



Article Authenticating *q*-Gram-Based Similarity Search Results for Outsourced String Databases

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Abstract: Approximate string searches have been widely applied in many fields, such as bioinformatics, text retrieval, search engines, and location-based services (LBS). However, the approximate string search results from third-party servers may be incorrect due to the possibility of malicious third parties or compromised servers. In this paper, we design an authenticated index structure (AIS) for string databases, which is based on the Merkle hash tree (MHT) method and the *q*-gram inverted index. Our AIS can facilitate verification object (*VO*) construction for approximate string searches with edit distance thresholds. We design an efficient algorithm named GS^2 for *VO* construction at the server side and search result verification at the user side. We also introduce an optimization method called GS^2 -opt that can reduce *VO* size dramatically. Finally, we conduct extensive experiments on real datasets to show that our proposed methods are efficient and promising.

Keywords: approximate string search; edit distance; query result authentication; database outsourcing; inverted index

MSC: 68P20

1. Introduction

The advancement of big data technologies for data collection, data storage, and data analytics has led to a dramatic increase in the volume of data managed. As a result, organizations, especially small- and medium-sized enterprises, are usually overwhelmed by the maintenance and processing of huge amounts of data. To tackle these challenges, cloud computing and database outsourcing have emerged in the past decade and have become the mainstream computing paradigm today. Database outsourcing (*DBO*) means that the data owners (*DO*) delegate their data to a third-party service provider (note that we also refer to it as the server in the following), and the server takes charge of data storage for *DO* and query processing for users.

Although the *DBO* paradigm offers many benefits to companies to cope with the challenges brought by big data, it also poses threats to data privacy and security since *DO* is usually reluctant to reveal their data to the public and the users do not want their query results to be tampered with. A potential threat is that the server may not answer user queries honestly in order to save computational costs; for example, the server may only process query requests on a small part of the data. Another risk is that the query results may be compromised by an untrusted server. Therefore, it is essential for the user to verify or authenticate whether the query results provided by the server are correct.

Many real-world databases contain a variety of textual data, such as personal information, blogs, emails, documents, scientific reports, genome data, etc. In certain scenarios, even the entire data records in a relational database are regarded as textual data during processing [1]. Some companies may resort to the *DBO* paradigm by outsourcing their



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Copyright: © 2023 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). textual data to third parties in order to reduce the overhead of database maintenance. On the other hand, some organizations choose to maintain textual data and process user queries on their own; for example, a tool called BLAST (basic local alignment search tools) is provided to the public to search for similar nucleotide sequences. Note that in either method of database maintenance, there may exist security risks, e.g., the server might be malicious or might have been hacked; hence, the query result returned from the server may not be trusted anymore.

In this paper, we consider the problem of result authentication for approximate string searches. Currently, there exists some research work on text/string search result authentication. For example, Pang et al. designed an elegant solution for text search result verification [2]; however, the solution focuses on authenticating TF-IDF-based text search results and is not suitable for string search result authentication. Dong et al. [3] proposed an authentication mechanism for an outsourced string similarity search using the MB tree method, which integrates the B^{ed} tree [4] and MHT methods [5]. Similarly, this work cannot be applied to scenarios where strings are indexed by a *q*-gram inverted index.

As a general case of string matching, approximate string searches aim to find *similar* strings with respect to a given string. Approximate string searches have a wide range of applications, such as search engines, duplicate detection, spell checking, plagiarism checking, and genome data analysis. Various metrics are employed to measure the similarity between strings [6], with edit distance being one of the most frequently utilized metrics. However, the time complexity of directly computing edit distance between two strings is high.

Many methods have been proposed to speed up the edit-distance-based computation of approximate string searches. One widely used approach is the *q*-gram method, which takes *q*-grams as signatures to filter the candidate set of similar strings, followed by a refinement step to verify the remainder in the candidate set. The *q*-gram method is efficient in that it can filter candidates quickly without having to scan the entire set of strings, thus speeding up the approximate string search process.

In this paper, we first design an authenticated index structure for strings, which is based on the Merkle hash tree (MHT) method [5] and the *q*-gram inverted index. Then, we introduce two schemes, namely GS^2 and GS^2 -opt, for authenticating approximate string search results, with the latter being more efficient in terms of communication overhead. Both GS^2 and GS^2 -opt comprise several procedures, including approximate string search processing, verification object (*VO*) construction, and result verification. Our contributions in this paper are summarized as follows:

- We design an authentication structure by incorporating MHT, a popular authenticated data structure, with *q*-gram-based inverted index [7], an index commonly used to accelerate similar string searches.
- We design an authentication method called *GS*² for approximate string searches. It is a general authentication scheme suitable for several string similarity measures, such as Edit distances, Hamming distances, and Cosine similarity.
- We develop an optimal method named *GS*²-*opt* that reduces the size of *VO* by combining multiple filtering techniques with an optimized scheme for edit-distance-based searches.
- We complement the proposed methods and conduct extensive experiments on real datasets to verify the effectiveness of our method. According to the experimental data, the *GS*²-*opt* approach can reduce the size of *VO* by 96.27% in comparison to the *GS*² scheme.

The paper is organized as follows. We introduce preliminaries and give a brief overview of related work in Section 2. In Section 3, we formally define the problem of authenticating the results of approximate string searches. In Section 4, we present our approaches, named GS^2 and GS^2 -opt, to solve the problem. Then, we analyse the security of our approach in Section 5. We present our experimental results which verify the effectiveness of our method in Section 6. Finally, we conclude the paper in Section 7.

2. Background

In this section, we will introduce the related work and preliminary knowledge relevant to the paper.

2.1. Approximate String Search

Given a query string s_Q , a set D of strings, and a threshold δ , the problem of approximate string searches (or string similarity searches) is to retrieve all strings $s \in D$ such that the similarity score (based on some specific measure) between s and s_Q is not less than δ . Due to its extensive applications, approximate string searches have garnered a significant amount of attention [4,8,9].

There are several challenges in processing approximate string searches, and the most challenging one is to process approximate string searches with scalability. Many approaches have been put forward, including some that introduce novel index structures [4,10–12], while others concentrate on optimizing the search process [8,13–17].

2.2. Edit Distance

Edit distance is the most commonly used measure of string similarity, which indicates the minimum number of single-character operations that transform one string into another. Generally, the permitted edit operations are insertion, deletion, and substitution, and one can assign a specific cost or weight to each of these operations. Currently, the simplest edit distance is the Levenshtein distance [18], where the cost of each operation is 1 (except for a character substitute itself, which is 0). Note that we use the Levenshtein distance as the edit distance in this paper.

Let $ed(s_1, s_2)$ be the edit distance between two strings s_1 and s_2 . Given two strings s_1 and s_2 , if $ed(s_1, s_2) \le k$, where k is a pre-specified threshold, then we say that s_1 and s_2 are similar. Wagner et al. [19] presented a classical algorithm for computing edit distance with a time complexity $O(|s_1| * |s_2|)$, where $|s_i|$ is the length of s_i .

2.3. q-Gram-Based Inverted Index

Since the calculation of the edit distance is time-consuming, preprocessing is usually employed to speed up the calculation process, during which features of strings are generated first and then indexed subsequently. Commonly used index structures include the *q*-gram inverted index, trie index, suffix tree index, and BWT index.

The *q*-gram inverted index [20] is one of the most commonly used indexing methods for strings. Basically, *q*-grams of a string *s* are all substrings (grams) of length *q*, and there are |s| - q + 1 such substrings in total. We call the set of those substrings the *q*-gram set , denoted by grams(*s*, *q*). For instance, consider a string s = "words"; the 2-gram set of *s* is {"wo", "or", "rd", "ds"}, i.e., grams(*s*, 2) = {"wo", "or", "rd", "ds"}. It is intuitive that similar strings share a certain number of *q*-grams.

The *q*-gram-based inverted index consists of two components, i.e., a dictionary of grams and a set of inverted lists (note that we also call it the *i*-list for brevity). Each entry of the dictionary includes a gram and a pointer to the corresponding *i*-list of the gram, and the *i*-list is a collection of string *IDs* (denoted by *Sid*), where each string contains the gram as a substring. To illustrate, an example string set is given in Table 1, and its corresponding 2-gram inverted index is shown in Table 2.

Currently, much research effort is being put into using the *q*-gram based inverted index to speed up string similarity searches [9,21–25]. Jokinen et al. [21] quantitatively revealed the characteristics of sharing grams between query strings and candidate strings (as presented in Lemma 1), thus laying down a foundation for fast candidate string filtering without scanning the entire set of strings.

Sid	String
1	arise
2	braise
3	fraise
4	praise
5	fraise praise raise
6	rise

Table 1. An example string set.

Table 2. 2-gram inverted index of Table 1.

Gram		Inverted List
ai	\mapsto	2, 3, 4, 5
ar	\mapsto	1
br	\mapsto	2
fr	\mapsto	3
is	\mapsto	1, 2, 3, 4, 5, 6
pr	\mapsto	4
ra	\mapsto	2, 3, 4, 5
ri	\mapsto	1,6
se	\mapsto	1, 2, 3, 4, 5, 6

Lemma 1. If $ed(s_i, s_Q) \le k$, then at least $|s_Q| - q + 1 - kq$ grams are shared by both s_i and s_Q [21], i.e.,

$$ed(s_i, s_Q) \le k \land |s_Q| - q + 1 - kq > 0$$

$$\Rightarrow |s_Q| - q + 1 - kq \le |grams(s_i, q) \cap grams(s_Q, q)|.$$
(1)

2.4. Result Authentication for String Search

To solve the problem of text search result authentication, Pang et al. proposed an elegant solution for text search engines based on an inverted index [2]. Later, inspired by this work, Goodrich et al. extended Pang's solution to support web content searches by authenticating web crawlers [26]. However, these approaches are not suitable for approximate string search authentication because they use similarity functions for documents, such as TF-IDF, which cannot be used to measure the similarity between strings.

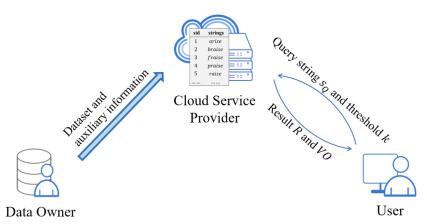
Dong et al. presented an authentication structure that integrates the B^{ed} tree and MHT methods to prove the soundness and completeness of approximate record-joining results [1] and for outsourced string similarity searches [3]. Their approach relies on a specific data structure, i.e., the B^{ed} tree structure, and cannot be employed directly to a scenario where strings are indexed by the *q*-gram inverted index.

3. Problem Formulation

In this section, we describe the system model, threat model, and design goals.

3.1. System Model

Our authentication scheme for approximate string searches involves three parties, i.e., the data owner (*DO*), the cloud service provider (or server), and the user (or client), as shown in Figure 1. Here, *DO* refers to the entity that possesses a textual dataset *D* and intends to outsource it to the server. To assist users to authenticate the search results from the server, the *DO* needs to upload additional auxiliary information along with the dataset *D* to the server. The server is responsible for database maintenance and query processing. When a user sends a query request with a query string s_Q and a similarity threshold *k* to the server, the latter processes the query against *D*, collects the result *R*, generates a verification object (*VO*), and returns *R* and *VO* to the user. After receiving *R* and *VO*, the user can



verify (1) whether the server has performed the query honestly and (2) whether the result R is correct.

Figure 1. The textual database outsourcing model.

3.2. Threat Model

The assumption made in this paper is that the server is not completely trustworthy, and thus, the results returned by the server may be incomplete or unsound. Therefore, the user must verify the results to guarantee both *completeness* and *soundness*. By *completeness*, we mean that the query result set *R* includes all the qualified strings from the dataset *D* based on the user's query. On the other hand, by *soundness*, we mean that all the strings in *R* indeed come from *D* without being tampered with and meet the query criteria.

To be more specific, given a query Q issued by the user, let R_{DO} and R_S be the query result set obtained at the DO side (Note that DO is always honest, so R_{DO} is complete and sound) and at the server side, respectively. We have the following inference:

- R_S is complete, if $\forall s_i \in R_{DO} \Rightarrow s_i \in R_S$ holds.
- R_S is *sound*, if $\forall s_i \in R_S \Rightarrow s_i \in R_{DO}$ holds.

Note that we also use *correct* to refer to the situation where both completeness and soundness are achieved.

We present some instances of incorrect results in Figure 2, where the string set is $D=\{"Alice", "Police", "Marialice"\}$, and the correct result for a query with $s_Q="Molice"$ and k=2 is $R_{DO}=\{"Alice", "Police"\}$. Figure 2a shows that an item in R_{DO} has been eliminated by the server. Figure 2b shows that an additional item that does not satisfy the query requirements has been appended to the result R' by the server. Figure 2c shows that the server has fabricated an item and adds it to the result R'. Figure 2d shows that the server has altered the item from "Police" to "Police", and, concretely, this case violates both the completeness and soundness properties.

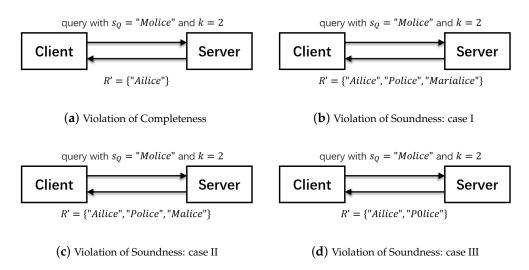


Figure 2. Examples of incorrect results.

3.3. Design Goals

There are two key goals for our solution, i.e., security and efficiency. Security means that our solution is capable of detecting incomplete or tampered query results for the user. Efficiency means that our solution should not require accessing the entire database for query result authentication. Meanwhile, our solution should also aim to minimize the size of the *VO* so as to cut down communication overhead between the server and user, as well as the time cost in verification at the user side.

4. Our Solutions

In this section, we first present GS^2 , an efficient authentication method designed for approximate string searches on outsourced string databases that are indexed by a *q*-grambased inverted index. Then, we introduce an optimization method called GS^2 -opt, which reduces *VO* size dramatically. Both GS^2 and GS^2 -opt can prove to the user that (1) any string in *D* that is similar to s_Q is included in *R*, and there is no qualified string in *D* left behind, and (2) all the strings in *R* are indeed similar to s_Q and are not tampered with.

Basically, our methods consist of three phases: (1) a preprocessing phase, during which the *DO* creates an authenticated data structure and signs it. Then, the *DO* uploads *D*, the root signature (*sig*) of the structure and parameter *q* to the server; (2) a searching phase, during which the server receives a query request from the user, performs a string search against *D* to find query result *R*, and constructs a *VO* with respect to *R*. The server then returns both *R* and *VO* to the user; and (3) an authentication phase, during which the user verifies the correctness of the query result *R* against *VO*.

$4.1.~GS^2$

We present the details of the three phases of GS^2 in the following.

4.1.1. Preprocessing Phase

During this phase, the *DO* constructs and signs the authenticated index structure (AIS) for *D* before outsourcing *D* to the server. Our AIS mainly consists of two parts, namely, a dictionary and an inverted index. Thanks to its efficiency and simplicity, we use MHT for authentication of the data structure construction. We summarize the construction process in Algorithm 1.

Algorithm 1: Generate signature for dictionary.	
Input: dataset <i>D</i> , private key <i>s</i> _k	
Output: Signature <i>S</i> _{dic} of the dictionary	
1 \\construct a MHT of D and then sign the root of the MHT ;	
2 $D \leftarrow sort(D)$; $id \leftarrow 0$; $nodes \leftarrow \emptyset$;	
s foreach $s \in D$ do	/* compute h_t */
4 $s.append(id + +);$	
5 $h_t(t) \leftarrow h(h(s.Sid) h(s.string));$	
6 nodes.append($h_t(t)$);	
7 while $len(nodes) > 1$ do	/* build MHT */
$s \mid new_nodes \leftarrow [];$	
9 foreach $node_l$, $node_r \in nodes$ do	
10 $ node_f \leftarrow h(H(node_l) H(node_r));$	
<pre>11</pre>	
12 $_$ nodes \leftarrow new_nodes;	
13 $S_{dic} \leftarrow sig_{s_k}(nodes);$	
14 return S _{dic} ;	

First, we show how to generate a signature for the dictionary. As shown in line 2 of Algorithm 1, the *DO* sorts the strings in *D* in alphabetical order and assigns each string a unique ID, i.e., *Sid*, as shown in Table 1. The reason to sort *D* is that the strings of a query result tend to share a common prefix; hence, sorting helps to cluster similar strings together, which can reduce *VO* size and thus save network costs.

The *DO* computes the tuple hash h_t for each tuple $t(Sid, string) \in D$ as follows:

$$h_t(t) = h(h(t.Sid)|h(t.string)),$$
(2)

where | stands for string concatenation, and $h(\cdot)$ is a one-way hash function. Lines 3 to 6 of Algorithm 1 show how to compute tuple hashes. We use the tuple hash as an identifier for a tuple.

The *DO* builds the MHT of *D* in a bottom-up manner. Specifically, if node *i* of the MHT is a leaf node, then its hash is simply $h(i) = h_t(t_i)$, where t_i is the data tuple corresponding to node *i*. On the other hand, if node *i* is an internal node, then its hash can be calculated as $h(i) = h(h_{node_l} | h_{node_r})$, where h_{node_l} and h_{node_r} are the hash values of the left and right children of node *i*, respectively. This process repeats until the root node of the MHT is obtained. This construction is illustrated in Figure 3, where *Hi* is the hash value of node *i*, and the "root hash" (denoted as H_r) is a digest of all the tuples in the Merkle hash tree. We summarize the calculation of MHT in lines 7 to 12 of Algorithm 1.

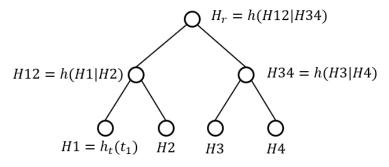


Figure 3. An example of MHT.

After the MHT is built, the *DO* generates a pair of keys (p_k, s_k) , where p_k and s_k refer to the public key and private key, respectively. Then, the *DO* signs the root hash h_r of the

MHT with the *DO*'s private key s_k , i.e., $S_{dic} = sig_{s_k}(H_r)$, where sig(.) is a standard digital signature algorithm, such as RSA and ECDSA.

Next, we explain how to compute the signature of the inverted index, as shown in Algorithm 2. Specifically, the *DO* creates an inverted index for the q-grams (lines 2 to 8 of Algorithm 2), where q is a hyper-parameter that can be pre-determined based on some statistics of dataset and user queries, such as the distribution of the alphabet and the average length of the strings.

Input: dataset D, private key s_k , gram length q Output: S_{inv} 1 $inv_gram \leftarrow \emptyset$; nodes $\leftarrow []$; 2 foreach $s \in D$ do /* build $inv_index */$ 3 grams \leftarrow grams (s, q) ; 4 foreach $g \in grams$ do 5 if $g \in inv_gram$ then 6 $inv_gram.g.append(s.Sid)$; 7 else 8 $inv_gram[g] \leftarrow [Sid]$ 9 foreach $g \in inv_gram$ do /* compute $h_t */$ 10 $h_{list} \leftarrow h(concat(g.Sid))$; 11 $h_t(gram) \leftarrow h(h(g) h_{list})$; 12 nodes.append($h_t(gram)$); 13 while $len(nodes) > 1$ do		
$i inv_gram \leftarrow \emptyset; nodes \leftarrow [];$ $2 \text{ foreach } s \in D \text{ do} /* \text{ build inv_index }*/$ $3 grams \leftarrow grams(s, q);$ $4 \text{ foreach } g \in grams \text{ do}$ $5 if g \in inv_gram \text{ then}$ $6 inv_gram.g.append(s.Sid);$ $7 else$ $8 [inv_gram[g] \leftarrow [Sid]$ $9 \text{ foreach } g \in inv_gram \text{ do} /* \text{ compute } h_t */$ $10 h_{list} \leftarrow h(concat(g.Sid));$ $11 h_t(gram) \leftarrow h(h(g) h_{list});$ $12 nodes.append(h_t(gram));$		
2 foreach $s \in D$ do /* build inv_index */ 3 grams \leftarrow grams (s,q) ; 4 foreach $g \in$ grams do 5 if $g \in inv_gram$ then 6 inv_gram.g.append $(s.Sid)$; 7 else 8 $\left\lfloor inv_gram[g] \leftarrow [Sid]$ 9 foreach $g \in inv_gram$ do /* compute h_t */ 10 $h_{list} \leftarrow h(concat(g.Sid))$; 11 $h_t(gram) \leftarrow h(h(g) h_{list})$; 12 $\left\lfloor nodes.append(h_t(gram))$;		
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10 $h_{list} \leftarrow h(concat(g.Sid));$ 11 $h_t(gram) \leftarrow h(h(g) h_{list});$ 12 $nodes.append(h_t(gram));$		
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11 $h_t(gram) \leftarrow h(h(g) h_{list});$ 12 $nodes.append(h_t(gram));$		
12 $nodes.append(h_t(gram));$		
12 while $len(nodes) > 1$ do /* build MHT */		
14 $new_nodes \leftarrow [];$		
15 foreach $node_l$, $node_r \in nodes$ do		
$16 \qquad node_f \leftarrow h(H(node_l) \mid H(node_r));$		
17 <i>new_nodes.append(node_f);</i>		
18 $nodes \leftarrow new_nodes;$		
19 $S_{inv} \leftarrow sig_{s_k}(nodes);$		
20 return S_{inv} ;		

For each *i*-*list* = $\langle Sid_1, Sid_2, ..., Sid_m \rangle$ of the inverted index (lines 9 to 12 in Algorithm 2), the *DO* computes the *tuple hash* h_t for the *i*-*list* as follows:

$$h_{i-list} = h(Sid_1|Sid_2|\dots|Sid_m), \tag{3}$$

$$h_t(gram - list) = h(h(gram)|h_{i-list}).$$
(4)

For instance, consider the gram "*ai*" and its corresponding list in Table 2. We thus have $h_{ai'',i-list} = h(3|4|5|6)$ and $h_t(ai'') = h(h(ai'')|h_{ai'',i-list})$. Note that $h_t(ai'')$ corresponds to h_1 in Figure 4.

Then, the *DO* computes the MHT of the inverted index. Specifically, the *gram-list* of the inverted index is sorted by grams in alphabetical order, and the MHT construction process of the inverted index is the same as MHT construction for the dictionary. When the MHT is built, the *DO* signs the root hash to generate signature S_{inv} of the inverted index (lines 13–19 of Algorithm 2). We give an example of MHT construction for the inverted index in Figure 4.

After the AIS is constructed as described above, the *DO* uploads dataset *D*, *q*, and two signatures, S_{dic} and S_{inv} , to the server. It is worth noting that in some existing outsourcing database models, the *DO* stores signatures locally (instead of uploading them to the server) and only sends them to users upon request. For example, [3] proposed a solution called $AutoS^3$, in which the *DO* is required to remain active to keep the system running prop-

erly. In contrast, our solution allows the DO to go offline without affecting the database outsourcing services .

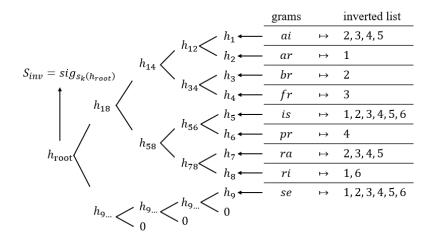


Figure 4. An MHT built on Table 2.

4.1.2. Searching Phase

In this section, we present the process of generating the *VO* for user queries. In general, the *VO* is composed of two components, i.e., the *VO* of the dictionary and the *VO* of the inverted index. First, we provide a brief description of how the server performs approximate string searches with respect to a user query, as summarized in Algorithm 3. Then, we show how to construct *VO* in detail.

Upon receipt of dataset *D*, *q*, and signatures from the *DO*, the server builds the *q*-gram inverted index based on *D* and *q*. Assume the user issues a query *Q* with string s_Q and threshold *k* to the server, which implies that the user wants to search for all strings whose edit distance from s_Q is less than or equal to *k*. After receiving s_Q and *k* from the user, the server calculates the minimum number of shared grams, denoted by τ , between s_Q and a string s_i . According to Lemma 1, we have

$$\tau = \max(s_i, s_Q) - q + 1 - kq. \tag{5}$$

Without loss of generality, τ is set to the minimum value of the above equation, i.e., $\tau = |s_Q| - q + 1 - kq$, as shown in line 2 of Algorithm 3.

Algorithm 3: Search for similar strings in <i>D</i> .
Input: dataset <i>D</i> , inverted index <i>inv_gram</i> , query string <i>s</i> _Q , threshold <i>k</i>
Output: R
1 e - $gram \leftarrow []$; ne - $gram \leftarrow []$; $R \leftarrow \emptyset$; $Cstr \leftarrow \emptyset$;
2 $ au = len(s_Q) - q + 1 - k imes q;$;
s_Q -grams \leftarrow grams (s_Q, q) ;
4 foreach $g \in s_Q$ -grams do
5 if $g \in inv_gram$ then e -gram.append(g);
6 else ne-gram.append(g);
7 $n_Sid \leftarrow counter(inv_gram[e-gram]);$
s foreach $Sid \in n_Sid$ do
9 \lfloor if Sid.num $\geq \tau$ then Cstr.add(D[Sid]);
10 foreach $s \in Cstr$ do
11 $\lfloor \text{ if } ed(s, s_Q) \leq k \text{ then } R.add(D[Sid]);$
12 return R;

The server extracts a set of s_Q -grams of $|s_Q| - q + 1$ grams from $grams(s_Q, q)$, as shown in Line 3 of Algorithm 3. Then, all the *i*-lists corresponding to grams in s_Q -grams are fetched from the inverted index. It is possible that some gram in the s_Q -grams may not have a corresponding *i*-list in the inverted index; hence, we divide the grams in s_Q -grams into the following two cases (lines 4 to 6 of Algorithm 3) :

- If a gram $g \in s_Q$ -grams has a corresponding *i*-list, g is called an *e*-gram;
- If a gram *g* ∈ *s*_Q*-grams* does not have a corresponding *i-list*, *g* is called an *ne-gram*.
 We refer to the *i-lists* that correspond to the *e-gram* as *e-i-lists*.

After that, the server counts the number of occurrences of *Sid* in *e-i-lists* and find those that appears no less than τ times. To count the number of occurrences, there exist many efficient algorithms, e.g., the Heap algorithm, MergeOpt [7], ScanCount, MergeSkip, and DivideSkip [9]. After the counting step, we obtain a set of candidate strings, denoted by the *C-string*, whose *Sid* appears at least τ times in the *e-i-lists*, as shown in lines 7 to 9 in Algorithm 3.

For each string $s_c \in C$ -string, the server computes the edit distance between s_c and s_Q , and those strings whose edit distance from s_Q is less than or equal to k are put into the result set R (as shown in lines 10 and 11 in Algorithm 3). Note that the server can directly use the dynamic programming technique to calculate the edit distance between two strings, or it can use the verification technique [1] or other filter-verification frameworks [27] to speed up the computation.

Example 1. Let us consider an example with s_Q ="arisen", k = 1, and the database D, as shown in Table 1, and its corresponding inverted index, as shown in Table 2. After receiving s_Q and k, the server first computes $\tau = |s_Q| - q + 1 - kq = 6 - 2 + 1 - 1 \times 2 = 3$ and then extracts s_Q grams = {"ar", "ri", "is", "se", "en"}, from which the server obtains e-gram={"ar", "ri", "is", "se"} and ne-gram={"en"}. Next, the server counts the frequency of occurrence of Sid in e-i-lists, and only the strings with Sid \in {1,6} satisfy the requirement of appearing at least τ times. Hence, we have C-string={"arise", "rise"}. Finally, the server computes the edit distance between the strings in the C-string and s_Q . Among them, only the string "arise" satisfies the threshold of edit distance k. Therefore, the result R = {"arise"} is returned to the user.

Next, we show how to construct *VO* with respect to a query result *R*. First, the server builds leaf nodes for all grams in the inverted index. For each *e-gram*, the leaf node contains the gram and its corresponding *i-list*. For each gram that is a neighboring list (the alphabetic predecessor and successor of a gram in the inverted index) of the *ne-gram*, the leaf node contains the gram and the hash value h_{list} of its *i-list*. For each of the remaining grams, the leaf node contains its tuple hash $h_t(t)$ (lines 1 to 12 of Algorithm 4). Note that if a gram in the set of the *ne-gram* is out of the boundary of the inverted list, then the neighbors of that gram only contain one gram. For example, as shown in Table 2, the neighbors of gram "*aa*" and "*sf*" are "*ai*" and "*se*", respectively, whereas the neighbors of gram "*fr*" are "*br*" and "*is*".

Similar to the process of forming *VO* on a traditional MHT, in our solution, the server calculates the hashes of the internal nodes (lines 13 to 18 of Algorithm 4). Similarly, the server builds the MHT for dataset *D* and maintains the information about the *C-string*, as shown in lines 19 to 25 of Algorithm 4.

The formation of *VO* for result set *R* is shown in line 26 of Algorithm 4, from which we can see that *VO* consists of (1) the *e-grams* and their corresponding *e-i-lists*, the neighboring grams of each *ne-gram* and their *i-list* hash values, and the hashes of other nodes; (2) *C-string* set along with their *Sid* and the hashes of other nodes; and (3) the signatures of the dictionary MHT and of the inverted index MHT, respectively. The server sends *R* and *VO* to the user for verification. Note that in scenarios where the user continuously issues queries, the server only needs to send the signatures once to reduce network communication costs.

It is worth mentioning that the elements in *VO* are organized in such a way that the user can easily recalculate the root hash of the dictionary MHT and the root hash of the inverted index MHT. Specifically, the elements in *VO* are organized according to where

they are located in the MHT, and a pair of parentheses is added around elements that share the same parent node in the MHT.

	Al	gorithm	4:	VO	construction
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Input: dataset *D*, inverted index *inv_gram*, *e-gram* list *e_l*, *ne-gram* list *ne_l*, query string s_{O} , k, C-string Cstr, signature S_{dic} and S_{inv} Output: VO 1 $n_ne_l \leftarrow []; i \leftarrow 1; ns \leftarrow [];$ 2 foreach $ne \in ne_l$ do n_ne_l.append(inv_gram.getneighbors(ne)); 3 4 foreach $n \in inv_gram$ do /* compute h_t */ if $n.gram \in e_l$ then 5 $ns.append([n.gram, n.list, flag \leftarrow 1, l \leftarrow 0]);$ 6 else if $n.gram \in n_ne_l$ then 7 $h_{list} \leftarrow h(concat(n.list));$ 8 9 $ns.append([n.gram, h_{list}, flag \leftarrow 1, l \leftarrow 0]);$ else 10 $h_t \leftarrow h(h(n.gram)|h(concat(n.list)));$ 11 $ns.append([h_t, flag \leftarrow 0, l \leftarrow 0]);$ 12 13 while $i < len(inv_gram))$ do /* compute internal node */ 14 $i \leftarrow i \times 2$; **foreach** $n_l, n_r \in ns$ **do** 15 **if** n_1 and n_r same level and both flag 0 **then** 16 $n_f \leftarrow [h(n_l \mid n_r), flag \leftarrow 0, l \leftarrow n_l.l + 1];$ 17 *ns.replace*([*n*₁, *n*_{*r*}], *n*_*f*); 18 19 $VO_{inv} \leftarrow ns; ns \leftarrow [];$ 20 foreach $n \in D$ do /* compute h_t */ if $n \in Cstr$ then 21 $ns.append([n, flag \leftarrow 1, l \leftarrow 0]);$ 22 else 23 $[ns.append([h_t(n), flag \leftarrow 0, l \leftarrow 0]);$ 24 25 $VO_{dic} \leftarrow compute_in_nodes(ns);$ 26 $VO \leftarrow \{VO_{inv}, S_{inv}, VO_{dic}, S_{dic}\};$ 27 return VO;

Example 2. Let us continue with the previous Example 1, where the result $R = \{\text{"arise"}\}$. Referring to Figure 4, it is clear that the neighbors of ne-gram="en" are "br" and "fr", and their corresponding i-list hashes are $h_{br.list} = h(2)$ and $h_{fr.list} = h(3)$, respectively. Similarly, after computing the other necessary hashes, i.e., h_1 , h_6 and h_7 , the server constructs the first part of VO as follows:

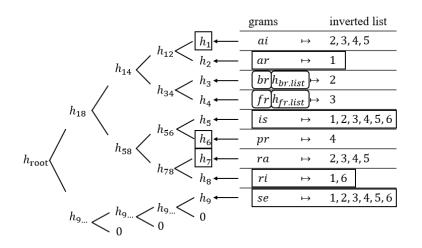
$$VO_{inv} = \{ (((h_1, ("ar", (1)))(("br", h_{br.list})("fr", h_{fr.list}))) \\ ((("is", (1, 2, 3, 4, 5, 6)), h_6)(h_7, ("ri", (1, 6))))) \\ (((("se", (1, 2, 3, 4, 5, 6))))) \}.$$
(6)

Recall that we have C-string={"arise", "rise"}, and after computing h_2 , h_{34} , and h_5 , the server constructs the second part of VO as follows:

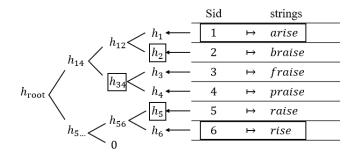
$$VO_{dic} = \{((((1, "arise"), h_2), h_{34})((h_5, (6, "rise"))))\}.$$
(7)

Figure 5 shows the elements that constructed VO_{dic} *and* VO_{inv} *. Finally, having obtained the above components, we have:*

$$VO = \{VO_{inv}, S_{inv}, VO_{dic}, S_{dic}\}.$$
(8)



(a) elements in VO_{inv}



(**b**) elements in VO_{dic}

Figure 5. An example of *VO* constructed by the GS^2 method.

4.1.3. Authentication Phase

Given a user query with s_Q and k, the server conducts query processing and returns the query result set R, along with its corresponding VO to the user. Upon receipt of R and VO, the user verifies whether the received R is correct. Note that the user obtains the DO's public key p_k via a secure channel, such as a certificate authority (CA). We summarize the authentication procedures in Algorithm 5.

At first, based on the *VO* received, the user computes the root hashes for both the dictionary and the inverted index. Specifically, by using VO_{inv} and VO_{dic} , the user can compute the hashes of internal nodes in a recursive manner until the root hashes h'_{root_i} (root hash of the inverted index) and h'_{root_d} (root hash of the dictionary) are obtained (lines 1 and 2 of Algorithm 5). Note that we assume that the *DO* shares with the user some necessary information, such as the hash function, the public key encryption algorithm used by the *DO*, and the parameter *q*, etc. Then, the user decrypts signatures S_{inv} and S_{dic} with p_k , as follows:

$$h_{root_i} = sig_{p_k}(S_{inv}), \tag{9}$$

$$h_{root_d} = sig_{p_k}(S_{dic}). \tag{10}$$

where h_{root_i} and h_{root_d} are compared against h'_{root_i} and h'_{root_d} , respectively. If either $h'_{root_i} \neq h_{root_i}$ or $h'_{root_d} \neq h_{root_d}$, then the user concludes that the query result *R* returned from the server fails the correctness verification, as shown in lines 2 to 6 of Algorithm 5.

Algorithm 5: Query result authentication.
Input: dataset <i>R</i> , query string s_Q , threshold <i>k</i> , verification object <i>VO</i> , public key
<i>pk,</i> gram length <i>q</i>
Output: isPass
1 $h'_{root_i} \leftarrow compute_inv_root(VO.VO_{inv});$
2 $h'_{root_d} \leftarrow compute_dic_root(VO.VO_{dic})$;
$h_{root_i} \leftarrow sig_{p_k}(VO.S_{inv});$
4 $h_{root_d} \leftarrow sig_{p_k}(VO.S_{dic})$;
5 if $h'_{root_i} \neq h_{root_i}$ or $h'_{root_d} \neq h_{root_d}$ then
$6 return \ is Pass \leftarrow False;$
7 Cstr, n_ne_g, e_g, e_g_list \leftarrow parse(VO);
s foreach $g \in grams(s_Q, q)$ do
9 if $g \notin e_g$ and $g's$ neighbors $\notin n_ne_g$ then
10 return is Pass \leftarrow False
11 $Cstr' \leftarrow []; \tau \leftarrow len(s_Q) - q + 1 - k \times q;$
12 $n_sid \leftarrow counter(e_g_list));$
13 foreach $sid \in n_sid$ do
14 $\lfloor \text{ if } sid.num \geq \tau \text{ then } Cstr'.append(Cstr[sid]);$
15 if $Cstr' \neq Cstr$ then return $isPass \leftarrow False$;
16 $R' \leftarrow [$];
17 foreach $s \in Cstr$ do
18 $\lfloor \text{ if } ed(s_Q, s) \leq k \text{ then } R'.append(s);$
19 if $R \neq R'$ then return is Pass \leftarrow False ;
20 return $isPass \leftarrow True;$

On the other hand, if *VO* is correct, then the user compares $grams(s_Q,q)$ with the gram set G_{VO} that consists of all grams appearing in VO_{inv} . For each gram g, where $g \in grams(s_Q,q)$ and $g \notin G_{VO}$ (i.e., g is an *ne-gram*), the user looks for its neighbors in G_{VO} , and the neighbors should be adjacent to each other in *VO*, unless there is only one neighbor. Otherwise, the user is certain that the *R* fails the completeness test.

For each gram g, where $g \in grams(s_Q,q)$ and $g \in G_{VO}$ (i.e., g is an *e-gram*), the user counts the number of occurrences of *Sid* in the *e-i-lists*. For those strings whose *Sid* appears at least $\tau = |s_Q| - q + 1 - kq$ times, the user needs to check whether they appear in VO_{dic} . If any *Sid* appears at least τ times but does not appear in VO_{dic} , then the user is certain that the *R* does not pass the completeness test (lines 7 and 16 in Algorithm 5).

Finally, the user extracts all strings in VO_{dic} and computes the edit distance between each of them and s_Q . If the edit distance of some string is not greater than k, but the string is not included in R, then the user concludes that the result set R fails the completeness test. Conversely, if a string is in R but it is missing during the comparison, then the user is certain that R fails the soundness test (lines 17-21 in Algorithm 5).

Example 3. Let us recall Example 2. We have $s_Q = \{\text{"arisen"}\}, k = 1, R = \{\text{"arise"}\}, VO_{inv}, VO_{dic}, and two signatures <math>S_{inv}$ and S_{dic} . At first, the user recalculates the root hash for VO_{inv} and VO_{dic} , respectively, as follows:

$$\begin{aligned} h'_{root_i} = &h(h(h(h_1|h(``ar''|h(1)))|h(h(``br''|h_{list1})|h(``fr''|h_{list2})))| \\ &h(h(h(h(``se''|h(1,2,3,4,5,6))|0)|0)|0)); \end{aligned}$$

$$h'_{root_{4}} = h(h(h(h(1|"arise")|h_{2})|h_{34})|h(h(h_{5}|h(6|"rise"))|0)),$$
(12)

where h'_{root_i} and h'_{root_d} are compared with the decoded signatures to verify the authenticity of VO.

Then, the user retrieves the set of e-grams and the set of neighbors of ne-gram from VO_{inv} and compares the set of e-grams against gram("arisen",2) in order to obtain ne-gram={"en"}. Based on the ne-gram, the user is certain that the neighboring gram "br" lies before {"en"}, and "fr" lies after {"en"} in the inverted index. Next, the user counts the number of occurrences of Sid and only strings in {1,6} whose occurrences are greater than or equal to $\tau = 3$. Finally, the user retrieves strings {"arise", "rise"} from VO_{dic} , where $Sid = \{1,6\}$, and computes the edit distance between the string $s \in \{$ "arise", "rise" $\}$ and s_Q . Here, only s = "arise" satisfies the threshold k, which is the same as R. Hence, the user confirms that the result R sent from the server is correct.

It is worth mentioning that by modifying the computation of τ (note that an example computing method can be found in [27]), our GS^2 model can be applied to other threshold-based similarity measures for strings, such as Hamming distance, Edit similarity, Jaccard similarity, Cosine similarity, and Dice similarity. For example, for the Hamming distance, τ can be calculated by $\tau = |s_Q| - q + 1 - k * q$, and the calculation of τ values for other distance measures can be found in [27].

 GS^2 is not a very cost-effective approach in terms of network communication because the size of the *VO* generated is large, resulting in a higher transmission cost. For most resource-constrained clients, especially mobile devices or IoT devices, the network bandwidth and battery are limited. Therefore, it is important to reduce the communication overhead between the server and the users. To address this issue, in the next section, we design an optimization scheme, which can significantly reduce the *VO* size.

4.2. Optimization Method for GS^2

In this section, we introduce an optimization scheme named GS^2 -opt. In GS^2 , VO_{inv} and VO_{dic} are the two components that take up the majority of the *VO* size, since the length of the *i*-list in VO_{inv} may be excessively long, and the number of *C*-strings may be large for a large string dataset.

In our GS^2 -opt model, we reduce the size of VO_{inv} and VO_{dic} by using the following strategies:

- 1. We employ a length-filtering technique to reduce the size of *VO*_{inv} and *VO*_{dic}.
- 2. We propose the *representative grams* to reduce the size of *VO*_{*inv*} at the cost of a slight increase in the size of *VO*_{*dic*}.
- 3. We design a solution to deal with the case where some parts of the result are empty during authentication.
- 4. We use compression techniques to further reduce the size of *VO*.

Note that the goal of the above optimization strategies is to shrink the *VO* size, and that some of them are not suitable for similarity functions other than the edit distance. Below, we explain these strategies in detail.

4.2.1. Length Filtering

Long strings tend to cover many grams, which causes them to frequently sneak into *C-strings*, but they are obviously different from s_Q . Therefore, we use a length-filtering technique [28] to further weed out strings that do not satisfy the edit distance constraint. The application of the length-filtering technique is based on the following lemma:

Lemma 2. If $ed(s_i, s_Q) \le k$, then their lengths cannot differ by more than k, i.e.,

$$ed(s_i, s_Q) \le k \Rightarrow \left\| |s_i| - |s_Q| \right\| \le k.$$
(13)

Proof of Lemma 2. The proof for Lemma 2 is straightforward. An insertion or deletion edit operation changes the string length by no more than 1, and the substitution operation does not change the string length at all. Therefore, within *k* edit operations, the string length changes by no more than *k*. \Box

To authenticate the strings pruned by length filtering, we modify some of the steps in GS^2 . Below, we will explain the details of these changes. Specifically, during the preprocessing phase, strings are sorted by length at first and then sorted in alphabetical order. For example, given a set of strings in Table 3, after sorting, they are listed in Table 4.

Table 3. Strings sorted in alphabetical order.

Sid	String
1	arise
2	braise
3	fraise
4	praise
5	raise
6	rise

Table 4. Strings sorted by length at first, then sorted in alphabetical order.

Sid	Strings
1	rise
2	arise
3	raise
4	braise
5	fraise praise
6	praise

Additionally, before building the MHT for the inverted index, the *DO* sorts each of the *i*-*list*, builds an MHT for each *i*-*list* in the inverted index (as shown in Figure 6), and takes the root hash as the *i*-*list* digest:

$$h_{i-list} = h_r. (14)$$

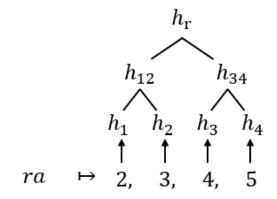


Figure 6. An Example of MHT for an *i*-list.

Figure 6 shows an example of building an MHT for an *i-list* of "*ra*", and its *tuple hash* is computed as follows:

During the searching phase, before constructing the *VO*, the server calculates the desired range of length $[|s_Q| - k, |s_Q| + k]$ and filters out all the *C*-strings that fall outside

this range. To facilitate authentication, when building the MHT for the dictionary, the strings that are immediately adjacent to the range borders are included in VO_{dic} . On the other hand, when building the MHT for the inverted index, for each *gram-list*, the server constructs an MHT and uses digests to represent the *Sid* of strings whose length is not in the range (except two boundary strings that are not in the range). In this process, we also introduce the similar concept of *buddy inclusion* [2] to further reduce the *VO* size.

In the verification phase, the user verifies that the length of the first string in the *C*-string is not greater than $|s_Q| - k - 1$ and that the length of last string in the *C*-string is not less than $|s_Q| + k + 1$. The user also checks that the range of *Sid* of each *e*-*i*-list covers the *Sid* of the boundary strings and that all of the *Sids* in the *e*-*i*-list are continuous. With the exception of the steps described above, all other procedures are the same as those in GS^2 .

4.2.2. Representative Grams

As discussed in [7], the frequency distribution of words in real-life datasets is often heavily skewed, with a significant portion of words appearing much more frequently than others. This phenomenon is also observed in the frequency distribution of string grams. For example, in the dataset *authors* obtained from DBLP (http://dblp.uni-trier.de/xml/ (accessed on 6 July 2022)), it has been observed that nearly one-third of string grams appear much more frequently, with a frequency over ten times higher than the other one-third. Therefore, in an inverted index, some grams tend to be associated with long *i-lists*. On the other hand, grams with a long *i-list* have a higher probability of becoming an *e-gram* in a query than grams with a short *i-list*. Therefore, if a query string s_Q covers some frequently occurring grams, it is likely that many strings that are very different from s_Q will be included in the *C-string*, leading to a large *VO*.

To address this problem, we propose the use of *representative grams*. The basic idea is that instead of returning all the *e-grams* and the corresponding *i-lists*, the server only returns a few grams that represent the grams covered by s_Q . If a string covers grams that appear multiple times, its *Sid* may appear several times in the same gram. To deal with this problem, we design a new format for the *gram-list*, i.e., a triplet (*gram*, *t*, [*list*]), where the element *t* denotes the number of times *Sid* occurs.

An example string set is given in Table 5, and the new format of the inverted index built on the string set can be seen in Table 6. In this example, the notation " $(ar,2) \mapsto 1, 2$ " means that the gram "ar" occurs twice in both strings with Sid = 1 and Sid = 2. Hence, the *tuple hash* is calculated as follows:

$$h_t(``ar, 2'') = h(h(ar|2)|h_{list}).$$
(16)

Table 5. An example string set.

Sid	Strings
1	arar
2	arari
3	rari

Table 6. Example of 2-gram inverted index with occurrence time.

Grams	t		Inverted List
(ar ,	1)	\mapsto	3
(ar ,	2)	\mapsto	1,2
(ra ,	1)	\mapsto	1,2,3
(<i>ri</i> ,	1)	\mapsto	2,3

In our previous description, if a string is considered a *C*-string, it must share at least $\tau = |s_Q| - q + 1$ grams with s_Q . In *GS*²-opt, we change this condition so that the sum of t for a string's grams must be greater than or equal to τ . This modification allows us to more

accurately evaluate the similarity of strings that may contain the same grams that appear multiple times.

For ease of discussion, we define the representative grams (i.e., those grams) to be included in VO_{inv}) as *r*-grams, the corresponding *i*-list as *r*-*i*-list, the sum of all *t* values for any set of grams as *st*, the sum of all *t* values for *e*-grams as *st_e*, the sum of all *t* values for *r*-grams as *st_r*, the subset of *e*-grams as *sub-e*-grams, and the sum of all *t* values for a string *s* in a certain *sub-e*-gram (i.e., how many times the *Sid* of the string *s* appears in the *sub-e*-gram) as *st_s*. Then, we have the following lemma:

Lemma 3. The server only needs to return those r-grams where $st_r \ge st_e - \tau + 1$ and the length of *i*-lists is the shortest among the e-grams. All strings with $st_s \ge \tau - (st_e - st_r)$ are included in the C-string. This allows us to reduce the number of *i*-lists included in VO_{inv} .

To illustrate how this method works, let us consider a string $s \in D$ that is similar to the query string s_Q . According to the previous similarity condition, s must share at least τ grams with s_Q . This indicates that for *e-grams*, st_s must be greater than or equal to τ . For any subset of the *e-grams* with $st = st_e - i$, if a string s is similar to s_Q , then its st_s must be bigger than or equal to $\tau - i$, $i \in [1, \tau - 1]$. If we set $i = \tau - 1$, it follows that for any subset of *e-gram* with the sum of t values, $st = st_e - \tau + 1$, and the *Sid* of s must appear at least once in their corresponding *i-list*. This means that any subset of *e-grams* with $st \ge st_e - \tau + 1$ can be considered a *representative gram* with respect to user query Q.

Next, we provide a formal proof of the completeness of the proposed method (the soundness proof is straightforward and is omitted).

Proof of Lemma 3. Given a string $s \in D$, where *s* contains |s| - 1 + q grams, there must be three cases, as shown below:

- Case 1: ∃ gram ∈ r-gram. In this case, if st_s ≥ τ − (st_e − st_r), then s will appear in the *C*-string, and the user will check the distance between s and s_Q to determine whether s should be included in *R*. On the other hand, if st_s < τ − (st_e − st_r), then we can infer that for the entire *e*-gram, we have st_s < τ. Therefore, it is impossible that s is similar to s_Q. The VO authentication will fail if there is any violation of these inequalities.
- Case 2: (∃ gram ∈ e-gram) ∧ (∀ gram ∉ r-gram). In this case, we have st_e − st_r ≤ τ − 1, which means that for the entire *e-gram*, we have st_s ≤ τ − 1. Therefore, it is impossible for *s* to be similar to s_O.
- Case 3: \forall gram \notin e-gram. In this case, s is not similar to s_0 .

Therefore, the server only needs to include the *r-i-list*, which is a subset of the *e-i-list*, into *VO* to guarantee the completeness of the results. It is worth noting that there are many possible combinations of *r-grams*, and the value of $st_r = st_e - \tau + 1$ may not be the optimal value to fit all scenarios. As such, the specific number of *r-grams* to be included in *VO* should be determined according to the dataset and the user query.

4.2.3. Dealing with Empty Results

When the query result is empty, the user only needs to authenticate its completeness. Specifically, we have the following lemma:

Lemma 4. For a query with s_Q and k, if $\alpha > k \times q$, where α is the number of ne-grams, then R is an empty set.

In this case, it is not necessary to include any *i*-*list* in *VO* because the result set is already known to be empty.

Proof. An insert or substitution operation in the edit distance creates at most q grams that contain the inserted or substituted character, while a deletion operation creates at most q - 1

grams. Given *k* edit operations, at most, *kq* grams can be created. The number of *ne-gram* α represents the number of grams that need to be created if a string $s \in D$ is transformed into s_0 . Therefore, if $\alpha > kq$, there are no strings in *D* that can be transformed into s_0 .

Thus, if a user query meets this condition (i.e., the number of the *ne-gram* is greater than $k \times q$), the server can include only the neighbours of *ne-grams* in VO_{inv} with the digests that are necessary for computing the root hash, omitting all *e-grams* and their associated *e-i-lists*. The authentication phase in this case is much easier than the authentication in GS^2 , and the size of *VO* is also reduced dramatically.

4.2.4. VO Compression

By analyzing the contents of *VO*, we found that English letters and digits, along with a few symbols such as "(", ")" and ",", are the most frequent characters appearing in *VO*, especially in VO_{inv} (in which the *e-i-list* and digest of other *gram-list* are the main components).

To cut down the communication overhead in *VO* transmission, we use the Lempel– Ziv–Markov chain algorithm (LZMA) [29] to compress *VO*. LZMA is a dictionary-based compression method, and it uses a range encoder to compress the data. After *VO* construction, the server uses LZMA to compress *VO* and then sends it to the user. Upon receipt of the compressed *VO*, the user decompresses it before performing the authentication procedure.

5. Security Analysis

Given a query string s_Q with threshold k, let R be the set of strings in D that are similar to s_Q and let R' be the set of similar strings returned by the server. An untrustworthy server may attempt to cheat the user by violating the *soundness* or *completeness* of the query result, as shown below.

- 1. *Violation of soundness:* following the definition in Section 3.2, the server returns R', where $\exists s \in R' \land s \notin R$. This means that the server is returning some strings that do not belong D or are not similar to s_0 .
- 2. *Violation of completeness:* the server returns R', where $\exists s \in R \land s \notin R'$. This means that the server is not returning all of the strings that are similar to s_Q .

Soundness. To facilitate the analysis, we divide this situation into two situations: (1) $\exists s \in R' \land s \notin R \land s \notin D$, and (2) $\exists s \in R' \land s \notin R \land s \in D$. For the first case, it is easy to detect this during the authentication procedures as the digest of *s* is different from the original strings, resulting in the root hash calculated by the user not matching the one decoded from the *DO*'s signature. Hence, the user is able to catch this violation during the step of comparing the root hashes.

For the second case, the server returns some strings that are not similar to s_Q . Apparently, this will also be identified by the user in the final step of the authentication procedure, when the user compares strings in R' with the strings (i.e., the *C-string*) that meet the similarity threshold.

Completeness. When a string $s \in R$ is excluded by the server from R', the following cheating actions may be taken by the server to convince the client to accept R':

- Case 1: The server claims that grams shared by both *s*_O and *s* do not exist.
- Case 2: The server does not deny the existence of the grams but forges the corresponding *i-list* to exclude the *Sid* of *s*.
- Case 3: The server returns the correct grams and their corresponding *i-lists* but omits *s* from the *C-string*.
- Case 4: The server only omits *s* from *R*′ but provides a correct *VO*.

For Case 1, the server omits the grams which it claims from VO_{inv} to create two sibling neighbours in order to deceive the user. However, no matter how the server alters the grams in the original inverted index, it is impossible for the root hash of the inverted index MHT computed by the user to match the one decoded from the *DO*'s signature. The user can detect this case during the authentication phase. For Case 2, this action will also change the root hash values of the inverted index MHT and will be detected by the user, similar to Case 1. For Case 3, with the correct inverted index, the user can filter out the *C-string*'s *Sid* on their own. If *s* does not appear in VO_{dic} , the user will conclude that *R* fails the completeness test. Finally, for Case 4, with the correct *C-string* set, the user can verify all strings in the *C-string* and compare them against those in *R'*. If a string *s* is similar to s_Q but does not appear in *R'*, then the user will find that *R'* is incorrect.

6. Experiment

6.1. Experiment Set-Up

Datasets and query workload. We used two real-world datasets in the experiments. The first one was the Frequently Occurring Surnames dataset (denoted by *LastName*) provided by the United States Census Bureau (https://www.census.gov/topics/population/genealogy/data/1990_census/1990_census_namefiles.html (accessed on 6 July 2022)), which contains 88,799 last names with an average length of 6.83 characters and a maximum length of 13 characters. The second one is the *author* dataset downloaded from DBLP (http://dblp.uni-trier.de/xml/ (accessed on 6 July 2022)), which includes 2,845,839 researcher names with an average length of 14.39 characters and a maximum length of 67 characters.

We built a query workload of single strings by randomly selecting 25 strings from each of the datasets, and we also artificially generated 25 strings. The reason we used synthetic strings was to ensure that the *ne-gram* exists. The results presented in the experiment part are the average over 10 trials.

Parameter setting. The parameters included (1) *q* for the *q*-gram inverted index and (2) the edit distance threshold *k*, as shown in Table 7, where the default values are in bold.

 Table 7. Parameter setting.

Parameter	Range of Values
q	2 , 3, 4, 5, 6
k	1, 2, 3, 4, 5

Experimental environment. We implemented both the GS^2 and GS^2 -opt methods in Python and employed the *SHA256* hash function from the Crypto Library. All the experiments were conducted on a machine with a 3.10GHz CPU and 128GB main memory, running an Ubuntu 18.04.3 LTS operating system.

To evaluate the performance of our methods, we adopted three metrics, i.e., *VO* construction time, *VO* size, and *VO* verification time, as described in Section 3.3. Note that the preprocessing time at the *DO*'s side and the network communication between the *DO* and the server are one-time costs that can be amortized over multiple queries and were thus not considered in our experiment. Meanwhile, we investigated the performance of each optimization technique in GS^2 -opt.

6.2. VO Construction Time

In this section, we measured the *VO* construction time at the server side. Figure 7 depicts the average *VO* construction time with various lengths of gram *q* and threshold *k*.

From Figure 7a, we can see that as q increases, the construction time of the GS^2 method first decreases slightly. The main reason is that as q increases, the length of *e-i-lists* in the VO_{inv} diminishes, thus reducing the time spent in counting *Sid*. As q increases further, however, the construction time goes up steadily. This is because the number of grams in the inverted index increases, leading to (1) increases in the time spent searching for the *ne-gram* and (2) increases in the time for the construction of VO_{inv} . Furthermore, we can see that when q=2, the processing time of GS^2 -opt is significantly larger than that of GS^2 . This is because the *i-lists* in the VO_{inv} part are relatively long, and GS^2 -opt needs to take a long time to build MHTs for each *i-list*. In other cases, the performance trends of GS^2 and GS^2 -opt are basically consistent.

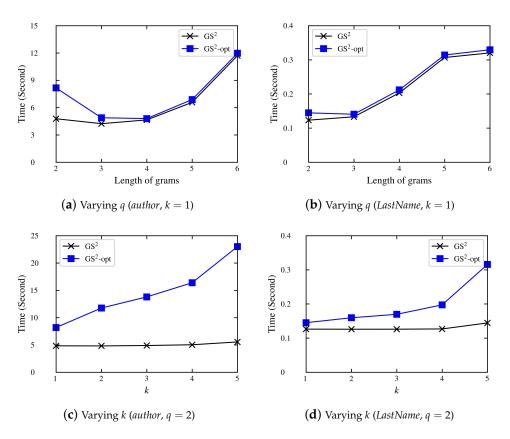


Figure 7. VO construction time versus q and k, respectively.

Figure 7b exhibits a similar trend to Figure 7a, but there are two differences. Firstly, the construction time of GS^2 always increases with q. The reason is that the *LastName* dataset is relatively small, and the time necessary to count the *Sid* is only a small proportion of the time taken to construct the whole *VO*. Therefore, the reduction in the *Sid* counting time imposes a negligible impact on the authentication time. Secondly, when q = 2, the gap between the two methods is not as significant as in Figure 7a. This is because the *LastName* dataset is relatively small and the *i-lists* in *VO*_{*inv*} are much shorter than those in the *author* dataset.

Next, we investigated the impact of edit distance threshold k on VO construction time. The results are presented in Figure 7c,d. From Figure 7c, we can see that for the *author* dataset, the VO construction time of the GS^2 method increases slightly with k. The rationale is that (1) the majority of the time cost is spent on constructing VO_{dic} , and (2) the number of *C-strings* has little impact on the construction time of VO_{dic} . We can see that in GS^2 -opt, as k increases, the construction time also increases. This is due to the fact that constructing the VO_{inv} part uses the majority of the total construction time, and as k increases, the number of *Sid* included in the *e-i-list* becomes longer, leading to an increase in the construction time also increases accordingly. We observe a similar performance trend on *LastName* dataset (Figure 7d).

6.3. VO Size

In this section, we investigate the impact of *q* and *k* on *VO* size. The experimental results are shown in Figure 8.

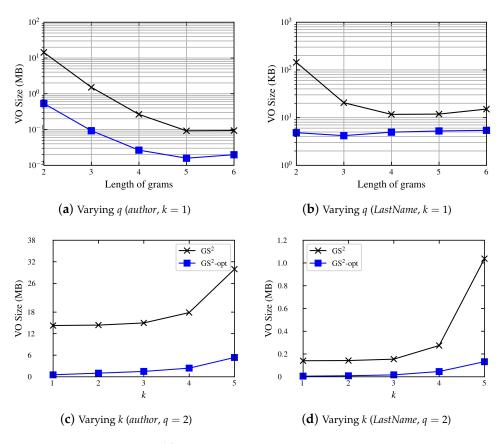


Figure 8. *VO* size versus *q* and *k*.

From Figure 8a, we can see that as q increases, the VO size of the GS^2 method decreases dramatically at first. This is because when the length of grams grows, the number of grams will increase dramatically, and the average length of *i*-*lists* in the invert index will decrease. Therefore, the number and average length of *e*-*i*-*lists* will decrease accordingly, leading to a decrease in the number of *C*-*strings* and both VO_{dic} and VO_{inv} . As q grows from 5 to 6, the VO size shows an increasing trend. The reason for this is that τ decreases as qgrows, resulting in an increase in the number of *C*-*strings* and the VO_{dic} size as well. From Figure 8a, we can also see that the performance trend of GS^2 -opt is similar to that of GS^2 , but the size of VO generated by GS^2 -opt is much smaller than that generated by GS^2 . The situation is similar in Figure 8b.

As shown in Figure 8a, when the length of a gram is large, e.g., q=5, the *VO* size will decrease drastically. However, the limitation of the threshold will be more strict with a greater gram length. If $\tau = |s_Q| - q + 1 - kq \le 0$, then the proposed methods will crash, and hence, the significant interval of *k* should be set to $[0, \frac{|s_Q|+1}{q} - 1]$. Therefore, during the preprocessing phase, the length of grams should be set to a reasonable value.

Figure 8c,d depicts the impact of k on VO size. We can see that in the GS^2 method, as k increases, the growth rate of the VO size becomes faster. This is because (1) as k increases, the volume of VO_{inv} remains constant, while the volume of the VO_{dic} increases rapidly; (2) VO_{dic} initially only accounts for a small proportion of VO, so its growth is not significant for the overall trend. However, when VO_{dic} accounts for a larger proportion, its increase imposes a greater impact on the trend of VO size. The trend of GS^2 -opt is similar to that of GS^2 despite the fact that both VO_{inv} size and VO_{dic} size increase in the GS^2 -opt method. However, the VO volume of GS^2 -opt is significantly smaller than that of GS^2 .

6.4. VO Verification Time

In this section, we study the impact of q and k on VO verification time. In general, the verification of VO involves several steps: (1) the decryption of signatures, (2) the verification of the *e-gram* and *ne-gram*, (3) the calculation of the edit distance for a *C-string*, (4) the verification of VO_{dic} , (5) the verification of VO_{inv} , and (6) re-counting the *Sid*. The total verification time is the sum of the time required for each of these six steps. It is worth noting that our experimental results show that time costs of the first three are negligible. Therefore, we only report the time consumed by the last three parts in our experiments.

When *q* increases, the verification time in the GS^2 method decreases at first and then increases, as shown in Figure 9a. This is because: (1) the size of VO_{inv} part decreases, and the time needed to verify it also decreases; (2) the time needed to re-count *Sid* decreases; (3) the time needed to verify VO_{dic} first decreases and then increases. The change in the verification time of GS^2 -opt is similar, but due to a greater decrease in the time necessary for the verification of VO_{inv} , the overall verification time shows a decreasing trend. For the *LastName* dataset, as shown in Figure 9b, a similar trend is observed.

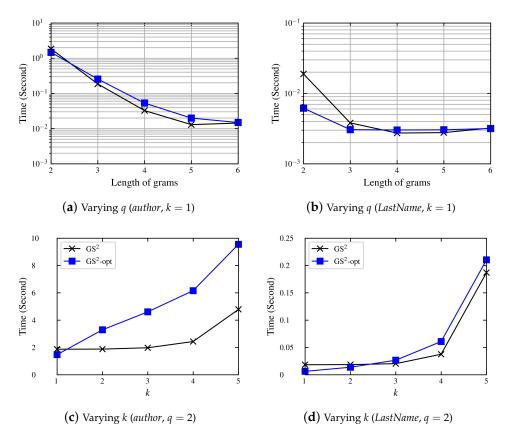


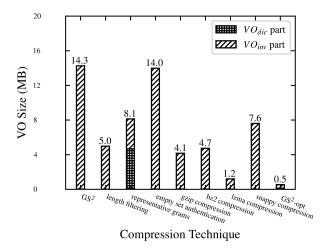
Figure 9. *VO* Verification time versus *q* and *k*.

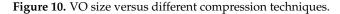
From Figure 9c,d we can see that when k is relatively small, the verification time of GS^2 on both datasets remains steady. However, when k is greater than 4, the verification time increases sharply. This is because the VO_{dic} size increases significantly with k, resulting in a significant increase in verification time. Since the proportion of time spent on validating VO_{dic} is very small at first, the increase in verification time has a marginal impact on the overall authentication time. On the other hand, as the proportion of VO_{dic} increases, it increases a greater impact on the overall authentication time. Note that in GS^2 , VO_{inv} does not change with k, so the time needed to verify it and recount *Sid* remains largely unchanged.

However, for the GS^2 -opt method, the size of VO_{inv} grows with an increase in k, and the time needed to verify it and recount *Sid* increases accordingly. Therefore, the growth of GS^2 -opt is more significant at first.

6.5. Performance of Techniques in GS²-opt

In this section, we investigate the effect of each component in GS^2 -opt on VO size, and the experimental results are shown in Figure 10. It is worth noting that the VO size displayed in bars 2 to 8 in the figure were obtained by applying each optimization technique individually on the GS^2 approach. The last bar shows the combined effect of these four techniques, and the compression scheme used is LZMA.





The first bar depicts the *VO* size of GS^2 . The second bar shows the *VO* size after applying the *length-filtering* technique on GS^2 . The third bar is the *VO* size after using the *representative grams* technique alone, which reduces the size of VO_{inv} and increases the size of VO_{dic} significantly. The fourth bar presents the result of using the *empty result improvement* technique, the optimization effects of which seem to be ordinary, and this is because the number of strings that meet the requirement for this technique in our query set is relatively small. However, for queries that meet the criteria, the *VO* size can be reduced by approximately 99.96% in the test. The next four bars show the *VO* sizes after compressing using different compression techniques, where snappy performs quite ordinarily and LZMA performs the best. The last bar gives the result of combining the previous four techniques, where the VO size is reduced to only 3.73% of its original size.

7. Conclusions

In this paper, we focused on the problem of ensuring the correctness of the results of approximate string searches on outsourced text databases in the context of cloud computing. To solve this problem, we have designed an authenticating structure for a *q*-gram-based inverted index and proposed an efficient method named GS^2 , with which we can efficiently build *VO* and verify the correctness of the query result. Furthermore, we proposed an optimal method named GS^2 -opt for more efficiency in terms of communication overhead. Extensive experiments on real datasets confirmed the effectiveness of our GS^2 and GS^2 -opt methods. In our future work, we will consider approximate string search result authentication with a freshness guarantee.

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