

A Hub-Based Labeling Algorithm for Shortest Paths on Road Networks

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Abraham et al. [SODA 2010] have recently presented a theoretical analysis of several practical point-to-point shortest path algorithms based on modeling road networks as graphs with low highway dimension. Among the methods they analyzed, the one with the best time bounds is the *labeling algorithm*. Their results suggest that the algorithm is interesting from a theoretical viewpoint, but leave open the existence of a practical implementation. This paper presents such an implementation and shows experimental evidence that it is actually faster than the fastest method previously studied.

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1 Introduction

Motivated by computing driving directions, the problem of finding point-to-point shortest paths in road networks has received significant attention in recent years. Even though Dijkstra’s algorithm [8, 14] solves the problem in almost linear time [17], applications to continent-size road networks require faster algorithms. Preprocessing makes sublinear-time algorithms possible.

There are too many preprocessing-based methods to describe in full—we refer the reader to [12] for a comprehensive overview. We present here only some highlights with the algorithms that are relevant to our study. Arc flags [23, 21] and landmark-based A* search [18] reduce the search space by directing the search towards the goal. Highway hierarchies [26, 27] and reach [20, 19] use hierarchical properties of road networks to sparsify the search. One of the key ingredients of efficient implementations of these methods is the notion of a *shortcut* (introduced by Sanders and Schultes [26]), which is a new arc representing a shortest path between its endpoints. Shortcuts by themselves became the basis of *contraction hierarchies* (CH) [16], an elegant algorithm that motivated much follow-up work. Another method is *transit node routing* (TNR) [3, 4], which is based on the observation that there is a small set of vertices that cover all sufficiently long shortest paths out of any region of a road network. This allows long-range queries to be reduced to a small number of table look-ups. The most efficient implementation of TNR uses CH during the preprocessing stage and to handle local queries [16]. One can combine goal-direction with hierarchical methods or TNR [19, 6, 5]. In particular, the fastest previously known point-to-point shortest path algorithm [6] combines elements from TNR (using CH) and arc flags. It is six orders of magnitude faster than Dijkstra’s algorithm for random (long-range) queries.

Although these algorithms have been shown to work well in practice, a theoretical analysis has been given only recently, in a paper by Abraham et al. [1]. It is based on modeling road networks as graphs with low *highway dimension*. The method with the best time bounds, presented by Abraham et al. as a variant of TNR, is actually a *labeling algorithm*. Labeling algorithms have been studied before in the distributed computing literature [15, 30]. In particular, [15] gives upper and lower bounds on label sizes for general graphs, trees, planar graphs, and bounded degree graphs.

Like most speedup techniques on road networks, the labeling algorithm works in two stages. The preprocessing stage computes for each vertex v a *forward label* $L_f(v)$ and a *reverse label* $L_r(v)$. The forward label consists of a set of vertices w , together with their respective distances $\text{dist}(v, w)$ from v . Similarly, the reverse label consists of a set of vertices u , each with its distance $\text{dist}(u, v)$ to v . The labels have the following *cover property*: For every pair of distinct vertices s and t , $L_f(s) \cap L_r(t)$ contains a vertex u on a shortest path from s to t . To emphasize that a labeling has the cover property, we call a labeling *valid*.

The query stage of the labeling algorithm is quite simple, given the cover property. Given s and t , find the vertex $u \in L_f(s) \cap L_r(t)$ that minimizes $\text{dist}(s, u) + \text{dist}(u, t)$ and return the corresponding path. One can think of a label for a vertex v as a set of *hubs* to which v has a direct connection. The cover property ensures that any two vertices share at least one hub on the shortest path between them.

The results of Abraham et al. [1] suggest that the labeling algorithm is interesting from a theoretical viewpoint, but leave open the existence of a practical implementation. In fact, as described in their paper, the algorithm appears impractical. First, preprocessing, although polynomial-time, is too slow for continent-size networks. Second, the worst-case bound on the

memory overhead per vertex is also too high for a practical implementation on such networks.

In this paper we give a practical implementation of the labeling algorithm. Our preprocessing algorithm uses the contraction hierarchies (CH) algorithm by Geisberger et al. [16], which we review in Section 3, and augments it with other techniques. We call our implementation HL (Hub-based Labeling algorithm). Our contributions are as follows:

- We observe that the sets of vertices visited by the forward and reverse searches of hierarchical and reach-based algorithms contain the corresponding labels. More precisely, visited vertices with exact distance values form valid labels. Similar observations have appeared for graphs of bounded tree-width [15] and for road networks [22] in an implicit manner; we make it explicit and take advantage of it.
- We use the above observation to implement a labeling algorithm by pruning the CH search space. We observe that this produces labels that are small enough to store in main memory even for continental-sized road networks.
- We show how to use ideas from “theoretical” preprocessing algorithms [1] to improve the quality of the fast heuristic CH preprocessing, leading to even smaller labels.
- We show how to implement efficiently the preprocessing and query stages of HL, including simple but effective techniques to accelerate long-range queries.
- We describe a label compression technique that significantly reduces the space requirements of the algorithm.
- We present experiments showing trade-offs of our implementation and comparing it to the fastest previous codes.

The results show that the labeling algorithm is not only theoretically efficient, but actually practical. In fact, when optimized for speed, for random queries our algorithm is about as fast as five random accesses to main memory.

The main contribution of this paper is to show that HL is practical and should be considered for real-life applications. An added bonus is that HL is currently the fastest algorithm for the problem. While simpler than the previous state of the art, the combination of TNR and arc flags (TNR+AF) discussed above, HL is faster by a factor of more than three for long-range queries (although HL memory usage is higher by a small constant factor). For short-range queries, the performance difference between HL and TNR+AF is even greater—an order of magnitude. By applying our compression techniques, HL has roughly the same memory footprint as TNR+AF, and still outperforms it.

This paper is organized as follows. Section 2 gives basic definitions and reviews Dijkstra’s algorithm. Section 3 gives a quick summary of the CH algorithm. Section 4 presents the basic version of our labeling algorithm. Section 5 shows how it can be improved, leading to an even more practical algorithm. We evaluate the algorithm experimentally in Section 6, and conclude in Section 7.

2 Preliminaries

The input to the preprocessing stage of a shortest path algorithm is a graph $G = (V, A)$ with length $\ell(a) > 0$ for every arc a . We denote the number of vertices in G by n . The length of any path P in G is the sum of the lengths of its arcs. The distance between vertices v and w , denoted by $\text{dist}(v, w)$, is the length of the shortest path between them. The query phase of the shortest path algorithm takes as input a source s and a target t , and returns $\text{dist}(s, t)$.

Dijkstra’s algorithm [14, 8] is the best-known method for computing shortest paths in our setting. It is an efficient implementation of the scanning method for graphs with non-negative arc lengths (see e.g. [29]). For every vertex v , it maintains the length $d(v)$ of the shortest path from the source s to v found so far, as well as the predecessor $p(v)$ of v on the path. Initially $d(s) = 0$, $d(v) = \infty$ for all other vertices, and $p(v) = \text{null}$ for all v .

Dijkstra’s algorithm maintains a priority queue of unscanned vertices with finite d values, the values serving as keys. At each step, the algorithm extracts a vertex v with minimum value from the queue and *scans* it: for each arc $(v, w) \in A$, if $d(v) + \ell(v, w) < d(w)$ it sets $d(w) = d(v) + \ell(v, w)$ and $p(w) = v$. The algorithm terminates when the target t is extracted, without scanning t .

3 Contraction Hierarchies

We now discuss *contraction hierarchies* (CH) [16], an acceleration technique that, on road networks, can find exact point-to-point shortest paths orders of magnitude faster than Dijkstra’s algorithm.

Most state-of-the-art shortest-path algorithms, including the CH algorithm, depend crucially on a very simple notion: *shortcuts* [26]. Given two vertices $u, v \in V$, a shortcut (or *shortcut arc*) is a new arc (u, v) with length $\text{dist}(u, v)$, the original distance in G between u and v . The *shortcut operation* deletes (temporarily) a vertex v from the graph and adds shortcut arcs between its neighbors to maintain the shortest path information. More precisely, for any neighbors u, w such that $(u, v) \cdot (v, w)$ is the only shortest path between u and w , we add the shortcut (u, w) with $\ell(u, w) = \ell(u, v) + \ell(v, w)$.

Given the notion of shortcuts, CH preprocessing is simple: define a total order among the vertices and shortcut them sequentially in this order, until a single vertex remains. The output of this routine is a graph $G^+ = (V, A \cup A^+)$ (where A^+ is the set of shortcut arcs created), as well as the vertex order itself. We denote the position of a vertex v in the ordering by $\text{rank}(v)$. Define $G^\uparrow = (V, A^\uparrow)$ by $A^\uparrow = \{(v, w) \in A \cup A^+ : \text{rank}(v) < \text{rank}(w)\}$. Similarly, $A^\downarrow = \{(v, w) \in A \cup A^+ : \text{rank}(v) > \text{rank}(w)\}$ and $G^\downarrow = (V, A \cup A^\downarrow)$.

During an s - t query, the algorithm performs a forward search from s and a reverse search from t . The *forward CH search* consists of running Dijkstra’s algorithm from s in G^\uparrow , stopping when there are no more vertices to scan. Similarly, the *reverse CH search* consists of running a reverse version of Dijkstra’s algorithm from t in G^\downarrow . With these searches, the CH algorithm computes upper bound estimates $d_s(v)$ and $d_t(v)$ on distances from s to v and from v to t for every $v \in V$. For some vertices, these estimates may be greater than the actual distances; in fact, many vertices are not visited by the search and (implicitly) have infinite estimates. However, as shown by Geisberger et al. [16], the maximum-rank vertex u on the s - t shortest path is guaranteed to be visited, $v = u$ will minimize $d_s(v) + d_t(v)$, and this will be the length of the shortest path from s to t . The path itself can be obtained from the concatenation of the s - u and u - t paths.

Note that queries are correct regardless of the contraction order, but query complexity and the number of shortcuts added may vary greatly from one permutation to the next. The best results reported in [16] are obtained by on-line heuristics that select the next vertex to shortcut based on its current degree and the number of new arcs added to the graph, among other factors.

We use a slightly different priority function for ordering vertices [9]. The priority of a vertex u is given by $2 \cdot ED(u) + CN(u) + H(u) + 5 \cdot L(u)$, where $ED(u)$ is the difference between the number of arcs added and removed (if u were shortcut), $CN(u)$ is the number of previously contracted neighbors, $H(u)$ is the total number of underlying arcs represented by all shortcuts added, and $L(u)$ is the level u would be assigned to.

The *level* of a vertex u , denoted by $level(u)$, is defined as follows. If all neighbors of u in G^+ have higher rank than u , $level(u) = 0$. Otherwise, $level(u) = level(v) + 1$, where v is the highest-level vertex among all lower-ranked neighbors of u .

4 Basic HL

In this section, we first review the theoretically justified labeling algorithm of Abraham et al. [1], then we describe our practical implementation.

4.1 The Theoretical Algorithm

The preprocessing of the theoretical labeling algorithm [1] is based on *shortest path covers* (SPCs). Intuitively, an (r, k) -shortest path cover S is a set of vertices that (1) hits every shortest path of length between r and $2r$ and (2) is *sparse*, in the sense that every ball of radius $2r$ contains at most k elements from S . Computing such covers is probably NP-hard, but the authors suggest a greedy algorithm to obtain an $O(\log n)$ approximation in polynomial time.

More precisely, the assumptions of [1] imply that for a fixed parameter h , (r, h) -SPCs exist for all r , and the greedy algorithm computes an $O(r, O(h \log n))$ -SPC. The parameter h , the *highway dimension* of the graph, is believed to be small for road networks.

As suggested by Abraham et al., preprocessing for the labeling algorithm uses the greedy algorithm to compute SPCs C_i for $r = 2^i$, $0 \leq i \leq \log D$, where D is the diameter of the graph. The paper proves that the following labeling is valid: For each v , take the union over i of C_i intersected with the ball of radius $2 \cdot 2^i$ around v . Note that the size of the label of v is $O(k \log n \log D)$.

As stated, the algorithm is not practical. First, the greedy algorithm for computing SPCs requires computing all-pairs shortest paths many times over, yielding preprocessing times of several months for continent-size networks. Second, the theoretical bound on the label size could be well into the thousands in practice, which would lead to unrealistic space requirements and uncompetitive query times.

4.2 A Practical Implementation

Before explaining our new practical implementation of the labeling algorithm, we need the notion of a *superlabel*. Given a vertex v , it is defined similarly to a label, except for the fact that a distance $d(v, w)$ stored within the label need not equal the real distance $\text{dist}(v, w)$. We only enforce $d(v, w) \geq \text{dist}(v, w)$. For simplicity, we refer to superlabels as labels except when we want

to emphasize that we are dealing with superlabels. When we want to stress that we deal with labels according to the original definition, we will refer to them as *strict labels*. We also use the same notation, $L_f(v)$ and $L_r(v)$, for superlabels. The cover property for superlabels is defined as follows: For every pair of distinct vertices s and t , $L_f(s) \cap L_r(t)$ contains a vertex u such that u is on a shortest path from s to t , $d(s, u) = \text{dist}(s, u)$, and $d(u, t) = \text{dist}(u, t)$. It is obvious that the query algorithm remains correct when superlabels are given.

The [idea](#) of our new implementation of the labeling algorithm is as follows. Given s and t , consider the sets of vertices visited by the forward CH search from s and the reverse CH search from t . CH works because the intersection of these sets contains the maximum-rank vertex u on the shortest s – t path. Therefore, we can obtain a valid superlabeling if we define, for every v , $L_f(v)$ and $L_r(v)$ to be the sets of vertices visited by the forward and reverse CH searches from v . (For undirected—i.e., symmetric—graphs, one label would be sufficient.) This is similar to the many-to-many algorithm of Knopp et al. [22], which during queries computes and stores the sets of vertices visited by the reverse search from the destinations, along with the distance labels. Note that the many-to-many algorithm implicitly uses the fact that these sets form superlabels.

On the road network of Western Europe (used for most of our experiments), [the average size of the labels computed this way is about 500](#). For a practical algorithm, we would like to have smaller labels. Note that if we have a forward superlabel $L_f(v)$, we can prune it by deleting from it vertices w with $d(v, w) > \text{dist}(v, w)$. We can prune $L_r(v)$ similarly. If before pruning u is a vertex in $L_f(s) \cap L_r(t)$ with $d(s, u) = \text{dist}(s, u)$ and $d(u, t) = \text{dist}(u, t)$, u will not be pruned. Therefore, superlabels remain valid after pruning. *Partial pruning* and *complete pruning* delete some and all vertices that can be pruned, respectively. On road networks, complete pruning reduces the label size significantly (about 80% on the benchmark instance).

We now consider a simple way of representing a label so as to allow efficient queries. We identify a vertex i with its (integer) ID. We describe the L_f labels; the L_r labels are symmetric. The label $L_f(v)$ is represented as the concatenation of three elements:

- N_v is an integer representing the number of vertices in the label.
- I_v is a zero-based array containing the IDs of all vertices in the label in ascending order.
- D_v is an array containing the distances from v to the vertices in its label (in the same order as I_v): $D_v[i] = \text{dist}(v, I_v[i])$.

With this representation, the query algorithm is straightforward. Given s and t , it must pick, among all vertices $w \in L_f(s) \cap L_r(t)$, the one minimizing $\text{dist}(s, w) + \text{dist}(w, t) = d_s(w) + d_t(w)$. The fact that the I_v arrays are sorted allows this computation to be performed in linear time, with a single sweep through the labels, similar to mergesort. We maintain array indices i_s and i_t (initially zero) and a tentative distance μ (initially infinite). At each step we compare $I_s[i_s]$ and $I_t[i_t]$. If these IDs are equal, we found a new w in the intersection of the labels, so we compute a new tentative distance $D_s[i_s] + D_t[i_t]$, update μ if necessary, then increment both i_s and i_t . If the IDs differ, we increment either i_s (if $I_s[i_s] < I_t[i_t]$) or i_t (if $I_s[i_s] > I_t[i_t]$). We stop when either $i_s = N_s$ or $i_t = N_t$, and return μ .

A crucial aspect of this algorithm is that each array is accessed sequentially. Good locality reduces the number of cache misses, which are the bottleneck of our algorithm.

For added efficiency, in practice we actually use pointer arithmetic (instead of maintaining indices) and use sentinels to detect when a particular array has been traversed in full. We discuss

further improvements to our basic algorithm in the next section.

5 Improvements

As we have seen, our basic preprocessing and query algorithms are simple, given an implementation of CH. In this section we describe improvements that speed up preprocessing and queries, or reduce space overhead of the algorithm.

When discussing algorithmic details, one should keep in mind that our fastest query implementation is less than five times slower than a random memory access, as Section 6 will show. To achieve this, we need to consider low-level implementation details carefully. Locality is particularly important, since each additional cache miss results in noticeable performance penalty.

5.1 Fast Preprocessing

We now describe a detailed implementation of the preprocessing algorithm outlined in Section 4, together with some improvements. Recall that we must compute forward and reverse labels for each vertex v .

Our algorithm works as follows. For every $v \in V$, start with $L_f(v) = \emptyset$. Run the forward CH search from v and add to $L(v)$ all vertices w with $d_s(w) = \text{dist}(v, w)$. Compute $L_r(v)$ similarly using the reverse CH search. Since a CH search is very fast (a fraction of a millisecond on Western Europe), doing the required $2n$ searches is practical.

However, while performing a search from v , for each vertex w visited we must check if its distance label $d(w)$ matches the actual distance $\text{dist}(s, w)$. We can use any efficient point-to-point shortest path algorithm for these *check queries*. In fact, as Section 5.2 will explain, one can bootstrap and use HL itself.

Even though HL is fast, it is not free. Therefore, we also use a fast heuristic modification to the CH search to identify most vertices with incorrect distance labels, thus avoiding some unnecessary check queries. The heuristic is similar to the stall-on-demand procedure used by CH and related algorithms [28]. We describe it for the forward search; the reverse search can be dealt with similarly. Suppose we are performing a forward CH search from v and we are about to scan w , which has distance label $d(w)$. The modification examines all incoming arcs $(u, w) \in A^\downarrow$. If $d(w) > d(u) + \ell(u, w)$, the distance label of w is provably incorrect. Therefore, we can safely remove w from the label, and we do not scan outgoing arcs from it. It is easy to see that the following holds:

Lemma 5.1 *Every vertex visited by the original forward CH search and assigned a correct distance label is visited by the modified CH search and assigned a correct distance label.*

We can generalize this approach for better pruning as follows. When scanning a vertex w with distance $d(w)$ during a forward CH search, we perform a graph search from w in A^\downarrow , limited to depth h . If we find a vertex u with $d(u) + d^*(u, w) < d(w)$, where $d^*(u, w)$ is the distance between u and w with respect to the graph search, we remove w from the label and do not scan outgoing arcs from w . We call this approach h -hop heuristic label pruning, where h is a user-defined parameter.

Our experiments show that only a small fraction of the vertices that remain (after the h -hop heuristic is applied) have incorrect labels, even with $h = 1$. This means one could skip the check

queries altogether to save preprocessing time, and use these partially pruned labels for queries. Still, for optimum query performance, we do perform the checks during preprocessing to obtain strict labels in most of our experiments.

Note that the labels of different vertices can be computed completely independently. Therefore, parallelizing the label computation is straightforward.

5.2 Bootstrapping

The partially pruned label preprocessing algorithm finds most vertices that can be pruned, but it may miss some. One can prune further to obtain strict labels using a point-to-point query algorithm. In this section we describe how to **bootstrap** HL, i.e., use HL itself as the query algorithm. In addition to eliminating the need to implement another algorithm, this is attractive because HL is one of the fastest point-to-point shortest path algorithms.

A simple way to do bootstrapping is as follows. First, compute partially pruned labels. Then, for every label $L_f(v)$ ($L_r(v)$), iterate through the vertices w in the label and use point-to-point queries to check if $d(w) = \text{dist}(v, w)$ ($d(w) = \text{dist}(w, v)$). If the test fails, delete w from the label.

One can do more efficient bootstrapping by pruning the labels as they are computed, eliminating the need to store bigger intermediate labels except for the current vertex. To do this, we compute the labels for vertices in descending level order. We describe how to compute the forward labels; the reverse case is similar. Suppose we have just computed the partially pruned label $L_f(v)$. We know that $d(v) = 0$ and that all other vertices w in $L_f(v)$ have higher level than v , which means $L_r(w)$ must have already been computed. We can therefore compute $\text{dist}(v, w)$ using $L_f(v)$ and $L_r(w)$, removing w from $L_f(v)$ if $d(w) > \text{dist}(v, w)$.

If we use bootstrapping during preprocessing, we have to be more careful for a parallel implementation. Computing the label of a vertex v in level i requires access to labels of vertices in levels higher than i . We can process vertices of the same level in parallel, but we have to synchronize the procedure after each level. Fortunately, the number of levels is small (about 150) on road networks. More importantly, most of the vertices are in the lowest five levels [9].

5.3 Label Ordering

Our preprocessing is CH-based and uses CH ordering of vertices. Once the labels are computed, however, we can reorder vertices (i.e., assign them new IDs) to speed up queries. We call the newly assigned ID of a vertex its *internal ID*. Our experiments show that certain label orderings yield better query times and compression rates.

Besides keeping the original input order, we also considered assigning new IDs in ascending or descending level order (keeping the input order within each level). Moreover, we tested using the CH-rank as internal ID, as well as the inverted CH-rank. Interestingly, as we shall see in Section 6, we obtained the best results by keeping the input order for all but the topmost (highest-ranked) k vertices, which are assigned internal IDs from 0 to $k - 1$. We considered two variants of this method, depending on how IDs are assigned among the top k vertices. In the *top k input* order, their relative order is the same as in the input; in *top k level* order, these vertices are sorted by level.

One optimization we apply to all label orderings is to assign ID $|V|$ to the most important vertex. Since it is in *every* label of the graph (assuming the graph is strongly connected, which

we do), this vertex then acts as a sentinel during our sweep through the labels, thus simplifying our termination checks.

5.4 Shortest Path Covers

As already mentioned, computing exact SPCs (or even approximations) is not practical for continent-size networks. However, variants of the greedy algorithm can be applied to the smaller contracted graph towards the end of CH preprocessing. In this section, we show how to do this. As we shall see in Section 6, this leads to a better CH vertex ordering and smaller labels.

CH tends to contract the least important vertices (those on few shortest paths) first, and the more important vertices (those on more shortest paths) later. The heuristic used to choose the next vertex to contract appears to work well at the beginning of the algorithm, when choosing unimportant vertices. However, we observed that it works poorly near the end of preprocessing, when it must order important vertices relative to one another.

We now propose an algorithm that uses shortest path covers to improve the ordering of important vertices. We modify our preprocessing algorithm as follows. We start by running the CH preprocessing with our original selection rule. Instead of running it to completion, however, we pause it when there are only t vertices left (a reasonable number is $t = 10\,000$). Let G_t be the graph at this point; it is an overlay of the original graph, i.e., it preserves the distances between the remaining vertices.

In a second step, we run a greedy algorithm to find a set C of good “cover” vertices, i.e., vertices that cover a large fraction of all shortest paths of G_t , with $|C| < t$ (we use $|C|$ between $t/20$ and $t/10$ in our experiments). Starting with $C = \emptyset$, at each step we add to C the vertex v that hits the most uncovered (by C) shortest paths in G_t .

Once C has been computed, we continue the CH preprocessing, but forbid the contraction of the vertices in C until they are the only ones left. This guarantees that the top $|C|$ vertices of the hierarchy will be exactly the ones in C , which are then contracted in reverse greedy order (i.e., the first vertex found by the greedy algorithm is the last one remaining).

Greedy Cover Algorithm. We now describe the greedy algorithm mentioned above in more detail. Let C be the (initially empty) set we are building, and suppose we need to compute the next vertex to add to it. We do so by keeping a counter $h(v)$ for each vertex $v \notin C$. Initially set to zero, this counter will eventually contain the number of uncovered shortest paths that are hit by v . We grow a shortest path tree T_r from each root $r \in G_t$. Let $h(r, v)$ be the number of shortest paths starting at r that are hit by v but not by C ; the summation of $h(r, v)$ over all r is $h(v)$.

To compute $h(r, v)$, remove from T_r all vertices already in C , together with their descendants; the paths from r to any of these vertices are already hit by C . Then traverse the resulting tree T'_r in bottom-up fashion. If v is a leaf of T'_r , set $h(r, v) = 1$; otherwise, set $h(r, v)$ to the sum of the $h(r, u)$ values of its children u in the tree, plus one (to account for the path from r to v itself).

As stated, the algorithm requires computing t full shortest path trees to insert each new vertex in C . We can accelerate this process by stopping the growth of the tree as soon as all of its labeled vertices are on paths already hit by C . This does not help much while C is relatively small (there is always an uncovered branch), but it becomes very effective as C increases in size. Still, the running time in the worst case is $O(Ct^2 \log t)$, assuming the graph has constant average degree.

However, we expect a significant boost in performance by computing shortest path trees with PHAST [9].

5.5 Label Compression

In this section we show how the labels can be compressed, reducing the memory consumption of HL. We consider two independent techniques: using fewer bits to represent small IDs in each label, and sharing common parts among different labels. We discuss each in turn.

5.5.1 Compressing Small IDs

The first compression technique is quite simple. Normally, we represent vertex IDs and distances as (separate) 32-bit integers. To compress the label, we represent each of the first 256 vertices as a single 32-bit word, with 8 bits allocated to the ID and 24 bits to the distance, which is enough for the instances tested. Of course, one could generalize this scheme by splitting each word into b bits for IDs and $\omega - b$ bits for distances, where ω is the word size.

This 8/24 compression technique changes the representation of only 256 out of millions of vertices. To take full advantage of this method, we must reorder vertices so that the most important vertices (which tend to be in most labels) are the ones with the lowest IDs. For this compression scheme, a top 256 input or level ordering is most suitable.

This compression scheme is simple and adds little overhead to queries; in fact, they actually get faster because of better locality.

5.5.2 Compressing Common Prefixes

We now propose a different compression technique. For it to be effective, we must reorder vertices so that the important ones are assigned the very lowest IDs. For concreteness, assume we reorder vertices in descending level order. For two nearby vertices in a road network, their forward (or reverse) CH trees are different near the root but are often the same when sufficiently away from the root. Furthermore, the vertices away from the root are the high-level (“important”) ones. By reordering vertices in descending level order, the labels of nearby vertices will often share long common prefixes, with the same sets of vertices.

The intuition for our compression scheme is to compute a dictionary of the common label prefixes and reuse them. One complication is that, even when two labels have prefixes with the exact same vertices, the corresponding distances are, in general, different. We overcome this issue by decomposing each dictionary prefix into a forest and storing distances relative to the roots.

We now describe this *k-prefix compression* method in detail. We explain how forward labels can be compressed; the reverse case is symmetric. First, for each label $L_f(v)$ we do the following. Let $P_k(v)$ be the *prefix* of the label, consisting of those vertices whose internal ID is lower than k . Let $S_k(v)$ be the *suffix* of the label, consisting only of the vertices whose internal ID is at least k . In other words, the vertices in the prefix are those among the k most important in the graph; the others are in the suffix. The parameter k is the *prefix threshold*.

Consider the forward CH search tree T from v . The subgraph of T induced by $S_k(v)$ either is empty or forms a tree containing v ; the subgraph induced by $P_k(v)$ is a forest. Consider this forest. First suppose that $S_k(v)$ is non-empty. For each tree T_i of the forest, let $b(T_i)$ (the *base* of T_i) denote the parent of T_i ’s root. By extension, the base of a vertex $w \in P_k(v)$, denoted by

$b(w)$, is defined as $b(T_i)$, if $w \in T_i$. Note that $b(w) \in S_k(v)$ by definition. In the special case in which $S_k(v)$ is empty (the complete tree T is in $P_k(v)$), we define $b(v) = v$.

For each w in $P_k(v)$, we store the distance (in the tree) between $b(w)$ and w ; we denote it by $\delta(w)$. We also store $b(w)$ itself, as the position $\pi(w)$ of $b(w)$ in $S_k(v)$ (to save space).

Altogether, each prefix consists of a list of triples $(w, \delta(w), \pi(w))$. We consider two prefixes to be *equal* if they consist of the exact same triples, and *different* if at least one triple does not match. We build a dictionary consisting of all *distinct* prefixes, which are stored sequentially in an array.

To represent a forward label $L_f(v)$, we combine three elements: its prefix $P_k(v)$ (represented as a reference to its position in the dictionary), the number of vertices in the prefix, and the suffix $S_k(v)$ (represented as before). Note that we do not use 8/24 compression to represent the top 256 vertices.

During a query from v , suppose w is in $P_k(v)$. Note that $\text{dist}(v, w) = \text{dist}(v, b(w)) + \text{dist}(b(w), w)$. We can compute the distance in constant time: we precomputed $\text{dist}(b(w), w) = \delta(w)$ and we know the position $\pi(w)$ of $b(w)$ in $S_k(v)$, where $\text{dist}(v, b(w))$ is stored explicitly.

5.5.3 Flexible Prefix Compression

The main drawback of the k -prefix compression is that we must use the same prefix threshold for all labels, even though some labels may share longer prefixes than others. In this section we show how to extend k -prefix approach to exploit this. In the *flexible prefix compression* scheme, we allow each label L to be split arbitrarily into a suffix and a prefix (there are $|L|$ ways to do so, if $|L|$ is the number of entries in the label). As before, common prefixes are represented only once and shared among labels.

Our goal is to minimize the total space consumption, considering the sizes of all (exactly n) suffixes and all (at most n) prefixes we actually keep. Obviously, there is an optimum way of splitting each label so as to minimize this value, but it is unclear how it can be found efficiently. Instead, we use heuristics to find a good solution.

To do so, we state our problem as an instance of *facility location* [25]. We can interpret each label L as a *customer* that must be represented (served) by a suitable prefix (facility). Deciding which prefixes to keep is equivalent to deciding which facilities to open. The cost of opening a facility is the size of the corresponding prefix. The cost of serving a customer (label) L by a prefix P is the size of the corresponding suffix ($|L| - |P|$). Each label L must be served by the available prefix that minimizes the service cost.

Although the facility location problem is NP-hard [2], there are good heuristics to solve it in practice [25]. For efficiency, we adopt a three-phase approach. First, we use a greedy algorithm to find an initial set of prefixes. Second, we use a simplified local search that greedily adds or removes individual prefixes from the current solution until its total cost no longer decreases. Finally, we run a more elaborate local search [25] which also allows swaps (i.e., simultaneous insertions and deletions) of prefixes. We stop when a local minimum is reached.

The quality of the final solution found by this algorithm does not depend much on the initial set of prefixes, but a good initial solution helps to reduce the time to reach a local optimum. To generate an initial solution, we start with an empty set and greedily add to it the prefix P that maximizes $(\text{avg}_{L \in C(P)}(|P|/|L|))^2 \cdot \log(|C(P)|)$, where $C(P)$ is the set of yet uncovered labels that have P as a prefix.

5.6 Partition Oracle

In this section we describe how to improve the performance of HL on long-range queries. In particular, this accelerates *random queries*, in which the source s and target t are picked uniformly at random. If the source and the target are far apart, the CH searches tend to meet at very important (high-rank) vertices. If we rearrange the labels such that more important vertices appear before less important ones, long-range queries can stop traversing the labels when sufficiently unimportant vertices are reached. To accomplish this, while maintaining correctness, we must make slight changes to the preprocessing algorithm.

During preprocessing, we first find a good partition of the vertices in the graph into cells of bounded size, while trying to minimize the total number b of boundary vertices. Intuitively, we would like the number k of cells to be large and b to be small, but there is a trade-off between these two in practice.

Second, we perform CH preprocessing as usual, but delay the contraction of boundary vertices until the contracted graph has at most $2b$ vertices. We do so by blocking the contraction of boundary vertices (during the CH algorithm) while the graph remains large, but restoring their normal priorities once the number of vertices is down to $2b$. Let B^+ be the set consisting of all vertices that have rank as least as high as that of the lowest-ranked boundary vertex. This set includes all boundary vertices and has size $|B^+| \leq 2b$.

Third, we compute labels in normal fashion, except that we store at the beginning of a label for v the ID of the cell v belongs to.

Fourth, for every pair of cells (C_i, C_j) , we run HL queries between every vertex in $B^+ \cap C_i$ and every vertex in $B^+ \cap C_j$, and keep track of the internal ID of their meeting vertex. Let m_{ij} be the maximum such ID over all queries made for this pair of cells. We then build an $k \times k$ matrix, with entry (i, j) corresponding to m_{ij} . Note that building this matrix requires up to $4b^2$ queries in total.

This concludes the preprocessing stage.

We run s - t queries as usual, looking at vertices in increasing order of internal ID, but stopping as soon as we reach (in either label) a vertex with internal ID higher than m_{ij} . We can do so because m_{ij} is an upper bound on the internal ID of the meeting vertex for any query between C_i and C_j . This strategy requires one extra memory access to retrieve m_{ij} , but for long-range queries we end up looking at a fraction of each label. Also note that this approach only unfolds its full potential if important vertices have low internal IDs.

5.7 Index-free Labels

To perform an s - t query, HL must bring from memory two labels, $L_f(s)$ and $L_r(t)$. To figure out where in memory these labels are, we need to access the entries for s and t in an *index array*. When applying all the speed-oriented optimizations described above, random s - t queries become so fast that these two accesses end up being a significant fraction the running time (about 20%).

We can avoid these accesses by essentially eliminating the index arrays, as follows. We reserve c bytes in each label array (forward and reverse) for each label. If a label $L_f(v)$ has size at most c , we store it starting at position $v \cdot c$ in the forward label array (the reverse case is similar). Otherwise we store as much of $L_f(v)$ as possible in the initial array, and the remaining entries in a third array (which we call *escape array*). In this case, the label array also stores an index to the escape array as part of each label.

During an s - t query, we can start reading the label arrays directly (with no need for an index). If, however, we realize that the sentinel is not among the first c bytes, we continue reading the label from the escape array. Note that this approach increases the memory footprint of HL, since short labels will not use their reserved space in full. On the other hand, if a label fits into its pre-allocated space, performance improves as we do not need to access its index. The choice of c determines the trade-off between memory consumption and query performance.

6 Experimental Results

Implementation Details. We implemented our algorithm in C++ and compiled it with Microsoft Visual C++ 2010. We use OpenMP for parallelization during preprocessing. The evaluation was conducted on a machine equipped with two Intel Xeon X5680 processors and 96 GB of DDR3-1333 RAM, running Windows 2008R2 Server. Each CPU has 6 cores clocked at 3.33 GHz, 6 x 64 kB L1, 6 x 256 kB L2, and 12 MB L3 cache.

Since query times are very small, we forced procedure inlining whenever appropriate (on our default machine, a function call takes about 150 ns, roughly the time of 3 memory accesses). For the same reason, we prefetch data to the L1 cache whenever appropriate.

For the uncompressed version of HL, we use four arrays, two for each direction. In the following, we explain the implementation of the forward labels. The *index array* has n 64-bit entries indicating the starting points of each label on the second array, called **data**. The **data** array has 32-bit unsigned integers storing both vertex IDs and distances. We store the label $L_f(v)$ of a vertex v in **data** in $2|L_f(v)|$ consecutive entries. The first entry is the distance to the sentinel, i.e., the most important vertex. The next $|L_f(v)|$ entries are the label IDs of the vertices of $L_f(v)$, in increasing order. The remaining $|L_f(v)| - 1$ entries are the corresponding distances, in the same order. The rationale for this approach is that we must access almost all IDs in a label, whereas distances only need to be checked when the IDs match. Separating IDs from distances leads to fewer cache misses. Another optimization we apply is to align each label to cache lines, which have 64 bytes in our machine.

Storing the distance to the sentinel as the first entry achieves two goals. First, we can quickly initialize an upper bound on the s - t distance by adding up the sentinel distances from s and to t . Second, it facilitates efficient implementation of partition oracles.

The 8/24 compression is implemented similarly to the uncompressed version. The only difference is that the vertices with small internal IDs (up to 255) are stored together with their distance label, right after the sentinel distance. In the remainder of the label, the high ID vertices are represented as before.

The suffixes of the prefix compression approach are stored as in the uncompressed variant. The only difference is that, at the very beginning (before the sentinel distance), we store a pointer (an index) to the beginning of the prefix. As discussed in Section 5.5, the prefix is a list of triples (distance to the forest root, ID, and offset). We store each triple in 64 consecutive bits: 32 bits for the ID, 8 bits for the offset, and 24 bits for the distance. Note that, unlike in our other implementations, we do not split IDs and distances. Moreover, since the prefix-based variant is optimized for memory consumption, it does not align each label with a cache line.

The partition oracle is implemented as an array of 32-bit integers. We tested using 16-bit integers instead, but the impact on running times is limited.

Methodology. We use two input graphs in our experiments, both taken from the webpage of the 9th DIMACS Implementation Challenge [13]. The *Europe* instance represents the road network of Western Europe, with 18 million vertices and 22.5 million road segments, and was made available by PTV AG [24]. The *USA* road network (generated from TIGER/Line data [31]) has 24 million vertices and 29.1 million road segments. Unless otherwise mentioned, we use the Europe instance as default.

HL uses contraction hierarchies during preprocessing. We implemented a parallelized version of the CH preprocessing routine, as discussed in [16], using the priority function described in Section 3 to order the vertices. With this priority term, preprocessing takes about three minutes (using all twelve cores) and generates upward and downward graphs with 33.8 million arcs each.

In the following, we report the (parallel) preprocessing time (without the CH preprocessing) and total space consumption in GB. Query performance is evaluated by running 100 000 000 s - t queries, with s and t picked (in advance) uniformly at random. In most experiments, we report average query times. Note that we use parallelization only during preprocessing; queries are executed on only one core. Also note that we consider the scenario in which only the shortest path distance is to be computed, and not the full path. If full descriptions of the shortest paths are needed, one could apply some of the path-expansion techniques used for TNR [3], for example.

The next section studies the impact of various design decisions on the performance of our algorithm. This will allow us to pick the best set of parameters for our algorithm in different scenarios. In Section 6.2, we compare these optimized versions of HL with other methods in the literature.

6.1 Trade-offs and Parameter Tuning

Impact of Label Pruning. Our first experiment studies how different label-pruning strategies affect label size and query performance. The input is the augmented graph produced by the CH algorithm; we build the labels by running upward searches on this graph. We tested the following pruning strategies: 1-hop heuristic, 2-hop heuristic, and 1-hop heuristic combined with bootstrapping. The results can be found in Table 1. Note that we do not use any compression in this experiment.

boot	hops	preprocessing			queries
		label time [s]	vertices /label	space [GB]	time [ns]
×	0	321	536.31	154.4	> 3000
×	1	385	133.03	37.0	937
×	2	491	113.05	31.5	834
✓	1	580	109.62	30.6	812

Table 1: Impact of label pruning on the performance of HL. We consider heuristic pruning with variable number of hops and exact pruning (bootstrapping).

The first row in the table estimates the performance of our algorithm with no pruning, i.e., if we used the entire CH search space as labels. Note that we did not actually run this variant in full, as it requires more memory than our machine has. The numbers provided are estimates based on sampling.

Among the pruned variants, observe that the 1-hop strategy already reduces the average label size by a factor of 4. Bootstrapping reduces the sizes of the labels even further, which also has a positive effect on query performance. Since bootstrapping does not slow down the preprocessing by too much, we use it by default, combined with the heuristic 1-hop pruning.

Impact of Label Ordering. As explained in Section 5, the order in which vertices are represented within the labels may have an impact on query performance. In Table 2, we consider six different strategies to assign internal IDs to vertices. The first (*input*) just keeps the initial input IDs. Second, we consider an *inverted level* assignment, in which higher-level vertices are assigned lower IDs, with the input order maintained within each level. The third and fourth strategies in the table consisting of “promoting” only the 256 highest-ranked vertices (according to the CH preprocessing), keeping the remaining vertices in input order. The two variants of this strategy differ on whether the 256 most important vertices are sorted in input or level order. Finally, we consider sorting all vertices in level and rank order.

ordering	preprocessing		queries	
	label time [s]	vertices /label	space [GB]	time [ns]
input	580	109.62	30.6	812
inverted level	705	109.62	30.6	1134
top 256 input	451	109.62	30.6	769
top 256 level	469	109.62	30.6	812
level	629	109.62	30.6	1155
rank	651	109.62	30.6	1143
inverse rank	599	109.62	30.6	1128

Table 2: Impact of label vertex ordering on the performance of HL.

Overall, keeping the original order for most vertices seems to be a good idea. Rearranging them by level (either ascending or descending) or rank has a negative impact on performance. This can be explained by the fact that there is locality in the original order: the IDs of nearby vertices in the graph are likely to be similar. Moreover, if two small regions are far from each other, it is often the case that all vertex IDs in one region are larger than all vertex IDs in the other. In particular, during an s - t query, this is often true for the regions around s and around t , which constitute a large portion of their respective labels. Recall that an HL query keeps one pointer for each label, advancing in each step the one pointing to the lowest ID. Because of the locality in the input, it often reaches the end of one label (thus stopping the search) while visiting a fraction of the other. Rearranging vertices by level or rank destroys this locality and decreases query performance.

We can, however, obtain additional performance gains by making the 256 most important vertices have the lowest IDs, while keeping the input order for the remaining vertices. For random queries, where s and t tend to be far apart, most searches meet only at these vertices. By keeping them close in the label, we improve locality when accessing actual distances. Unless otherwise stated, we use a top 256 input ordering as default for the remainder of this paper.

Impact of Shortest Path Covers. We now consider the impact of using shortest path covers (SPCs) to improve CH ordering during preprocessing. We test three variants. The first is the plain CH ordering already described, which takes about 3 minutes of preprocessing. The second variant (512/10k) computes 512 cover vertices picked from the topmost 10 000 vertices of the contraction hierarchy, and takes 25 minutes. Finally, the third variant (2048/25k) picks 2048 vertices from the topmost 25 000, and takes 151 minutes. Table 3 reports the results for HL, and Figure 1 plots the distribution of label sizes in each case.

SPC	preprocessing			query
	label time [s]	vertices /label	space [GB]	time [ns]
none	451	109.62	30.6	769
512/10k	482	85.79	24.2	598
2048/25k	489	84.74	23.9	594

Table 3: Impact of shortest path covers. We use top 256 input ordering. Note that preprocessing time does not include the time for computing the shortest path covers.

Computing shortest path covers during CH preprocessing has a significant impact on the performance of HL. Labels shrink by about 20%, and queries become about 30% faster. It appears that CH preprocessing does a poor job when picking the most important vertices. This has been observed before [16]: a variant of transit node routing based on CH yielded worse locality filters than previous algorithms. In the following, we use the 2048/25k variant as default, since our emphasis is on query time. If preprocessing time is an issue, the 512/10k variant is a good alternative.

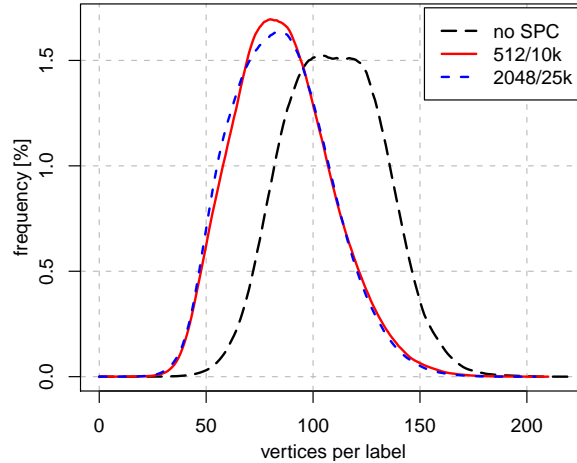


Figure 1: Distribution of label sizes when different shortest path covers are used during CH preprocessing.

Impact of Compression. With the labels optimized both by pruning and shortest path covers, we still have to store about 85 vertices per label, resulting in a total space consumption of 23.9 GB. Next, we check how well compression works.

We start by studying the k -prefix compression scheme, with k fixed for all labels. Figure 2 reports what happens when we vary the prefix threshold, which indirectly bounds the number of entries in the prefix. (Recall that if the threshold is k the prefix will have all vertices with ID lower than k .) The first plot shows the average total space consumption in GB. The second shows the average suffix size (in number of vertices) and the average number of labels that share each prefix, given by $2n/(\text{number of distinct prefixes})$. As the threshold increases, the average suffix length gradually decreases (which helps the compression), but the average prefix is shared by fewer and fewer vertices. For the instance tested, the best trade-off is obtained when the threshold is about 2^{16} (65 536): the space consumption is minimized at about 8.0 GB, a third of the original value.

Table 4 compares the performance of this variant with the uncompressed version, the simple 8/24 compression scheme, and the flexible prefix compression technique. As expected, both prefix implementations are slower than the uncompressed variant. Not only must they perform more memory accesses, they also use inverse rank ordering; the other variants use 256 top input ordering, which is better. Both variants of prefix compression (fixed and flexible) have the same query performance, but the flexible approach needs significantly less space. Unsurprisingly, preprocessing for the flexible approach is much more time-consuming.

In contrast, the 8/24 compression actually improves query performance. Because the labels are smaller, we have better locality. Since the 8/24 compression is clearly superior in terms of query times, we use this variant as default for the remainder of the paper.

Impact of the Partition Oracle. As discussed in Section 5, we can use a partition oracle to speed up long-range queries. In this experiment, we use the PUNCH algorithm [10] to partition the graph into cells with up to U vertices each, while attempting to minimize the number of boundary arcs. We tested values of U ranging from 5 000 to 50 000. On Europe, PUNCH needs 2 to 3 minutes to generate such partitions, with the number of boundary arcs ranging from 13 680 (with $U = 50\,000$) to 60 883 (for $U = 5\,000$).

Then, we compute a contraction hierarchy with delayed boundary vertices, as described in

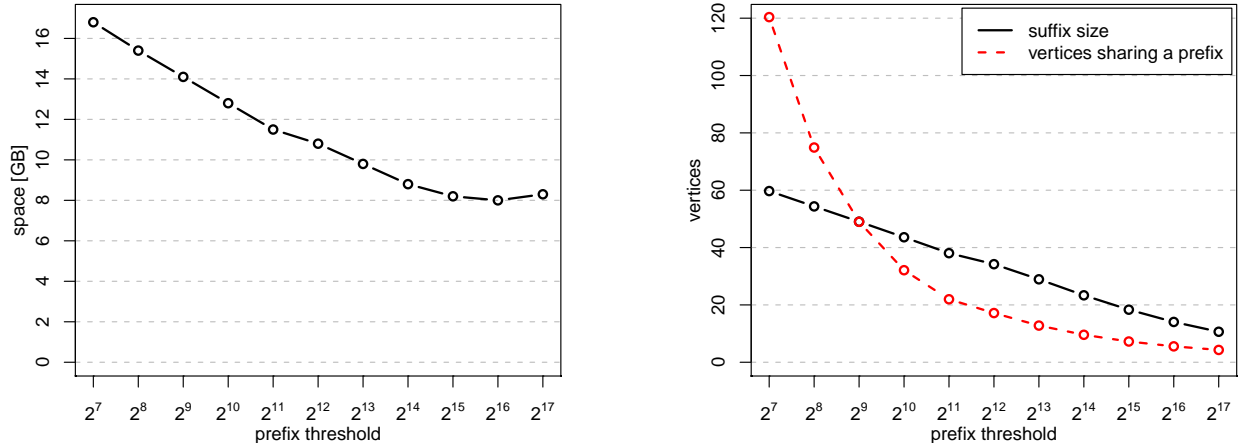


Figure 2: Label properties when varying the prefix threshold. On the left, we show the total space consumption in GB. On the right, we show the average number of vertices per suffix, as well as the average number of vertices that share each prefix.

compression	preprocessing		query
	label time [s]	space [GB]	time [ns]
none	489	23.9	594
8/24	471	20.1	572
prefix 65536	502	8.0	1172
prefix flexible	2002	5.6	1170

Table 4: Impact of compression. We use 256 top ordering for “none” and “8/24,” and inverse rank ordering for both prefix compressions.

Section 5. Finally, we compute the labels and the oracle table. Table 5 reports the results for HL with different cell sizes. Note that preprocessing times include neither partitioning the graph (1–2 minutes) nor constructing the contraction hierarchy (about 2.5 hours).

U	preprocessing			query	
	label time [s]	oracle time [s]	vertices /label	space [GB]	time [ns]
5000	425	1440	86.22	20.6	360
10000	441	589	85.97	20.6	358
20000	464	243	85.63	20.5	357
50000	443	72	86.44	20.7	366

Table 5: Impact of partition oracle for varying cells sizes. We use 8/24 compression, a top 2048 level ordering, and 2048/25k shortest path covers. Preprocessing times do not include partitioning the graph or building the hierarchy.

We observe that, below 50 000, the sizes of the cells have almost no impact on query performance. The reason for this is that with decreasing cell sizes, the quality of the oracles improve, but at the price of a bigger array. If the oracle array is small, the data is more likely to be in cache. However, when we increase U to 50 000, we see a penalty in performance. Since computing the oracle is much faster for cells with $U = 20\,000$, we settle for this parameter. In general, we observe an additional speedup of about 1.6 over HL without a partition oracle.

Index-free Labels. Next, we evaluate the performance of the index-free variant of HL. Table 6 reports space consumption and query performance when varying the number of reserved bytes per label, with and without a 20k partition oracle. Note that we reserve only multiples of a cache line, which is 64 bytes on our default machine.

The table clearly shows that removing the index does lead to additional speedups. Without an oracle, however, this only holds if the number of reserved bytes is large enough. In contrast, with a partition oracle the number of reserved bytes has almost no impact on query performance. The reason for this is that we often only look at the beginning of a label when performing random queries. In fact, we observe the lowest query times when reserving only 512 bytes per label, which keeps the additional overhead for removing the indexing structure very low (less than 5%). Hence, we consider this version of HL as our default.

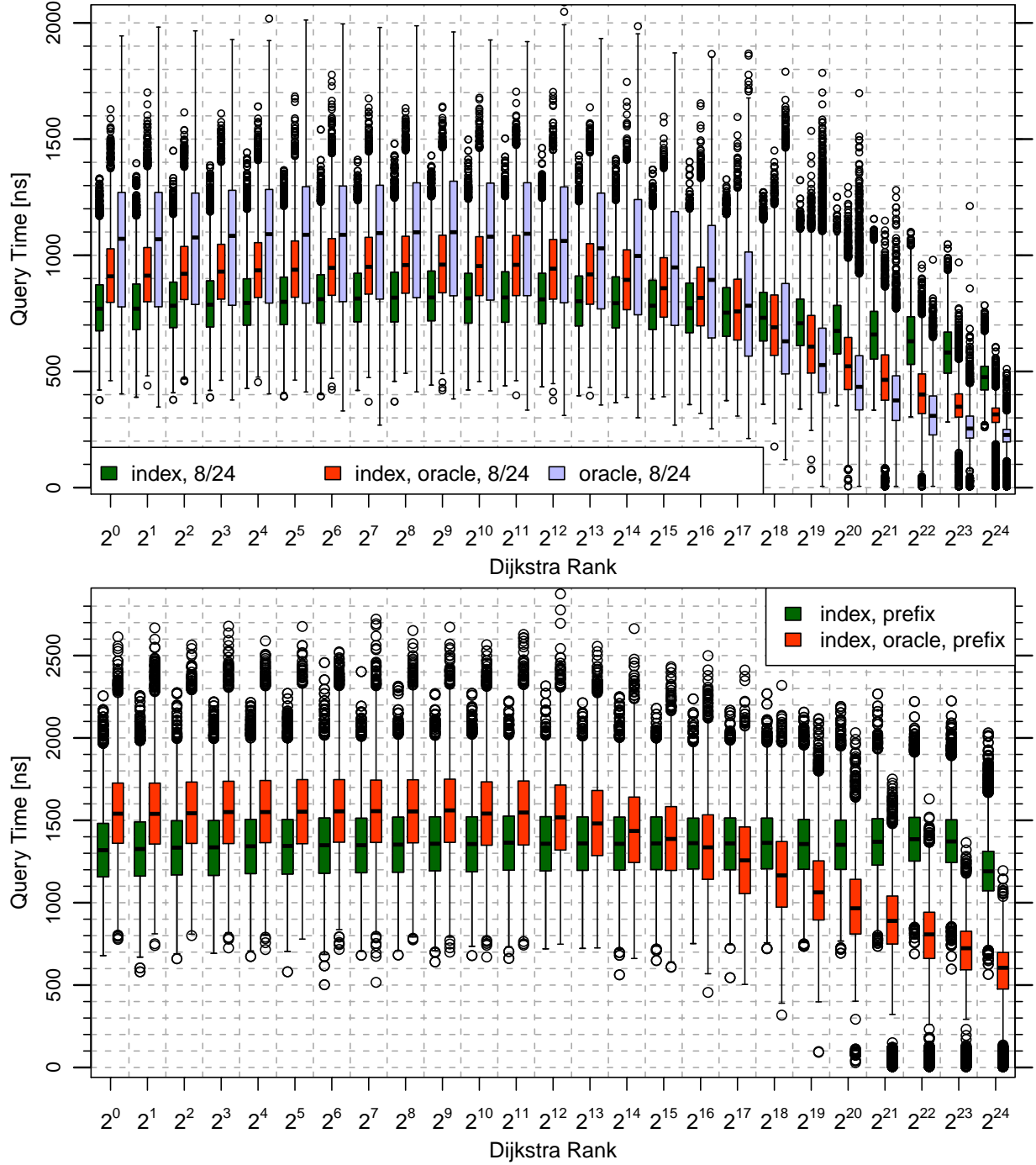


Figure 3: Comparison of different HL variants using the Dijkstra rank methodology [26]. The results are represented as box-and-whisker plot: each box spreads from the lower to the upper quartile and contains the median, while the whiskers extend to the minimum and maximum value omitting outliers, which are plotted individually. Note that the scale on the vertical axis differs slightly between the plots.

	NO ORACLE					20K ORACLE				
	preprocessing			query		preprocessing			query	
	reserved bytes	labels split [%]	label time [s]	space [GB]	time [ns]	labels split [%]	label time [s]	oracle time [s]	space [GB]	time [ns]
448		78.6	477	20.6	689	79.7	420	263	20.9	282
512		61.8	568	20.9	650	64.3	533	265	21.3	276
576		43.8	492	21.7	606	47.4	453	255	22.0	282
640		28.1	547	22.8	602	31.6	442	253	23.1	283
704		16.2	494	24.3	553	18.9	441	252	24.5	282
768		8.5	470	26.1	509	10.3	443	253	26.2	280
832		4.0	494	28.1	497	5.1	498	253	28.1	283
896		1.8	589	30.1	491	2.3	433	256	30.1	284
960		0.7	492	32.2	487	0.9	443	256	32.2	284
1024		0.2	556	34.4	479	0.3	445	258	34.4	280

Table 6: Performance of index-free HL for varying reserved bytes for the labels. We use 8/24 compression and a 256 top input ordering when using no oracle. Otherwise, we use a 2048 top level ordering.

Local Queries. Up to now, we have only evaluated the performance of HL using random queries, which are mostly long-range. In real-world applications, however, most queries tend to be local. The fastest existing methods (based on TNR) are one order of magnitude slower for local queries than for long-range ones. As we shall see, this is not the case for HL. To evaluate our algorithm on queries of various ranges, Figure 3 plots query times against Dijkstra rank [26]. For a search from s , the Dijkstra rank of a vertex v is i if v is the i -th vertex scanned when Dijkstra’s algorithm is run from s . We run 10 000 queries per rank, with s chosen uniformly at random. Since running times are very low, we have to be careful when running this test. First of all, in order to reduce cache effects, we choose a different set of 10 000 sources for each rank. Moreover, we use the internal operation counter of the CPU to determine the execution time of a single run (this is no problem because we pin queries to one core). On our 3.33 GHz machine, this yields a resolution of 0.3 ns for our timer. Finally, we repeat the whole experiment nine times and report the median (over all nine executions) of each individual query. Again, in order to reduce cache effects, we perform 100 million random queries between two subsequent runs.

The figure contains five different versions of our algorithm, differing in which optimizations (index, oracle) are activated and which compression technique is applied (8/24, flexible prefix). The first plot contains methods optimized for speed (with 8/24 compression), while the second contains methods with lower space usage.

As the first plot shows, HL has limited dependence on the locality of a query compared to TNR. Without an oracle, long-range queries are a factor of two faster than short- and mid-range queries. The median is always well below 1 μ s and only outliers take 1.5 μ s or more. All queries are answered below 2.2 μ s.

As expected, using an oracle only accelerates long-range queries. In fact, some queries are answered almost instantaneously (about 5 ns): in rare cases, all relevant information (oracle entry and labels) is by luck in L1 cache; if the oracle predicts that the search meets at the sentinel, it stops immediately. Short- and mid-range queries are actually slower with an oracle,

due to the different label ordering (top 2048 level with oracle, top 256 input without). Moreover, accessing the oracle array can be (relatively) costly. Removing the index slows down short-range queries even more, since these queries tend to access the escape array.

Comparing both plots, we observe that using flexible prefix instead of 8/24 compression slows down queries by about 500 ns, independent of the Dijkstra rank. Still, all queries are answered in less than 3 μ s. Interestingly, with prefix compression and no oracle, query times are almost independent of the Dijkstra rank: Since we use inverted ranks to set the internal IDs, both labels are traversed almost to the end on all queries.

To summarize, the index-free version with partition oracle is the best choice for long-range queries, with local queries four to five times slower. The version with index and without partition oracle is the fastest for local queries; it has almost the same performance for all query types.

6.2 Comparison with Existing Algorithms

In this section we compare HL to five previously known fast algorithms. The first one, CH [16], has been discussed in Section 3. The second algorithm, CHASE, is a combination of CH and arc flags [6].

The third algorithm is High-Performance Multi-Level Routing (HPML) [11]. This is the fastest previous algorithm for local queries. During preprocessing, it computes a hierarchy of graph separators, and uses it to compute and store many small (sparsified) graphs. Level graphs encode the distances between separators of a component on the same level, while upward (downward) graphs store the distances between separators and more (less) important separators in the same component. The query algorithm of HPML first connects the appropriate sequence of small graphs and then performs a (linear-time) search on the resulting graph. While the preprocessing is very time-consuming, this was the first approach to handle random queries in a matter of tens of microseconds. Unfortunately, the published implementation does not support very local queries.

The next algorithm in our study is transit node routing (TNR), introduced in [3]. As already mentioned, this approach uses a distance table to answer long-range queries very efficiently. For each vertex u , TNR preprocessing computes its *access nodes*, a set of nodes that covers all sufficiently long shortest paths starting or ending at u . Moreover, the distances between u and its access nodes are precomputed. Finally, a full distance table between all access nodes (collectively known as *transit nodes*) is computed and stored. On a continent-size road network, one can pick around 10 000 transit nodes such that each vertex has less than 10 access nodes on average.

The query algorithm uses a Euclidean-based *locality filter* to determine if a query is local or global. Local queries are handled by a Dijkstra-based algorithm (CH in the fastest version [16]). For a global s - t query, the shortest path contains an access node of s and one of t . Therefore, the query examines all pairs x, y where x and y are access nodes of s and t , respectively, and finds a pair that minimizes $d(s, x) + d(x, y) + d(y, t)$.

Long-range queries are dominated by distance table lookups, of which there are less than 100 on average. This is significantly faster than a typical local query implemented by CH. The most efficient version of TNR introduces several *layers* of transit nodes to accelerate mid-range queries.

The last algorithm we consider is the combination of TNR and arc flags (TNR+AF) [6]. Arc flags reduce the average number of table lookups by an order of magnitude, to less than four [6], making this the fastest previously known algorithm for random queries.

We compare these algorithms with three variants of HL, chosen based on the experiments

in the previous section. They differ in which optimizations (index, 20k oracle) are applied, the label ordering, and which compression technique is used (8/24, flexible prefix). The *prefix* variant is optimized for space consumption, the *local* version is optimized for fast short- and mid-range queries, while the *default* variant is optimized for random (long-range) queries. All techniques rely on a contraction hierarchy enhanced by shortest path covers. Table 7 details the differences between these main variants of HL.

variant	label ordering	compression	oracle	index	optimized for
HL prefix	inverse rank	flexible prefix	✓	✓	space consumption
HL local	top 256 input	8/24	×	✓	local queries
HL	top 2048 level	8/24	✓	×	random queries

Table 7: Overview of our main HL variants.

Note that our test machine is faster than the one used to test the other algorithms [6, 11]. To make a fair comparison, we also ran HL on an older machine, a dual AMD Opteron 250 running Windows 2003 Server clocked at 2.4 GHz, with 16 GB of RAM and 2 x 1 MB of L2 cache. It is known [19] to have roughly the same performance as the one in [6, 11], but unfortunately it does not have enough memory to preprocess all variants of HL for Europe. Hence, we evaluated two subgraphs of our standard inputs, representing California (1.89M vertices, 4.65M arcs) and Belgium (0.46M vertices, 1.18M arcs). A random query (HL local) on California takes 390 ns on our default machine, but 747 ns on the AMD, a factor of 1.915 slower. The slowdown for Belgium is very similar (1.908). Based on these figures, when reporting (random) query performance in Table 8 for other techniques, we also report their *scaled* running time, which is the published value divided by 1.915. The running times in Figure 4 are scaled as well.

method	EUROPE				USA			
	preprocessing		query		preprocessing		query	
	time [h:m]	space [GB]	time [ns] real scaled		time [h:m]	space [GB]	time [ns] real scaled	
CH	0:25	0.4	180000	93995	0:27	0.5	130000	67885
CHASE	1:39	0.6	17300	9034	3:48	0.7	19000	9922
HPML	≈ 24:00	3.0	18800	9817	≈ 24:00	5.1	19300	10078
TNR	1:52	3.7	3400	1775	1:30	5.4	3000	1566
TNR+AF	3:49	5.7	1900	992	2:37	6.3	1700	888
HL prefix	0:45	5.7	527	527	0:40	6.4	542	542
HL local	0:08	20.1	572	572	0:07	22.7	627	627
HL	0:14	21.3	276	276	0:18	25.4	266	266
Table Lookup	> 14:01	1 208 358.7	56	56	> 29:52	2 293 902.1	56	56

Table 8: Comparison of different speedup techniques. For HL, preprocessing times do not include the more than two hours required for enhanced CH preprocessing. Moreover, our preprocessing is parallelized, while previous methods have sequential preprocessing. For the table lookup implementation, the preprocessing time is only a lower bound, since it does not include the time to bring the distances computed on the GPU to main memory.

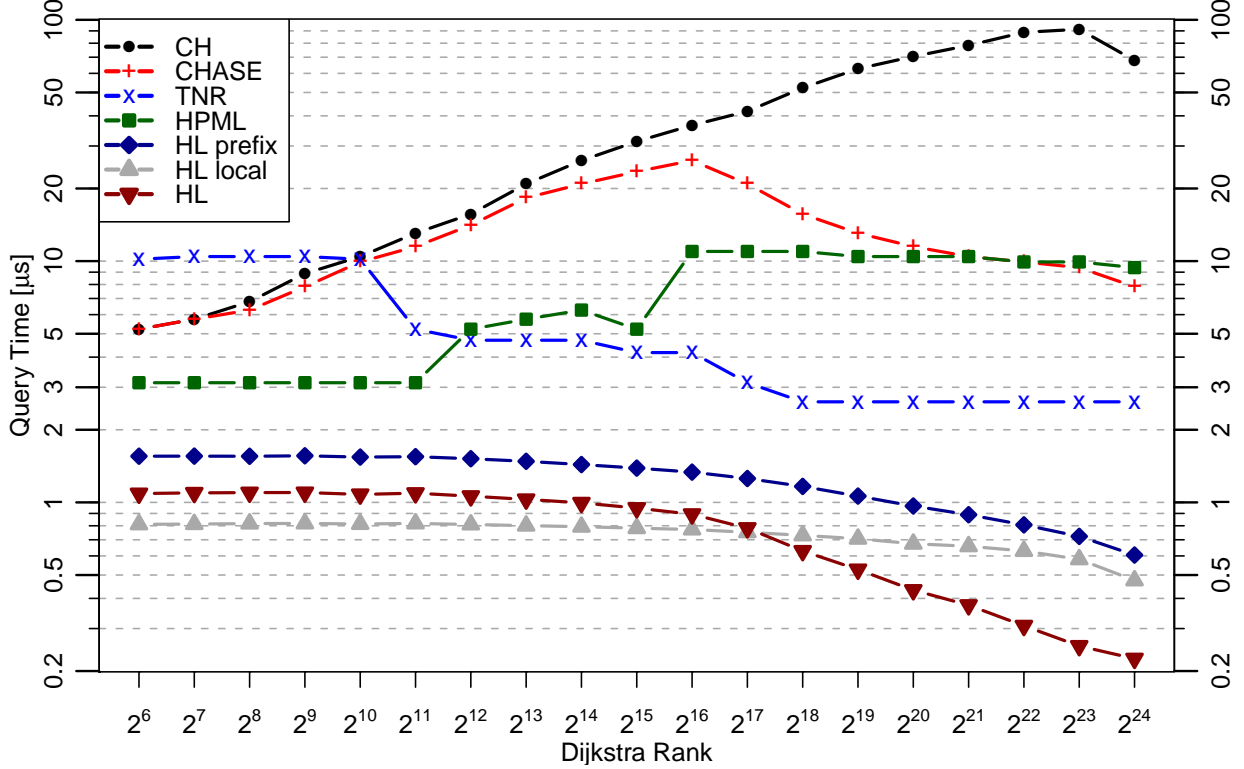


Figure 4: Performance of our main HL variants compared to various speedup techniques with respect to the Dijkstra rank. For each rank we report the median over 10 000 queries. Note that the (scaled) data for CH, CHASE, and TNR is taken from [12]. The data for HPML is also scaled and is taken from [11].

The table also compares HL to a (hypothetical) implementation of table lookups, which would run all-pairs-shortest-paths during preprocessing and store a distance table. A query would then consist of a single access to the appropriate position in the table. While preprocessing times are feasible with GPHAST [9] (on a GPU), the space consumption of this approach is way too big. We estimate its query performance using the time to access a random element of an array with 4 billion 32-bit integers.

For random queries, we observe that all variants of HL outperform all previous techniques. The default version of HL is about 3.5 times faster than TNR+AF, and six times faster than pure TNR. More importantly, as Figure 4 shows, TNR falls short when running short- or mid-range queries, where it takes 4 μ s to 10 μ s. For such queries, HL local is over an order of magnitude faster than TNR. Since TNR+AF only accelerates long-range queries compared to TNR, this observation also holds for TNR+AF. (Unfortunately, there are no published values for short- and mid-range TNR+AF queries.) Although HL performs more operations than TNR+AF, because of better locality it makes fewer accesses to the main memory, which explains why it is faster.

For short-range queries, HPML is the fastest previous algorithm. However, it is a factor of four slower than local HL and almost three times as slow as our default algorithm. Furthermore, HPML is more than one order of magnitude slower than HL for long-range queries and cannot

handle some short-range queries at all.

We also observe that compression succeeds; for Europe, HL prefix has roughly a four times lower space consumption than HL but is only twice as slow (for all types of queries). Even so, this version outperforms TNR+AF as well.

Comparing HL to a hypothetical table lookup implementation, we observe that the space consumption of HL is 56 730 (Europe) to 90 311 (USA) times lower, and it has much more practical preprocessing times. Still, random queries are only five times slower. In other words, with our optimizations a random HL query is as fast as five random accesses to main memory. For short- and mid-range queries, however, HL local is about 13 times slower than a random access.

7 Concluding Remarks

We have presented an efficient implementation of Hub Labels (HL), the algorithm with the best theoretical bounds among those studied by Abraham et al. [1]. We have demonstrated that HL is practical; in fact, it is the fastest known speedup technique for computing point-to-point distances on road networks. Although ours is not a strict implementation of the theoretical algorithm, it makes use of shortest path covers (the basis of the theoretical analysis) to obtain additional speedups.

In addition to an efficient implementation of the basic algorithm, we introduced two label compression techniques, a distance oracle to accelerate long-range (and random) queries, and the index-free variant of the algorithm. These techniques can be combined, resulting in many variants of HL. Three variants are the most interesting ones. The variant with index, no distance oracle, and 8/24 compression is simple and the fastest on local queries. Our default variant, which adds distance oracles and is index-free, is the fastest on random queries, a convenient benchmark that is used extensively in the literature. Adding prefix compression decreases the space requirements of our algorithm at the expense of the running time. The resulting algorithm uses as much space as TNR+AF, but is still faster on random queries and an order of magnitude faster on local queries.

Due to the excellent query performance, it is often feasible to run many point-to-point HL queries to solve basic problems in road networks. For example, for our Europe instance, a very efficient implementation of Dijkstra’s algorithm [17] needs 2.21 seconds to compute the distances from one vertex to all others. Computing these distances with HL takes only 0.93 seconds. Note that we need less time per entry (51.8 ns) than random queries do, since the forward label of the root vertex stays in L1 cache and reverse labels are traversed sequentially. Another advantage of running HL queries instead of Dijkstra’s algorithm is its easy parallelization. However, HL cannot catch up with PHAST [9], which needs less than 1 ns per distance entry.

We note that some of the ideas presented in this paper could probably be used to accelerate other techniques as well. In particular, a variant of our partition oracle could be used within transit node routing as a locality filter. During preprocessing, we first partition the graph into k regions. We then create a $k \times k$ boolean matrix in which entry (A, B) is true iff every s – t query with $s \in A$ to $t \in B$ is *global*, i.e., if it can be solved using transit nodes. To build this table, we need to run queries among the boundary vertices and external access nodes of all regions.

Finding better shortest path cover algorithms is an interesting research topic. On the theoretical side, we have a recent (unpublished) result showing how to improve the approximation bounds of [1] using an algorithm for finding hitting sets in low VC-dimension [7]. On the practical side,

it may be possible to improve the speed or the quality of cover computations, which could lead to smaller labels. A sufficiently fast algorithm may allow CH preprocessing based exclusively on shortest paths covers, which could lead to a better CH ordering. Similarly, computing HL labels based exclusively on shortest path covers is an obvious research direction. We expect that improved SPC algorithms and better orderings will have a positive effect on the query performance of HL.

Of course, it is also interesting to improve HL itself. Further improvements in speed are always desirable, but a more practical goal would be to further reduce the space usage of our algorithms. One could also think of how HL can be adapted to work in augmented scenarios like time-dependent networks or multi-criteria optimization. Memory consumption is again an issue, but compression and approximation techniques may help.

Another interesting question is minimizing the average label size. One can ask for both upper and lower bounds, both experimental and theoretical. On the theoretical size, the problem of finding the optimal labeling is probably NP-hard, but it is unclear whether a constant-factor approximation can be obtained in polynomial time.

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