not a space  $S$  can be enumerated with the memory resources available.

**Enumeration.** A sequential listing of all the states of  $S$ , or of a subset  $S|t$  consisting of all the target states  $s$  for which  $t(s) = true$ .

**Output-sensitive enumeration algorithm.** It is often difficult to estimate a priori the size of the output of an enumeration. An appropriate measure of the time required by an enumeration is therefore output-sensitive, whereby one measures the time required to produce and output one state, the next in the output sequence.

**Search.** The task of finding one, or some, but *not necessarily all*, states  $s$  that satisfy some constraint, such as  $t(s) = true$  or  $f(s)$  is optimal.

**Exhaustive search.** A search that is guaranteed to find all states  $s$  that satisfy given constraints. An exhaustive search need not necessarily visit all of  $S$  in every instance. It may omit a subspace  $S'$  of  $S$  on the basis of a mathematical argument that guarantees that  $S'$  cannot contain any solution. In a worst case configuration, however, exhaustive search is forced to visit all states of  $S$ .

**Examples of exhaustive search.** Searching for a key  $x$  in a hash table is an exhaustive search. Finding  $x$ , or determining that  $x$  is not in the table, is normally achieved with just a few probes. In the worst case, however, collisions may require probing the entire table to determine the status of a key. By contrast, binary search is not exhaustive; when the table contains 3 or more keys, binary search will never probe all of them. Common exhaustive search algorithms include backtrack, branch-and-bound, and sieves, such as Erathostenes' prime number sieve.

## 5 Reverse Search

The more information is known a priori about a graph, the less book-keeping data needs to be kept during the traversal. Avis and Fukuda [2] present a set of conditions that enable graph traversal without auxiliary data structures such as stacks, queues, or node marks. The amount of memory used for book-keeping is constant, i.e. independent of the size of the graph. Their *reverse search* is a depth-first search (DFS) that requires neither stack nor node markers to be stored explicitly — all necessary information can be recomputed on the fly. Problems to which reverse search applies allow the enumeration of finite sets much larger than would be possible if a stack and/or markers had to be maintained. Such enumeration is naturally time-consuming. But computing time is an elastic resource—you can always wait “a little bit longer” — whereas memory is



inelastic. When it is full, a stack or some other data structure will overflow and stop the search. Thus, exhaustive search is often memory-bound rather than time-bound.

Three conditions enable reverse search to enumerate a state space  $S = (V, E)$ :

1. There is an *adjacency operator* or “*oracle*”  $A: S \rightarrow 2^S$ , the powerset of  $S$ .  $A$  assigns to any state  $s$  an ordered set  $A(s) = [s_1, \dots, s_k]$  of its neighbors. Adjacency need not be symmetric, i. e.  $s' \in A(s)$  does not imply  $s \in A(s')$ . The pairs  $(s, s')$  with  $s' \in A(s)$  define the set  $E$  of directed edges of  $S$ .
2. There is a *gradient function*  $g: S \rightarrow S \cup \{\text{nil}\}$ , where *nil* is a fictitious state (a symbol) not in  $S$ . A state  $s$  with  $g(s) = \text{nil}$  is called a *sink* of  $g$ .  $g$  assigns to any state  $s$  a unique successor  $g(s) \in S \cup \{\text{nil}\}$  subject to the following conditions:
  - for any state  $s$  that is not a sink, i. e.  $g(s) \neq \text{nil}$ , the pair  $(g(s), s) \in E$ , i. e.  $s \in A(g(s))$ ,
  - $g$  defines no cycles, i. e.  $g(g(\dots g(s)\dots)) = s$  is impossible—hence the name *gradient*.

Notice that when  $A$  is not symmetric,  $g$ -trajectories point in the opposite direction of the arcs of  $E$ . The *no cycles* condition in a finite space  $S$  implies that  $g$  superimposes a forest, i. e. a set of disjoint trees, on  $S$ , where tree edges are a subset of  $E$ . Each sink is the root of such a tree.

3. It is possible to efficiently *enumerate all the sinks* of  $g$  before exploring all of  $S$ .

The motivation behind these definitions and assumptions lies in the fact that  $A$  and  $g$  together provide all the information necessary to manage a DFS that starts at any sink of  $g$ . The DFS tree is defined by  $A$  and  $g$  as follows: The children  $C(s) = [c_1, \dots, c_f]$  of any node  $s$  are those nodes  $s'$  culled from the set  $A(s) = [s_1, \dots, s_k]$  for which  $g(s') = s$ . And the order  $[c_1, \dots, c_f]$  of the children of  $s$  is inherited from the order defined on  $A(s)$ .

A DFS usually relies on a stack for walking up and down a tree. An explicit stack is no longer necessary when we can call on  $A$  and on  $g$ . Walking up the DFS tree towards the root is accomplished simply by following the gradient function  $g$ . Walking down from the root is more costly. Calling the adjacency oracle from any node  $s$  yields a superset  $A(s)$  of the children of  $s$ . Each  $s'$  in  $A(s)$  must then be checked to see whether it is a child of  $s$ , as determined by  $g(s') = s$ .

Similarly, no data structure is required that marks nodes already visited. The latter can always be deduced from the order defined on the set  $C(s)$  of children and from two node identifiers: the current state and its immediate predecessor in the DFS traversal.

We explain how reverse search works on a simple example where every step can be checked visually. Fig. 1 shows a hexagon (drawn as a circle) with vertices labeled 1 . . . 6. Together with all 9 interior edges it makes up the complete graph  $K_6$  of 6 vertices shown at the top left. The other 14 copies of the hexagon, each of them with 3 interior edges, show all the distinct triangulations of this

labeled hexagon. The double-tipped arrows link each pair of states that are neighbors under the only operator that we need to consider in this problem: a *diagonal flip* transforms one triangulation into a neighboring one by exchanging one edge for another. This, and the term *diagonal*, will be explained after we introduce the notation for identifying edges and triangulations.

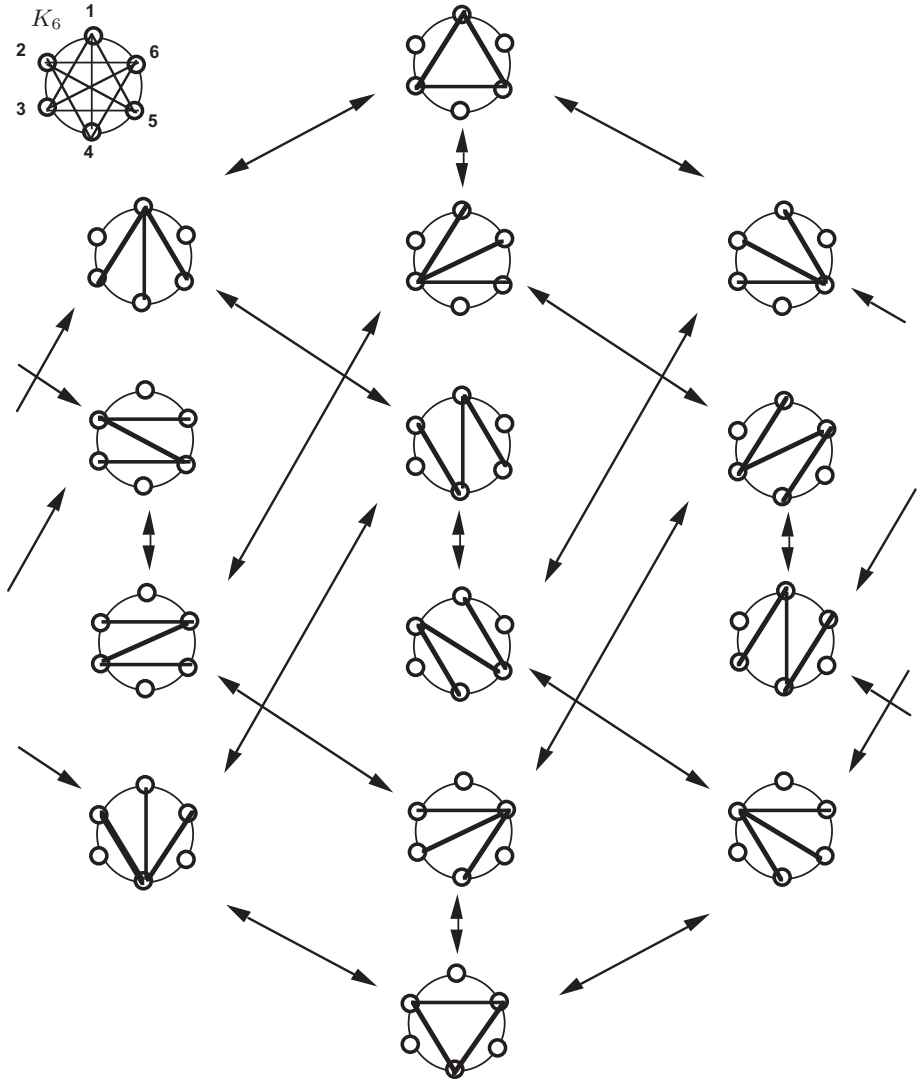


Fig. 1. The state space  $S$  of triangulations of a labeled hexagon

The second ingredient required to implement reverse search is a suitable gradient function  $g$ . We obtain this by imposing an arbitrary total order on the state space  $S$ , accepting the fact that it breaks the elegant symmetry of Fig. 1. Label an edge  $(i, j)$  with the ordered digit pair  $ij$ ,  $i < j$ . We label the 3 interior edges  $x, y, z$  of a triangulation with the triple  $x.y.z$  of edge labels (digit pairs) ordered as  $x < y < z$ . This labeling scheme assigns to each triangulation of the labeled hexagon a unique identifier  $x.y.z$ . As shown in Fig. 2, we order the 14 triangulations lexicographically. When  $x.y.z$  is interpreted as an integer, lexicographic and numerical order coincide.

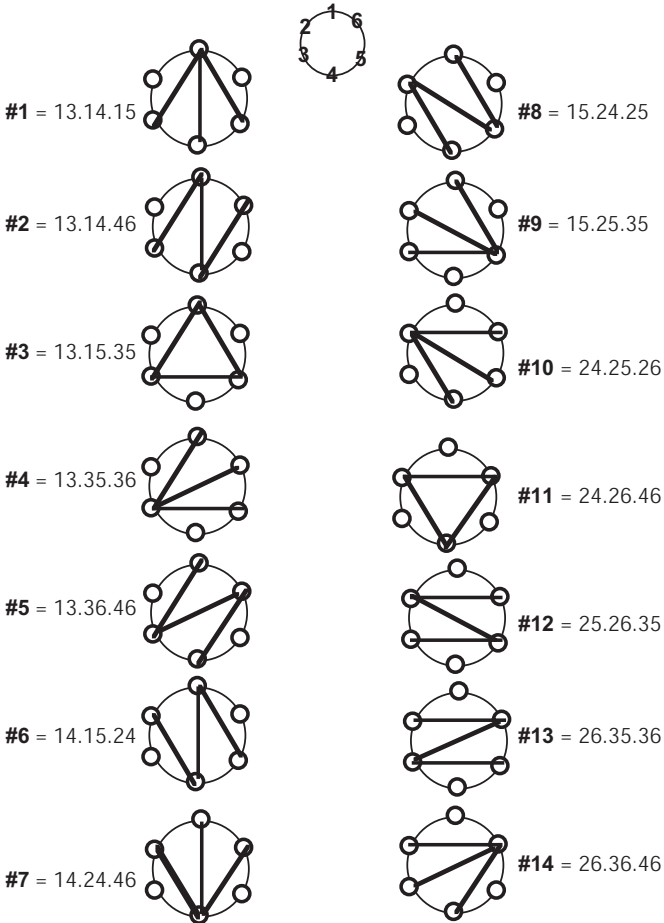


Fig. 2. The state space  $S$  sequenced by lexicographic order

To understand the operator *diagonal flip*, observe that in every triangulation, each interior edge is the *diagonal* of a quadrilateral, i. e. a cycle of length 4. In triangulation #1 = 13.14.15, for example, 13 is the diagonal of the quadrilateral with vertices 1, 2, 3, 4. By flipping diagonals in this quadrilateral, i. e. replacing 13 by its diagonal mate 24, we obtain triangulation #6 = 14.15.24. Thus, each of the 14 triangulations has exactly 3 neighbors under the operator *diagonal flip*. The 14 triangulations as vertices and the  $14 * 3/2 = 21$  neighbor relations as edges define the state space  $S = (V, E)$  of this enumeration problem.

Based on the total order shown in Fig. 2, define the gradient function  $g$  as follows:  $g(s)$  is the “smallest” neighbor  $s'$  of  $s$  in lexicographic order, provided  $s' < s$ .  $g$  is defined on all states of  $s$  except on triangulation #1 = 13.14.15, which has no smaller neighbor. In order to handle this case we define  $g(\#1) = s_0$ , where  $s_0$  is the “sentinel” introduced in the definition  $g: S \rightarrow S \cup \{s_0\}$ . This gradient function  $g$  defines the search tree shown in Fig. 3.

The third ingredient required by reverse search, an efficient way to construct all the sinks of  $g$ , is trivial in this example. The triangulation 13.14.15 contains the three *smallest* interior edges; it is therefore the smallest triangulation and the sink of all  $g$ -trajectories. We construct triangulation #1 and start a DFS traversal at this root of the search tree.

Consider the typical case when DFS arrives at node #4. The adjacency oracle returns #4’s three neighbors: #3, #5 and #13. The gradient function applied to these three neighbors identifies #13 as #4’s only child. DFS must now decide to either descend to its child #13 or to backtrack to its parent #3 =  $g(\#4)$ . How can we tell?

In addition to the current node #4, DFS retains the identifier of the immediate predecessor node in the traversal. If the predecessor is #3, then DFS is on its way down and proceeds to visit child #13. If, on the other hand, the predecessor is #13, then DFS is on its way back up and proceeds to re-visit its parent #3. A similar logic lets DFS take its next step in every case. Consider the case when DFS is currently at the root, vertex #1. If the predecessor is *nil*, i. e. our fictitious sentinel  $s_0$ , DFS knows that it is just starting its traversal. If the predecessor is #3, DFS has to visit its next child, in order, i. e. #6. If the predecessor is #6, i. e. the last child of the root, DFS is done.

## 6 Best Euclidean Spanning Trees: A Case Study in Geometry, Combinatorics and Enumerative Optimization

Problems about discrete spatial configurations involve a challenging interaction between the continuum characteristic of geometry and the discreteness of combinatorics. Each of these disciplines has evolved its characteristic techniques that exploit the nature of the objects treated, continuous or discrete. When used together to attack a geometric-combinatorial problem, surprises lurk beneath the surface. A seemingly minor change in a geometric specification may clash

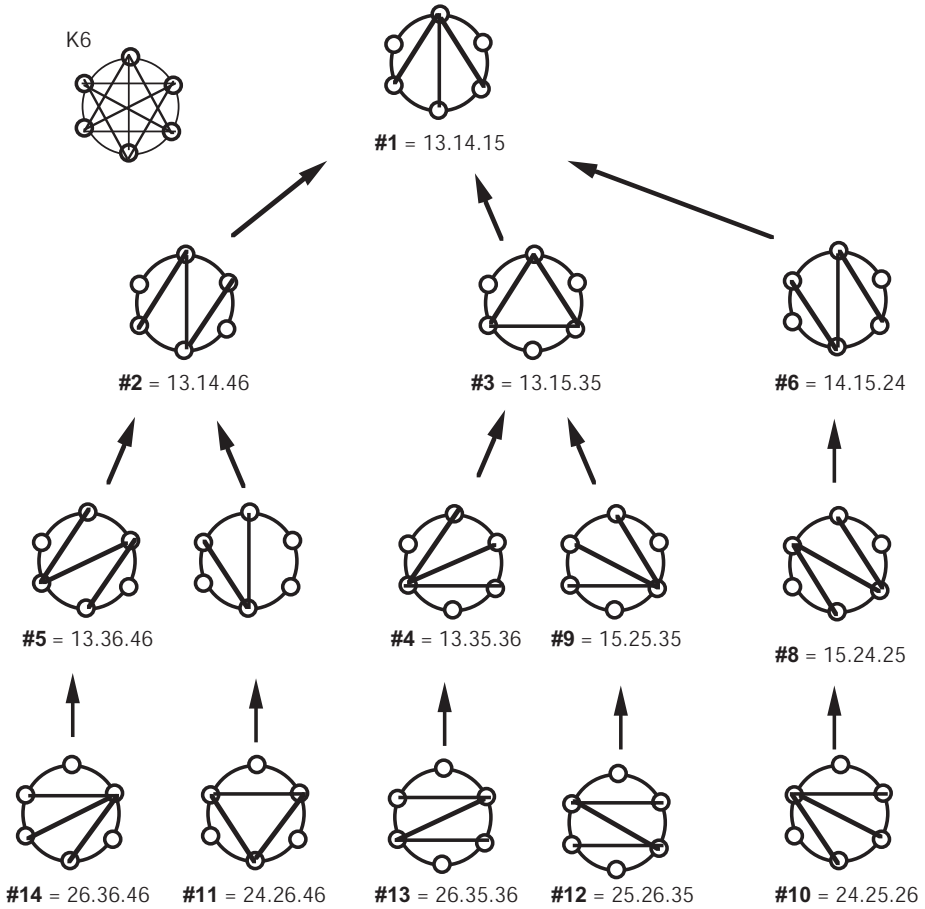


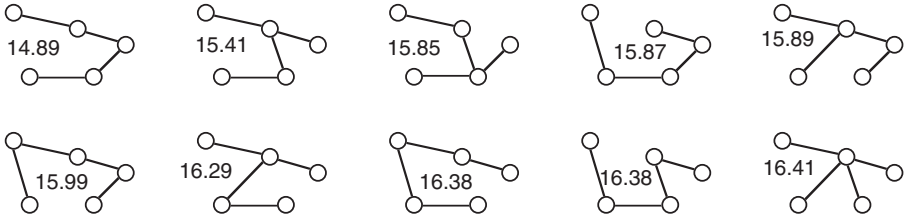
Fig. 3. Search tree defined by the gradient function  $g$

with combinatorial assumptions, and vice versa. This interaction between geometric and combinatorial constraints makes it difficult to predict, on an intuitive basis, which problems are amenable to efficient algorithms, and which are not.

The difficulties mentioned are exacerbated when we aim at enumerating all spatial configurations that meet certain specifications, not in any arbitrary order convenient for enumeration, but rather in a prescribed order. Search techniques impose their own restrictions on the order in which they traverse a search space, and these may be incompatible with the desired order.

We present an example of a simple geometric-combinatorial search and enumeration problem as an illustration of issues and techniques: the enumeration of plane spanning trees over a given set of points in the plane, i. e. those trees constructed with straight line segments in such a manner that no two edges

intersect [8]. [2] present an algorithm for enumerating all plane spanning trees, in some uncontrolled order that results from the arbitrary labeling of points. We attack this same problem under the additional constraint that these plane spanning trees are to be enumerated according to their total length (i. e. the sum of the lengths of their edges), from short to long. We may stop the enumeration after having listed the  $k$  shortest trees, or all trees shorter than a given bound  $c$ . Fig. 4 lists the 10 shortest plane spanning trees among the 55 on the particular configuration of 5 points with coordinates  $(0, 5), (1, 0), (4, 4), (5, 0), (7, 3)$ .



**Fig. 4.** The 10 shortest plane spanning trees over five given points, in order of increasing length

In trying to apply reverse search to a “ $k$  best problem” we face the difficulty that the goal is not just to enumerate a set in some arbitrary, convenient order, as we did in Section 5. In the case of enumerative optimization the goal is to enumerate the elements of  $S$  from best to worst according to some objective function  $f$  defined on  $S$ . One may wish to stop the enumeration after the  $k$  best elements, or after having seen all elements  $s$  with  $f(s) \leq c$ .

### 6.1 The Space of Plane Spanning Trees on a Euclidean Graph

Consider a graph  $G = (V, E, w)$  with  $n$  vertices  $p, q, r, \dots$  in  $V$ , weighted edges  $e = (p, q)$  in  $E$ , and a weight function  $w: E \rightarrow Reals$ . The set of spanning trees over  $G$  has a well-known useful structure that is exploited by several algorithms, in particular for constructing a minimum spanning tree (MST). The structure is based on an exchange operator, an *edge flip*, and on a monotonicity property.

Let  $T$  be any spanning tree over  $G$ ,  $e'$  an edge not in  $T$ ,  $Ckt(e', T)$  the unique path  $P$  in  $T$  that connects the two endpoints of  $e'$ , and  $e$  any edge in  $P$ . The edge flip  $T' = T - e + e'$  that deletes  $e$  from  $T$  and replaces it by  $e'$  creates a new spanning tree  $T'$  that is *adjacent to*  $T$ . If  $w(e) > w(e')$  this edge flip is profitable in the sense that  $|T'| < |T|$ , where  $|T|$  denotes the total length of  $T$ . The remarkable fact exploited by algorithms for constructing a minimum spanning tree (MST) is that in the space of all spanning trees over  $G$ , any local minimum is also a global minimum. This implies that any greedy algorithm based on profitable edge flips or on accumulating the cheapest edges (e. g. Prim, Kruskal) converges towards an MST.

In this paper we study *Euclidean* graphs and *plane* spanning trees. A Euclidean graph is a complete graph whose vertices are points in the plane, and whose edge weights are the distances between the endpoints of the edge. For  $p, q \in V$  and  $e = (p, q)$ , let  $|e|$  denote the length of edge  $e$ . For a Euclidean graph it is natural to consider “plane” or non-crossing spanning trees, i. e. trees no two of whose edges cross. It is well known and follows directly from the triangle inequality that any MST over a Euclidean graph is plane, i. e. has no crossing edges.

In the following section, we define a search tree for enumerating all plane spanning trees, in order of increasing total length, over a given point set in the plane. This search tree is presented in such a form that standard tree traversal techniques apply, in particular reverse search [2]. Specifically, we define a unique root  $R$  which is an MST; and a monotonic gradient function  $g$  that assigns to each tree  $T \neq R$  a tree  $g(T)$  with  $|g(T)| \leq |T|$ . The gradient function  $g$  has the property that, for any  $T \neq R$ , some iterate  $g(\dots g(T))$  equals  $R$ , i. e.  $R$  is a sink of  $g$ ; hence  $g$  generates no cycles. For efficiency’s sake, both  $g$  and its inverse can be computed efficiently.

For simplicity of expression we describe the geometric properties of the search tree as if the configuration of points was non-degenerate in the sense that there is a unique MST, and that any two distinct quantities (lengths, angles) ever compared are unequal.

Unfortunately, the space of *plane* spanning trees over a Euclidean graph does not exhibit as simple a structure as the space of *all* spanning trees. It is evident that if  $T$  is a plane tree, an edge flip  $T' = T - e + e'$  may introduce cross-overs. Thus, the geometric key issue to be solved is an efficient way of finding edge flips that shorten the tree and avoid cross-over. We distinguish two cases:

**6.1.1 Flipping edges in the Gabriel Graph.** Consider any set  $C$  of non-crossing edges over the given point set  $V$ , and any plane tree  $T$  contained in  $C$ . Trivially, any edge flip limited to edges in  $C$  cannot introduce any cross-over. We seek a set  $C$ , a *skeleton of  $G$* , that is dense enough so as to contain a sufficient number of spanning trees, and has useful geometric properties. Among various possibilities, the Gabriel Graph of  $V$  will do.

**Definition 1.** The Gabriel Graph  $GG(V)$  over a point set  $V$  contains an edge  $(p, q)$  iff no point  $r$  in  $V - \{p, q\}$  is in the closed disk  $Disk(p, q)$  over the diameter  $(p, q)$ .

The useful geometric properties mentioned include:

1.  $GG(V)$  has no crossing edges.
2. Consider any point  $x$  (not in  $V$ ) that lies inside  $Disk(p, q)$ . Then  $|x, p| < |p, q|$ ,  $|x, q| < |p, q|$ ,  $\angle(p, x, q) > 90^\circ$ .
3. Any MST over  $V$  is contained in  $GG(V)$ .

(*Proof:* consider any edge  $e$  of an MST. If there was any point  $p \in V$  inside  $Disk(e)$ ,  $e$  could be exchanged for a shorter edge.)

These geometric properties lead to a first rule.

**Rule 1.** Let  $T$  be a spanning tree over  $V$  that is not the (uniquely defined) MST  $R$ . If  $T$  is contained in  $GG(V)$ , let  $g(T) = T - e + e'$ , where  $e'$  is the lexicographically first edge of  $R$  not in  $T$ , and  $e$  is the longest edge in  $Ckt(e', T)$ .

Obviously,  $g(T)$  is closer to  $R$  than  $T$  is, and if the MST is unique, then  $|g(T)| < |T|$ .

**6.1.2 Flipping edges not in the Gabriel Graph.** As a planar graph, the Gabriel Graph  $GG(V)$  has a sparse set of edges. Thus, the vast majority of spanning trees over  $V$  are not contained in  $GG(V)$ , and hence Rule 1 applies mostly towards the end of a  $g$ -trajectory, for spanning trees near the MST  $R$ . For all other spanning trees, we need a rule to flip an edge  $(p, r)$  not in  $GG(V)$  in such a way that the spanning tree gets shorter, and no cross-over is introduced.

Consider a tree  $T$  not contained in  $GG(V)$ , and hence is not an MST. Among all point triples  $(p, q, r)$  such that  $(p, r)$  is in  $T$ , select the one whose  $\angle(p, q, r)$  is maximum. The properties of the Gabriel Graph imply the following assertions:  $\angle(p, q, r) > 90^\circ$ ,  $(p, r)$  is not in  $GG(V)$ ,  $|(p, q)| < |(p, r)|$  and  $|(q, r)| < |(p, r)|$ .

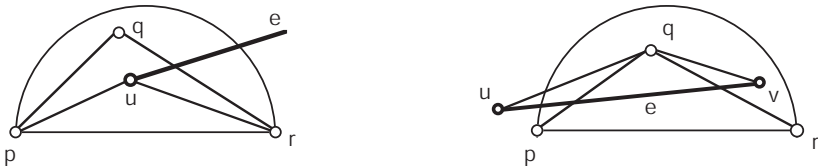
**Rule 2.** With the notation above, let  $g(T) = T - (p, r) + e'$ , where  $e'$  is either  $(p, q)$  or  $(q, r)$ , chosen such that  $g(T)$  is a spanning tree.

As mentioned,  $|g(T)| < |T|$ . Fig. 5 illustrates the argument that this edge flip does not introduce any crossing edges. At left, consider the possibility of an edge  $e$  one of whose endpoints  $u$  lies in the triangle  $(p, q, r)$ . This contradicts the assumption that  $\angle(p, q, r)$  is maximum. At right, consider the possibility of an edge  $e = (u, v)$  that crosses both  $(p, q)$  and  $(q, r)$ . Then  $\angle(u, q, v) > \angle(p, q, r)$ , again a contradiction. Thus, neither  $(p, q)$  nor  $(q, r)$  cause a cross-over if flipped for  $(p, r)$ .

These two rules achieve the goal of finding edge flips that

- shorten the tree, and
- avoid cross-over.

Thus, after a problem-specific detour into geometry, we are back at the point where a general-purpose tool such as reverse search can take over.



**Fig. 5.** The assumption of new cross-overs contradicts the choice of “angle(p,q,r) is maximum”



## 7 The Craft of Attacking Hard Problem Instances

Exhaustive search is truly a creation of the computer era. Although the history of mathematics records amazing feats of paper-and-pencil computation, as a human activity, exhaustive search is boring, error-prone, exhausting, and never gets very far anyway. As a cautionary note, if any is needed, Ludolph van Ceulen died of exhaustion in 1610 after using regular polygons of  $2^{62}$  sides to obtain 35 decimal digits of  $p$ —they are engraved on his tombstone.

With the advent of computers, *experimental mathematics* became practical: the systematic search for specific instances of mathematical objects with desired properties, perhaps to disprove a conjecture or to formulate new conjectures based on empirical observation. Number theory provides a fertile ground for the team *computation + conjecture*, and Derrick Lehmer was a pioneer in using search algorithms such as sieves or backtrack in pursuit of theorems whose proof requires a massive amount of computation [6]. We make no attempt to survey the many results obtained thanks to computer-based mathematics, but merely recall a few as entry points into the pertinent literature:

- the continuing race for large primes, for example Mersenne primes of form  $2^p - 1$ ,
- the landmark proof of the “four-color theorem” by Appel and Haken [1],
- more recent work in Ramsey theory or cellular-automata [5].

For such cases we need a complexity measure that applies to problem instances, rather than to over-sized problem classes. Counting individual operations and measuring the running time of numerous procedures is a laborious exercise. Again we marvel at the surprising practical effectiveness of the asymptotic complexity analysis of algorithms—nothing of comparable elegance is in sight when we attack hard problem instances.

The algorithm designer faces different challenges when attacking an instance as compared to inventing an algorithm that solves a problem class. For the second case we have many paradigms such as divide-and-conquer, greedy algorithms, or randomization. The designer’s problem is to discover and prove mathematical properties that greatly reduce the number of operations as compared to a brute-force approach. When attacking a problem instance we expect a priori that there is nothing much cleverer than brute-force, and that we will use one of half a dozen general purpose search algorithms. The main difficulty is algorithm and program optimization.

Unfortunately, the discipline of algorithm and program optimization so far has resisted most efforts at systematization. Several recent Ph.D. thesis’ have attempted to extract general rules of how to attack compute-intensive problem instances from massive case studies [4,3,7]. Data allocation on disk is a central issue, trying to achieve some locality of data access despite the combinatorial chaos typical of such problems. In problems involving retrograde analysis (e.g. [11,13]), where every state (e.g. a board position in a game) in the state space is assigned a unique index in a huge array, construction of a suitable index function is critical. Since such computations may run for months and generate

data bases of many GigaBytes, independent verification of the result is a necessity. Some of the experience gained is summarized in [10].

Attacking computationally hard problem instances has so far never been near the center of algorithm research. It has rather been relegated to the niche of puzzles and games, pursued by a relatively small community of researchers. The experience gained is more a collection of individual insights rather than a structured domain of knowledge. As computing power keeps growing and people attack harder problems, we will often encounter problems that must be tackled as individual instances, because no algorithm that applies to a large class will be feasible. Thus, it is a challenge to the computing research community to turn individual insights into a body of knowledge, and to develop a complexity theory of problem instances.

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