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Publication date: 2014

Document Version
Early version, also known as pre-print

Citation for published version (APA):
Faster Exact Algorithms for Computing Steiner Trees in Higher Dimensional Euclidean Spaces

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Abstract. The Euclidean Steiner tree problem asks for a network of minimum total length interconnecting a finite set of points in \(d\)-dimensional space. For \(d \geq 3\), only one practical algorithmic approach exists for this problem — proposed by Smith in 1992. A number of refinements of Smith’s algorithm have increased the range of solvable problems a little, but it is still infeasible to solve problem instances with more than around 17 terminals. In this paper we firstly propose some additional improvements to Smith’s algorithm. Secondly, we propose a new algorithmic paradigm called branch enumeration. Our experiments show that branch enumeration has similar performance as an optimized version of Smith’s algorithm; furthermore, we argue that branch enumeration has the potential to push the boundary of solvable problems further.

Keywords: Steiner tree problem, \(d\)-dimensional Euclidean space, exact algorithm, computational study

1 Introduction

Given a finite set of \(n\) points \(N\) in a \(d\)-dimensional space \(\mathbb{R}^d\), \((d \geq 2)\), the \textit{Euclidean Steiner tree problem} (ESTP) asks for a network \(T^*\) of minimum length interconnecting \(N\). The vertices of \(T^*\) corresponding to points of \(N\) are called terminals while possible additional vertices in \(T^*\) are called Steiner points. \(T^*\) necessarily is a tree, and is referred to as a \textit{Euclidean minimum Steiner tree} (EMST) for \(N\). The ESTP in the plane \((d = 2)\) has a history that goes back more than two centuries [4], and is known to be NP-hard [9], even when the terminals are restricted to lie on two parallel lines [16]. The generalisation to more than two dimensions was introduced by Bopp [3] in 1879; more recent mathematical treatments can be found in [10, 13]. The problem in \(\mathbb{R}^d\), \(d \geq 3\), has applications to areas such as phylogenetics [6, 7, 5] and to the structure and folding of proteins [18, 20].

An EMST \(T^*\) can be viewed as a union of full EMSTs whose terminals have degree 1. Steiner points in \(T^*\) (in any dimension) have degree 3. The three
incident edges at a Steiner point are co-planar and each pair meets at an angle of 120°. These degree conditions imply that full EMSTs spanning \(k\) terminals, \(1 \leq k \leq n\), have \(k - 2\) Steiner points.

The first exact algorithms for the ESTP in the plane (\(\mathbb{R}^2\)) were based on the following common framework [12]. Subsets of terminals are considered one by one. For each subset, all its full EMSTs are determined and the shortest is retained. Several tests are applied to these retained full EMSTs to decide if they can belong to an EMST for \(N\). Surviving full EMSTs are then concatenated in all possible ways to obtain trees spanning \(N\). The shortest of them is \(T^*\).

The main bottleneck in this approach is the generation of full EMSTs. It has been observed [24] that substantial improvements can be obtained if full RMSTs are generated simultaneously across various subsets of terminals. Very powerful geometrical pruning tests identifying non-optimal full EMSTs can then be applied not after but during their generation. As a consequence of this speed-up, the concatenation of full EMSTs became a bottleneck of the EMST computation. A remedy was based on the observation that the concatenation of full EMSTs can be formulated as a problem of finding a minimum spanning tree in a hypergraph with terminals as vertices and subsets spanned by full EMSTs as hyperedges [21]. In practice, this problem can be solved efficiently using branch-and-cut techniques. The dramatic improvements of both the generation and concatenation of full EMSTs led to the development of GeoSteiner [22, 23, 25] which can routinely solve problem instances with thousands of terminals in \(\mathbb{R}^2\) in a reasonable amount of time. A similar methodology has been applied to other metrics and generalisations [11, 15, 26, 27].

Significantly less improvement has been made on exact algorithms for the problem in \(\mathbb{R}^d\), \(d \geq 3\). Currently, only one practical algorithmic approach exists. It was proposed by Smith [19] in 1992. A couple of recent contributions [8, 14], all based on Smith’s algorithm, have pushed the boundary of solvable problems a little, but in practice it is still infeasible to solve problems for \(d \geq 3\) with more than 17 terminals.

As observed by Fampa and Anstreicher [8], the bounds used in [19] do not correspond to rigorous lower bounds on the solution values of these problems, but are instead obtained from putatively near-optimal solutions. In addition, when a node fails to fathom the algorithm has no means to estimate the objective values associated with its potential children. As a result the terminal nodes are added in a fixed order, even though varying the order has the potential to substantially reduce the size of the branch-and-bound tree.

**Our Contribution.** In this paper we first describe and evaluate some improvements to Smith’s algorithm. Then we describe a novel exact algorithm, a so-called a branch enumeration approach. The branch enumeration algorithm deviates significantly from Smith’s algorithm. One of the advantages of the branch enumeration algorithm is that it is possible to apply stronger pruning tests early in the enumeration, e.g. based on the notion of bottleneck Steiner distances. We evaluate our new branch enumeration algorithm and our improved version of Smith’s algorithm experimentally on a set of benchmark instances.
GeoSteiner \cite{22, 23, 25} in the Euclidean plane can be viewed as a branch
enumeration (to obtain a superset of full Steiner trees in the optimal solution)
followed by the concatenation of these full Steiner trees (using branch and cut to
solve the minimum spanning tree problem in a hypergraph). Prunning techniques
in GeoSteiner are extremely powerful and make it possible to efficiently prune
nonoptimal full Steiner trees. A branch enumeration algorithm has also been
suggested to compute Steiner minimum trees in Hamming metric space \cite{1}. Our
approach to find Steiner minimum trees in high-dimensional Euclidean spaces is
similar. However, pruning techniques in the Hamming metric space seem to be
much stronger than in high-dimensional Euclidean spaces (but far from being as
strong as in GeoSteiner when applied to the problems in the Euclidean plane).

Organisation of the Paper. In Section 2 we present some preliminaries on
the ESTP. Smith’s algorithm is introduced in Section 3. Its improvements are
discussed in Section 4. Then in Section 5 we describe the branch enumeration
algorithm. Computational results are presented in Section 6, and concluding
remarks are given in Section 7.

2 Preliminaries

Consider a tree $T$ in $\mathbb{R}^d$ that interconnects a set $N$ of $n$ points. We assume that
$T$ consists of a set of vertices (a superset of $N$) and a set of edges (straight line
segments) connecting pairs of vertices. The length of $T$, denoted by $|T|$, is the
sum of its edge lengths. The length of an edge $(u,v)$ connecting points $u$ and $v$
is denoted by $|uv|$. The given points of $N$ are called the terminals, and the other
vertices, if any, are called Steiner points (see Fig. 1). The ESTP is to determine
the shortest tree $T^*$ for $N$. Steiner points in $T^*$ have degree 3 while terminals
have degree at most 3. Any tree satisfying these degree constraints is called a
Steiner tree for $N$.

![Fig. 1: The EMST of five terminals in $\mathbb{R}^2$. The terminals are shown as filled black
circles, and the Steiner points are shown as smaller red circles. The topology $T^*$
of $T^*$ is shown to the right. Terminal 4 is connected to a Steiner point by a
zero-length edge and therefore the Steiner point overlaps with terminal 4.](image)

A Steiner topology $T$ of a Steiner tree $T$ for $N$ is a specification of the inter-
connections in $T$, disregarding the positions of its vertices. Any Steiner topology
can be transformed into a full Steiner topology (FST) where all terminals have
degree 1. If $T$ has a terminal $t$ adjacent to two vertices $v_1$ and $v_2$, a new Steiner point $s$ connected to $t$, $v_1$ and $v_2$ is added. If $T$ has a terminal $t$ adjacent to three vertices $v_1$, $v_2$ and $v_3$, a pair of Steiner points $s_1$ and $s_2$ is added. Steiner point $s_1$ is connected to $s_2$, $v_1$ and $t$. Steiner point $s_2$ is connected to $s_1$, $v_2$ and $v_3$.

The shortest Steiner tree for a given topology, $T$, is called a relatively minimal tree (RMT) of $T$. It always exists and is unique [12]. Its Steiner points may overlap with terminals and with other Steiner points because of zero length edges. Clearly, $T^*$ is an RMT for its FST $T^*$. Unfortunately, $T^*$ is not known beforehand. Unless $P=NP$, the only feasible way to find it seems to be the enumeration of all FSTs, the determination of their RMTs and the selection of the shortest one as $T^*$.

The bottleneck edge between two terminals $t_i$ and $t_j$ of $N$ is the longest edge on the path from $t_i$ to $t_j$ in the minimum spanning tree of $N$. The length of the bottleneck edge between $t_i$ and $t_j$ is referred to as the bottleneck distance and is denoted by $\beta(t_i,t_j)$. Bottleneck distances can be determined in polynomial time in a preprocessing phase. It is well-known [12] that no edge on a path between a pair of terminals $t_i$ and $t_j$ in the EMST can be longer than $\beta(t_i,t_j)$.

Consider a FST $T$ for $N$. Let $s$ be any of its Steiner points. Let $r_i$, $r_j$ and $r_k$ denote its adjacent vertices. When $s$ is deleted, $T$ breaks into three rooted binary trees or branches $B_i$, $B_j$ and $B_k$ rooted at $r_i$, $r_j$ and $r_k$, respectively, see Fig. 2. Note that roots have degree 2 unless they are terminals. Furthermore the branches are disjoint in the sense that they do not share any terminals. The depth of a terminal in a branch is the number of edges separating it from the root of the branch.

Consider the FST $T_i$ obtained by splicing away a non-terminal root $r_i$ of $B_i$ and connecting its adjacent vertices with each other. For notational convenience, let $\text{RMT}(B_i)$ denote $\text{RMT}(T_i)$. Similar splicing away can be applied to non-terminal roots of $B_j$ and $B_k$.

A pair of disjoint branches $B_i$ and $B_j$ rooted at respectively $r_i$ and $r_j$ can be combined into a branch $B = B_i \oplus B_j$ by adding a new Steiner point $r$ as a root of $B$ adjacent to $r_i$ and $r_j$.

A triplet of disjoint branches $B_i$, $B_j$ and $B_k$ rooted at respectively $r_i$, $r_j$ and $r_k$ can be combined into a FST $T = B_i \oplus B_j \oplus B_k$ by adding a new Steiner point $r$ adjacent to $r_i$, $r_j$ and $r_k$.

A new algorithm based on the enumeration of branches rather then the enumeration of FSTs for $N$ will be described in Section 5. Root splicing will be used to prune branches that cannot be in the FST $T^*$ of the EMST $T^*$ of $N$. Disjoint triplets of the remaining feasible branches (whose union spans $N$) will generate feasible FSTs of $N$, including $T^*$.

3 Smith’s algorithm (Smith)

Smith’s algorithm [19] enumerates FSTs of $N$. RMTs are then determined for each such FST. The shortest RMT encountered is an EMST $T^*$ of $N$. The
Fig. 2: Left: A FST that has been split in 3 branches $B_i$, $B_j$, and $B_k$ by removing the Steiner point $s$. Right: The RMTs of each branch.

The enumeration of FSTs is achieved by the following expansion procedure. Terminals are assumed to be in some fixed order $t_1, t_2, \ldots, t_n$. Assume that a FST $T_k$ for $t_1, t_2, \ldots, t_k$, $3 \leq k < n$, is given. Note that only one FST exists for $k = 3$. Expand $T_k$ into $2k-3$ FSTs, each with $k+1$ terminals, by inserting a new Steiner point $s_{k-1}$ into every edge $e$ of $T_k$. In each expanded FST, $s_{k-1}$ is adjacent to $t_{k+1}$ and to the two vertices of $e$. This expansion process stops when $k = n$. It can be shown that every FST of $N$ corresponds to exactly one unique sequence of such expansions. As described, the FSTs are generated in a breadth-first fashion. However, in order to obtain RMTs for all $n$ terminals as quickly as possible, FSTs in [19] are generated in the best-first manner.

Given a FST $T$ of $N$, arbitrary initial positions are assigned to its $n-2$ Steiner points. These positions are then recomputed iteratively by solving a system of $d(n-2)$ equations with $d(n-2)$ unknowns (corresponding to the locations of Steiner points). It can be shown [19] that the length of the tree reduces with each iteration and converges to $RMT(T)$. The iterative procedure terminates if edges meet at Steiner points at angles within the interval $[2\pi/3 - \epsilon, 2\pi/3 + \epsilon]$ for an arbitrarily small constant $\epsilon > 0$. The reader is referred to [17] for the justification that a good approximation on the angles gives a good approximation of the length.

This iterative procedure can also be applied during the expansion process. Let $T_k$ be a FST spanning $k$ terminals, $3 \leq k < n$. Determine $RMT(T_k)$. Suppose that it is not shorter than the shortest RMT of all $n$ terminals encountered so far. The expansion of $T_k$ can be stopped since any such expansion can only increase the lengths of RMTs of expanded FSTs. This algorithm will be referred to as Smith.

Two improvements for Smith have previously been suggested. Fampa and Anstreicher [8] used a conic formulation for finding a lower bound on a particular topology that eliminates the need to explicitly compute RMTs of children in the branch-and-bound tree. This was used to tighten lower bounds and guide the search. In the result-section we denote this method Smith-Fampa. Laarhoven and Anstreicher suggested a series of geometric criteria based on Voronoi-regions, bottleneck distances and lune-properties that could be used to discard partial topologies. Additionally they suggested exploring terminals starting with the
one furthest from the point set centroid and going toward the center. In the result-section we denote this method Smith-Laarhoven.

4 Smith’s modified algorithm (Smith*)

Smith’s algorithm adds terminals in fixed but arbitrary order when expanding FSTs. If the low index terminals are close to each other, the corresponding FSTs will have short RMTs. As a consequence, the expansion procedure will rarely stop before \( k = n \). It has been suggested to index the terminals by their non-decreasing distance to their centroid [14]. We suggest a different distance-based indexing. Terminals \( t_1, t_2 \) and \( t_3 \) have to maximize the sum of their pairwise distances. The terminal \( t_k, k = 4, 5, \ldots, n \), is farthest away from \( t_1, t_2, \ldots, t_{k-1} \).

To obtain a reasonable upper bound (needed to stop the expansion process), the FST \( T_M \) of the minimum spanning tree of \( N \) is determined. This is achieved by introducing \( c-1 \) Steiner points at terminals of degree \( c, c \geq 1 \). If \( c \geq 3 \), there are several ways of interconnecting these \( c-1 \) Steiner points with each other and with the terminal. In the current implementation, an arbitrary interconnection pattern is chosen. The length of \( RMT(T_M) \) yields a good upper bound, denoted by \( UB \), on the length of \( T^* \). Given such good upper bound, the expansion of FSTs in the depth-first manner is no longer essential. Best-first expansion based on the lower bound together with the distance-based ordering of terminals results in the generation of fewer FSTs. This method is referred to as Smith*.

The quality of lower bounds could be improved by using the conic formulations from the Smith-Fampa-method [8]. While smaller number of FSTs would be generated, the computation time increases significantly. Geometric criteria based on the lune-properties combined with the bottleneck distances were also used to speed up the expansion process by eliminating non-optimal FSTs [14]. However, the extra time spent on geometric computations makes the improvements minimal when compared to the distance-based sorting of terminals. None of these improvements are therefore included in the Smith* algorithm.

5 Branch enumeration algorithm (Branch)

The approach of GeoSteiner [22] seems at first sight to be applicable to the ESTP in \( \mathbb{R}^d, d \geq 3 \). GeoSteiner has two phases. In the generation phase, full Steiner trees for all FSTs of all subsets of terminals are generated. Naturally, full Steiner trees that cannot be in \( T^* \) are pruned away. When FSTs are generated across different subsets, there are large groups of them that are very similar. The power of GeoSteiner rests partly in its ability to generate full Steiner trees with similar FSTs in a common pass and partly in its ability to prune away partially constructed FSTs. In the concatenation phase, GeoSteiner selects a subset of not pruned full Steiner trees that span all terminals and has the minimum total length. This problem can be formulated as the minimum spanning tree problem in a hypergraph. While this problem is NP-hard, a branch-and-cut algorithm
seems to be quite efficient to solve problem instances involving full Steiner trees not pruned away during the generation phase.

Unfortunately, this is not the case. First of all, the generation of full Steiner trees for all subsets of terminals in \( \mathbb{R}^d, d \geq 3 \), is much more difficult than in \( \mathbb{R}^2 \). The determination of a full Steiner tree of a given FST with more than 3 terminals requires solving high-degree polynomials \([19]\). As a consequence, iterative numerical approaches are the only possibility. Such numerical approaches seem to block the generation of full Steiner trees across various subsets of terminals (unlike the \( \mathbb{R}^2 \) case). Finally, the geometrical non-optimality tests seem to be much weaker than in \( \mathbb{R}^2 \).

While the generation of full Steiner trees is very challenging for \( d \geq 3 \), the concatenation phase could be applied without any significant modifications. However, the lack of efficient pruning tests in the generation phase permits a huge number of non-optimal RMTs to survive causing the branch-and-cut concatenation algorithm to choke. These congestion problems become more and more serious as \( d \) grows.

The branch enumeration algorithm that is proposed here can be seen as a compromise between the approach used by GeoSteiner (generation of full Steiner trees of all subsets of terminals) and the numerical approach of Smith [19] described in Section 3 (generation of RMTs for FSTs with \( n \) terminals). Rather than enumerating RMTs for all FSTs for \( N \), branches involving subsets of terminals are generated. Consider a FST \( \mathcal{T} \) of \( N \). As already mentioned in Section 2, removing a Steiner point breaks \( \mathcal{T} \) into 3 branches. As will be seen in Subsection 5.3, every FST with \( n \) terminals contains a Steiner point whose removal creates three branches with at most \( \lfloor \frac{n}{2} \rfloor \) terminals. This significantly reduces the number of branches that need to be generated.

The branch enumeration algorithm consists of three phases (see Algorithm 1). The preprocessing phase computes EMSTs for subsets with up to 8 terminals. These EMSTs are used in subsequent phases to prune away non-optimal branches and RMSTs. The second phase generates branches containing up to \( \lfloor \frac{n}{2} \rfloor \) terminals. EMSTs obtained in the preprocessing phase together with RMTs of root spliced branches are used to prune away branches that cannot be in \( \mathcal{T}^* \). The third phase generates FSTs with \( n \) terminals by concatenating triplets of disjoint branches. The iterative procedure described in Section 3 is then used to determine their RMTs. The shortest RMT encountered is \( \mathcal{T}^* \).

5.1 Preprocessing phase

The maximum size \( \kappa \) of subsets of terminals for which ESMTs are determined in the preprocessing phase has a large effect on the number of branches that will be pruned away during their generation described in Subsection 5.2. As branches of up to size \( \lfloor \frac{n}{2} \rfloor \) are generated, setting \( \kappa = \lfloor \frac{n}{2} \rfloor \) ensures that for any branch it is possible to find a disjoint preprocessed tree. However, choosing \( \kappa > 8 \) makes the preprocessing phase computationally expensive. ESMTs are therefore generated using Smith* for all terminal subsets of size 1, 2, 3, \ldots, \( \kappa \), stored in \( \mathcal{P} \), and finally sorted by non-increasing lengths.
Algorithm 1 Branch enumeration algorithm

**Input:** Set $N$ of $n$ terminals in $\mathbb{R}^d$.

**Output:** ESMT $T^*$ of $N$.

1. **Preprocess:** Determine EMSTs of all subsets of terminals of size up to $\min\{8, \lceil n^2 \rceil\}$, store them in $P$, and order them by non-increasing length.
2. **Generate:** Enumerate sets $B_k$ of feasible branches with $k$ terminals, $k = 1 \ldots \lfloor n^2 \rfloor$ (pruning away non-optimal branches).
3. **Concatenate:** Combine triplets of disjoint feasible branches spanning $N$. Compute corresponding RMTs and let $T^*$ be a shortest one.

### 5.2 Generating branches

Branches are generated by increasing number of spanned terminals. Branches of size $k$, $k = 1, 2, \ldots \lfloor n^2 \rfloor$, are stored in a set $B_k$. It will be shown in Subsection 5.3 that only branches spanning up to $\lfloor n^2 \rfloor$ need to be generated. A single terminal is a branch containing 1 terminal and is therefore stored in $B_1$. The branches in $B_k$, $2 \leq k \leq \lfloor n^2 \rfloor$, are generated by combining branches in $B_l$ with branches in $B_{k-l}$, $l = 1 \ldots \lfloor k/2 \rfloor$. When $l = k - l$, care is taken to avoid generation of duplicate branches.

There are several criteria that can be used to reject a branch $B_k = B_l \oplus B_{k-l}$. First of all, $B_l$ and $B_{k-l}$ must be disjoint. This can be efficiently verified by bitwise AND on bit-strings representing subsets of terminals. Similarly, constructing the bitstring of $B_k$ can be efficiently done using bitwise OR.

Assume next that $t_i$ is a terminal in $B_l$ while $t_j$ is a terminal in $B_{k-l}$. Furthermore, assume that the depth of $t_i$ in $B_l$ is $d_i$ and the depth of $t_j$ in $B_{k-l}$ is $d_j$. Let $d = d_i + d_j + 2$. Suppose that $|t_it_j|/d > \beta(t_i, t_j)$ (see Fig. 3). Then $B_k$ cannot be a part of $T^*$ as it would contain an edge between $t_i$ and $t_j$ longer than the bottleneck distance between $t_i$ and $t_j$.

![Fig. 3: A branch $B_k$ with terminals $t_i$ and $t_j$ being $d = 4$ edges apart cannot be a part of $T^*$ if any of these $d$ edges are longer than the bottleneck distance $\beta(t_i, t_j)$. This will certainly be the case if $|t_it_j|/d > \beta(t_i, t_j)$.](image-url)
Fig. 4: The length of $\text{RMT}(B)$ added to the EMST of any subset of terminals disjoint from $T$ is a lower bound on $|T^*|$

Finally, suppose that $\text{RMT}(B_k)$ spans the set of terminals $N_k$ and the EMST $T^+$ for some non-empty subset of $N \setminus N_k$ satisfies

$$|\text{RMT}(B_k)| + |T^+| \geq UB$$

where $UB$ is an upper bound on the length of $T^*$ (see Section 4), then $B_k$ cannot be a part of $T^*$. This is an immediate consequence of the following lemma and the fact that $UB \geq |T^*|$.

**Lemma 1** Let $B_k$ be a branch of $T^*$. Let $N_k$ denote the terminals of $B_k$. Let $T^+$ be the EMST of a non-empty subset of $N \setminus N_k$. Then $|T^*| \geq |\text{RMT}(B_k)| + |T^+|$. 

**Proof.** If $|N_k| = 1$ then $|\text{RMT}(B_k)| = 0$ and the inequality trivially holds. Assume that $|N_k| \geq 2$. Let $B_k = B_i \oplus B_j$ with $r_k$, $r_i$ and $r_j$ being the roots of $B_k$, $B_i$ and $B_j$ respectively. Let $T_k^*$, $T_i^*$ and $T_j^*$ be the parts of $T^*$ corresponding to $B_k$, $B_i$ and $B_j$ respectively. Then

$$|T_k^*| = |T_i^*| + |r_i r_k| + |r_j r_k| + |T_j^*| \geq |T_i^*| + |r_i r_j| + |T_j^*| \geq |\text{RMT}(B_k)|$$

where the first inequality is due to the triangle inequality and the second inequality is due to the fact that $\text{RMT}(B_k)$ is the RMT of the FST obtained by splicing away $r_k$ from $B_k$. Hence,

$$|T^*| = |T_k^*| + |T^* \setminus T_k^*| \geq |\text{RMT}(B_k)| + |T^+|$$

since $T^* \setminus T_k^*$ is a tree spanning $N \setminus N_k$. \hfill $\square$

### 5.3 Concatenation of branches

Consider a FST $\mathcal{T}$ of a set $N$ of $n$ terminals, $n \geq 3$. Let $s$ be any of its $n - 2$ Steiner points. When $s$ (and its three incident edges) are removed, $\mathcal{T}$ splits into three branches $B_i$, $B_j$ and $B_k$ with respectively $n_i$, $n_j$, and $n_k$ terminals, $n = n_i + n_j + n_k$. Assume that $n_i \geq n_j \geq n_k$.

**Lemma 2** $\mathcal{T}$ has a splitting Steiner point $s$ such that $n_i \leq n_j + n_k$. 


Algorithm 2 Generate branches containing up to $\left\lceil \frac{n}{2} \right\rceil$ terminals

Input: Set $N$ of $n$ terminals in $\mathbb{R}^d$.
Input: Sorted list $P$ of EMSTs for small subsets of $N$.
Input: An upper bound $UB$ on $|T^*|$
Output: $B_k, k = 1 \ldots \left\lceil \frac{n}{2} \right\rceil$

1: $B_1 \leftarrow N$
2: for $k = 2$ to $\left\lfloor \frac{n}{2} \right\rfloor$ do
3:   $B_k \leftarrow \emptyset$
4:   for $l = 1$ to $\left\lfloor \frac{k}{2} \right\rfloor$ do
5:     for all $B_i \in B_l$ do
6:       for all $B_{k-l} \in B_{k-l}$ do
7:         if $B_l$ and $B_{k-l}$ have a common terminal then
8:           $B_l \oplus B_{k-l}$ cannot be a part of $T^*$
9:         end if
10:     end for
11:   end for
12:   $B_k \leftarrow B_k \oplus B_{k-l}$
13: end for
14: end for
15: for all $B_i \in B_k$ do
16:   Find the longest ESMT $T^+ \in P$ disjoint from $B_k$
17: end for
18: if $|\text{RMT}(B_k)| + |T^+| \geq UB$ then
19:   $B_k$ cannot be a part of $T^*$
20: end if
21: Add $B_k$ to $B_k$.
22: end for
23: end for
24: end for
25: end for
26: return $B_k, k = 1 \ldots \left\lfloor \frac{n}{2} \right\rfloor$

Proof. Assume that $n_i > n_j + n_k$ for every Steiner points in $T$. Pick a Steiner point $s$ minimizing $n_i$. Let $s' \in B_i$ denote the Steiner point adjacent to $s$ in $T$. It exists since $n_i \geq 2$. Let $n'_i, n'_j$, and $n'_k$ denote the number of terminals obtained by splitting $T$ at $s'$. Then $n'_i = n_j + n_k$, $n'_j = x$ for some $x$, $0 < x < n_i$, and $n'_k = n_i - x$. Hence, $n'_i, n'_j$, and $n'_k$ are all less than $n_i$, contradicting the choice of $s$. □

A split of any FST with $n_i \geq n_j \geq n_k$, $n = n_i + n_j + n_k$ and $n_i \leq n_j + n_k$ is called a canonical split.

Lemma 3 $\left\lceil \frac{n}{3} \right\rceil \leq n_i \leq \left\lceil \frac{n}{2} \right\rceil$ in a canonical split of any FST with $n$ terminals.

Proof. To obtain the first inequality, observe that $n = n_i + n_j + n_k \Rightarrow n \leq 3n_i \Rightarrow \left\lceil \frac{n}{3} \right\rceil \leq n_i$. To obtain the second inequality, observe that $n_i \leq n_j + n_k = n - n_i \Rightarrow 2n_i \leq n \Rightarrow n_i \leq \left\lceil \frac{n}{2} \right\rceil$. □
Lemma 4  An FST $\mathcal{T}$ with $n$ terminals has exactly one canonical split unless $n$ is even and $n_i = n/2$ in which case $\mathcal{T}$ has two canonical splits at adjacent Steiner points.

Proof. Let $B_i$, $B_j$ and $B_k$ denote three branches of the canonical split at a Steiner point $s$. Consider another Steiner point $s' \in \mathcal{T}$. Assume first that $s' \in B_j$. Let $n_i' \geq n_j' \geq n_k'$ denote the number of terminals in the branches of this split at $s'$. Hence, $n_i' \geq n_i + n_k$ and $n_j' + n_k' \leq n_j$. Now $n_i' \geq n_i + n_k > n_j \geq n_j' + n_k'$ implies that the split is not canonical. Similar argument applies if $s' \in B_k$. Assume finally that $s' \in B_i$. Hence, $n_i' \geq n_j + n_k$. Now $n_i' \geq n_j + n_k \geq n_i \geq n_j' + n_k'$ implies that this split is canonical iff $n_i' = n_j' + n_k'$. This can happen if and only if $n_i' = n_i = n/2$. Hence, $n$ has to be even and $s$ and $s'$ must be adjacent in $\mathcal{T}$. $\square$

It is therefore only necessary to concatenate triplets of disjoint branches $(B_i, B_j, B_k)$ with $n = n_i + n_j + n_k$, and whose individual sizes satisfy $\left\lceil \frac{n}{3} \right\rceil \leq n_i \leq \left\lfloor \frac{n}{2} \right\rfloor$, $n_i \leq n_j + n_k$ and $n_i \geq n_j \geq n_k$. See Algorithm 3 for a detailed description of the concatenation of triplets of branches.

Algorithm 3  Concatenate triples of branches to obtain EMST $T^*$

| Input: | Sets of branches $B_i, p = 1 \ldots \left\lfloor \frac{n}{2} \right\rfloor$ |
| Input: | An upper bound $UB$ on $|T^*|$ |
| Output: | $T^*$ |

1: $|T^*| \leftarrow UB$
2: for $i = \left\lceil \frac{n}{2} \right\rceil$ to $\left\lceil \frac{n}{3} \right\rceil$ do
3:  for $j = \min\{i, n - i - 1\}$ to $\left\lceil \frac{n - i}{2} \right\rceil$ do
4:    $k \leftarrow n - i - j$
5:    for all disjoint triples of branches $B_i, B_j, B_k$ with $B_i \in B_i, B_j \in B_j$ and $B_k \in B_k$ do
6:      $T \leftarrow \text{RMT}(B_i \oplus B_j \oplus B_k)$.
7:      if $|T| < |T^*|$ then
8:        $T^* \leftarrow T$
9:      end if
10: end for
11: end for
12: return $T^*$

6  Computational experiments

Smith’s algorithm (SMITH) described in Section 3, its modified version (SMITH*) described in Section 4 and the branch enumeration algorithm (BRANCH) described in Section 5, were implemented in C++ and run on an Intel Xeon X5550 2.67GHz CPU. All runs were stopped after 24 hours if they had not terminated or if they consumed more than 10GB of memory.
Fig. 5: Average CPU-times for Smith, Smith*, and Branch on instances from the Carioca set. Each column represents an average for each method run on five problem instances with a specific set of terminals \(n\) and dimensions \(d\). All values can be found in the appendix tables.

Figure 5 shows average CPU-times for the 40 Carioca instances for each \(d = 3, 4, 5\) from the OR-library [2] (Carioca set). Tables 3 to 5 in the appendix shows the CPU times in more details as well as reported lengths of ESMTs and the total number of RMTs generated by each of the three exact algorithms. Smith* and Branch consistently outperform the original Smith algorithm\(^3\). In a few instances Branch is faster than Smith* and for some large instances it terminates using the prespecified time and memory while Smith* does not.

All three algorithms construct RMTs for various subsets of terminals in order to obtain reasonable lower bounds permitting pruning of non-optimal configurations as early as possible. In particular, Branch computes many such RMTs with few terminals in its preprocessing and generation phases. Note, that the very straightforward way that preprocessing is currently done, the RMT of a particular topology may be computed multiple times. Additionally, the RMT of two distinct branches may also be equivalent. Using appropriate look-up tables to avoid redundant RMT computations would no doubt speed up Branch significantly.

\(^3\) For \(n = 14, d = 4, 5\) the average of Smith is lower because only one fast run terminated while Smith* and Branch succesfully solved several instances.
Table 1: The effect of different pruning criteria on the number of generated branches/full topologies in BRANCH algorithm. The level of a branch is the number of terminals it contains. 'Nothing' is a theoretical approach where all combinations of branches are used in enumeration, including those that overlap. The Overlap-pruning discards all branches where the same terminal appears twice. Bottleneck-pruning additionally uses the bottleneck criterion to prune. Finally, Lowerbound and Preprocessing also calculates lower bounds and compares them with the upper bound even in the branch generation phase. The Preprocessing pruning strengthens the lower bound of each branch by adding the lengths of disjoint optimal solutions from the preprocessing phase.

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Table 1 illustrates the strength of the different pruning criteria used in BRANCH. The 'Nothing' column indicates the number of branches that are explored if all pruning is disabled. The 'Preprocessing' column indicates the number of branches that are explored when all pruning methods are enabled. Using RMTs of branches as a lower bound is essential to be able to prune in the final concatenation phase (bottom rows in the table) but it is only when combined with tighter lower bounds obtained from the preprocessing stage that branches can be pruned while being generated.

The main reason why SMITH* and SMITH were terminated was because they consumed all memory of the machine. Even when switching to a depth-first-search, the size of the branch-and-bound tree grows extremely large. BRANCH never used more than 1GB of memory, but for large instances it was stopped because it exceeded the computation time.

Table 1 illustrates the strength of the different pruning criteria used in BRANCH. The 'Nothing' column indicates the number of branches that are explored if all pruning is disabled. The 'Preprocessing' column indicates the number of branches that are explored when all pruning methods are enabled. Using RMTs of branches as a lower bound is essential to be able to prune in the final concatenation phase (bottom rows in the table) but it is only when combined with tighter lower bounds obtained from the preprocessing stage that branches can be pruned while being generated.

Table 2 compares our implementation with the three previous algorithms that solve the d-dimensional geometric Steiner tree problem. The values for SMITH-FAMPA and SMITH-LAARHOVEN are taken from [14]. For most instances both SMITH* and BRANCH are superior to the previously suggested algorithms.

Figure 6 illustrates the effects of changing the number of terminal subsets that are preprocessed. The BRANCH algorithm was run on three point sets with $d = 3$ and $n = 12, 13, 14$ and different values of $\kappa$. In all cases, setting $\kappa = \lceil n/2 \rceil$ gives the lowest CPU time. Lower values of $\kappa$ result in very fast preprocessing,
but weak lower bounds that makes the concatenation phase explode. Larger
values of $\kappa$ make the lower bound extremely strong, but forces us to spend a
lot of time on preprocessing. As a lot of redundant computations are performed
in the preprocessing phase, this indicates that speeding up preprocessing and
increasing $\kappa$ might substantially improve the Branch algorithm.

Fig. 6: The CPU time in seconds for 3D instances from the carioca set where
preprocessing finds EMSTs for subsets up to different values of $\kappa$. No matter
what $n$ is, setting $\kappa = \lceil n/2 \rceil$ gives the lowest CPU time.

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Table 2: Performance of known Steiner tree algorithms. Numbers indicate the
improvement factor in nodes explored and CPU-time over Smith's original algo-

but weak lower bounds that makes the concatenation phase explode. Larger
values of $\kappa$ make the lower bound extremely strong, but forces us to spend a
lot of time on preprocessing. As a lot of redundant computations are performed
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Fig. 6: The CPU time in seconds for 3D instances from the carioca set where
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what $n$ is, setting $\kappa = \lceil n/2 \rceil$ gives the lowest CPU time.
7 Conclusions

Two new exact algorithms for solving the Euclidean Steiner tree problem in dimensions $d \geq 3$ have been proposed. One is an extension of the seminal algorithm by Smith while the other constructs branches in a bottom-up fashion. Computational studies show that both methods are faster than the original but the improved version of Smith’s algorithm is, in general, the fastest.

It is worth noting that the use of branches in the branch enumeration method provides more specific information about partial solutions than the expanding topologies in Smith’s algorithm. This property has the potential to expose different and possibly stronger methods for fathoming partial solutions early. For instance, Laarhoven and Anstreicher [14] present a combination of lune-properties and bottleneck distances that could directly be employed and be used to prune in the very first steps of the enumeration. In general our findings suggest a critical need to identify tighter lower bounds and geometric criteria for excluding partial solutions in higher dimensions.
References


### A CPU-times and RMTs

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Table 3: Results for 40 $R^3$ instances from the Carioca set (OR-library). \[EMST\] is the length of the ESMT, \#RMT$_n$ is the number of RMTs spanning all $n$ terminals. CPU-times are given in seconds. Runs were terminated if not completed within 24 hours or if more than 10GB of memory was consumed.
Table 4: Results for 40 $\mathbb{R}^4$ instances from the Carioca set (OR-library). $|EMST|$ is the length of the ESMT, $\#RMT_n$ is the number of RMTs spanning all $n$ terminals, CPU-times are given in seconds. Runs were terminated if not completed within 24 hours or if more than 10GB of memory was consumed.
Table 5: Results for 40 R^5 instances from the Carioca set (OR-library). |EMST| is the length of the ESMT, #RMT_n is the number of RMTs spanning all n terminals, CPU-times are given in seconds. Runs were terminated if not completed within 24 hours or if more than 10GB of memory was consumed.

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