Zips: Mining Compressing Sequential Patterns in Streams

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ABSTRACT

We propose a streaming algorithm, based on the minimal description length (MDL) principle, for extracting non-redundant sequential patterns. For static databases, the MDL-based approach that selects patterns based on their capacity to compress data rather than their frequency, was shown to be remarkably effective for extracting meaningful patterns and solving the redundancy issue in frequent itemset and sequence mining. The existing MDL-based algorithms, however, either start from a seed set of frequent patterns, or require multiple passes through the data. As such, the existing approaches scale poorly and are unsuitable for large datasets. Therefore, our main contribution is the proposal of a new, streaming algorithm, called Zips, that does not require a seed set of patterns and requires only one scan over the data. For Zips, we extended the Lempel-Ziv (LZ) compression algorithm in three ways: first, whereas LZ assigns codes uniformly as it builds up its dictionary while scanning the input, Zips assigns codewords according to the usage of the dictionary words; more heavily used words get shorter code-lengths. Secondly, Zips exploits also non-consecutive occurences of dictionary words for compression. And, third, the well-known space-saving algorithm is used to evict unpromising words from the dictionary. Experiments on one synthetic and two real-world large-scale datasets show that our approach extracts meaningful compressing patterns with similar quality to the state-of-the-art multi-pass algorithms proposed for static databases of sequences. Moreover, our approach scales linearly with the size of data streams while all the existing algorithms do not.

1. INTRODUCTION

Mining frequent patterns is an important research topic in data mining. It has been shown that frequent pattern mining helps finding interesting association rules, or can be useful for classification and clustering tasks when the extracted

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patterns are used as features [1]. However, in descriptive data mining the pattern frequency is not a reliable measure. In fact, it is often the case that highly frequent patterns are just a combination of very frequent yet independent items.

There are many approaches that address the aforementioned issues in the literature. One of the most successful approaches is based on data compression which looks for the set of patterns that compresses the data most. The main idea is based on the *Minimum Description Length Principle* (MDL) [5] stating that the best model describing data is the one that together with the description of the model, it compresses the data most. The MDL principle has been successfully applied to solve the redundancy issue in pattern mining and to return meaningful patterns [4, 6].

So far most of the work focussed on mining compressing patterns from static datasets or from modestly-sized data. In practice, however databases are often very large. In some applications, data instances arrive continuously with high speed in a streaming fashion. In both cases, the algorithms must ideally scale linearly with the data size and be able to quickly handle fast data updates. In the streaming case, the main challenge is that whole data cannot be kept in memory and hence the algorithm has to be single-pass. None of the approaches described in the literature scales up to the arbitrarily large data or obey the single-pass constraint.

In this work, we study the problem of mining compressing sequential patterns in a data stream where events arrive in batches, e.g. like stream of tweets. We first introduce a novel encoding that encodes sequences with the help of patterns. Different from the encodings used in recent work [2, 3, 7], the new encoding is online which enables us to design online algorithms for efficiently mining compressing patterns from a data stream. We prove that there is a simple algorithm using the proposed online encoding scheme and achieving a near optimal compression ratio for data streams generated by an independent and identical distributed source, i.e. the same assumption that guarantees the optimality of the *Huffman* encoding in the offline case [9].

Subsequently, we formulate the problem of mining compressing patterns from a data stream. Generally, the data compression problem is NP-complete [11]. Under the streaming context with the additional single pass constraint, we propose a heuristic algorithm to solve the problem. The proposed algorithm scales linearly with the size of data. In the experiments with one synthetic and two real-life large-scale datasets, the proposed algorithm was able to extract mean-

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ingful patterns from the streams while being much more scalable than the-state-of-the-art algorithms.

2. RELATED WORK

The SubDue system [6] is the first work exploiting the MDL principle for mining useful frequent subgraphs. In the field of frequent itemset mining, the well-known Krimp algorithm [4] was shown to be very good at solving the redundancy issue and at finding meaningful patterns.

The MDL principle was first applied for mining compressing patterns in sequence data in [2, 3] and in [7]. The GoKrimp algorithm in the former work solved the redundancy issue effectively. However, the first version of the GoKrimp algorithm [2] used an encoding scheme that does not punish large gaps between events in a pattern. In an extended version of the GoKrimp algorithm [3] this issue is solved by introducing gaps into the encoding scheme based on Elias codes. Besides, a dependency test technique is proposed to filter out meaningless patterns. Meanwhile, in the latter work the SQS algorithm proposed a clever encoding scheme punishing large gaps. In doing so, the SQS was able to solve the redundancy issue effectively. At the same time it was able to return meaningful patterns based solely on the MDL principle.

However, a disadvantage of the encoding defined by the SQS algorithm is that it does not allow encoding of overlapping patterns. Situations where patterns in sequences overlap are common in practice, e.g. message logs produced by different independent components of a machine, network logs through a router etc. Moreover, neither the GoKrimp algorithm nor the SQS algorithm were intended for mining compressing patterns in data streams. The encodings proposed for these algorithms are *offline encodings*. Under the streaming context, an offline encoding does not work because of the following reasons:

- 1. Complete usage information is not available at the moment of encoding because we don't know the incoming part of the stream
- 2. When the data size becomes large, the dictionary size usually grows indefinitely beyond the memory limit. Temporally, part of the dictionary must be evicted. In the latter steps, when an evicted word enters the dictionary again we loose the historical usage of the word completely.
- 3. Handling updates for the offline encoding is expensive. In fact, whenever the usage of the word is updated, all the words in the dictionary must be updated accordingly. On one hand, this operation is expensive, on the other hand, it is impossible to update the compression size correctly for the case that part of the dictionary has been evicted.

In contrast to these approaches, the Zips algorithm proposed in this work inherits the advantages of both state-ofthe-art algorithms. It defines a new *online encoding* scheme that allows to encode overlapping patterns. More importantly, under reasonable assumptions, it provably scales linearly with the size of the stream making it the first work in this topic being able to work efficiently on very large datasets. Our work is tightly related to the *Lempel-Ziv*'s data compression algorithm [9]. However, since our goal is to mine interesting patterns instead of compression, the main differences between our algorithm and data compression algorithms are:

- 1. Data compression algorithms do not aim to a set of patterns because they only focus on data compression.
- 2. Encodings of data compression algorithms do not consider important patterns with gaps. The *Lempel-Ziv* compression algorithms only exploit repeated strings (consecutive subsequences) to compress the data while in descriptive data mining we are mostly interested in patterns interleaved with noises and other patterns.

3. DATA STREAM ENCODING

In this work, we assume that events in a data stream arrive in batches. This assumption covers broad types of data streams such as tweets, web-access sequences, search engine query logs, etc. This section discusses online encodings that compress a data stream by a set of patterns. For education reasons, we first start with the simplest case when only singletons are used to encode the data. The generalized case with non-singletons is described in the next subsection.

3.1 Online encoding using singletons:

We discuss an online encoding that uses only singletons to compress the data. Since this encoding does not exploit any pattern for compress the data, we consider the representation of the data in this encoding as an uncompressed form of that data. Let $\sum = \{a_1, a_2, \dots, a_n\}$ be an alphabet containing a set of characters a_i , the online data encoding problem can be formulated as follows:

DEFINITION 1 (ONLINE ENCODING PROBLEM). Let A be a sender and let B be a receiver. A and B communicate over some network, where sending information is expensive. A observes a data stream $S_t = b_1b_2\cdots b_t$. Upon observing a character b_t , A needs to compress the character and transmit it to the receiver B, who must be able to uncompress it. Since sending information on the network is expensive, the goal of A and B is to compress the stream as much as possible to save up the network bandwidth.

In the offline scenario, i.e. when S_t is finite and given in advance, one possible solution is to first calculate the frequency of every item a (denoted as f(a)) of the alphabet in the sequence S_t . Then assign each item a a codeword with length proportional to its entropy, i.e. $-\log f(a)$. It has been shown that when the stream is independent and identically distributed (i.i.d) this encoding, known as the *Huffman* code in the literature, is optimal [9]. However, in the streaming scenario, the frequency of every item a is unknown and the codeword of a must be assigned at the time a arrives and B must know that codeword to decode the compressed item.

We propose a simple solution for Problem 1 as follows. First, in our proposed encoding we need codewords for natural numbers. This work uses the *Elias Delta* code [8] denoted E(n) to encode the number n. The Elias was chosen because it was provable as near optimal when the upper bound on n is unknown in advance. The length of the codeword E(n) is $\lfloor \log_2 n \rfloor + 2 \lfloor \log_2 (\lfloor \log_2 n \rfloor + 1) \rfloor + 1$ bits.

A first notifies B of the size of the alphabet by sending $E(|\sum|)$ to B. Then it sends B the dictionary containing all



Figure 1: A first sends B the alphabet *abcd* then it sends the codewords of gaps between consecutive occurrences of a character. B decodes the gaps and uses them to refer to the characters in part of the stream having been already decoded. The reference stream is E(3)E(1)E(6)E(5)E(2)E(4).

characters of the alphabet \sum in the lexicographical order. Every character in the dictionary is encoded by a binary string with length $\lceil \log_2 | \sum | \rceil$. Finally, when a new character in the stream arrives A sends the codeword of the gap between the current and the most recent occurrence of the character. When B receives the codeword of the gap it decodes the gap and uses that information to refer to the most recent occurrence of the character which has been already decoded in the previous step. Since the given encoding uses a reference to the most recent occurrence of a word to encode its current occurrence we call this encoding the *reference encoding* scheme. We call the sequence of encoded gaps sent by A to B the *reference stream*.

EXAMPLE 1. Figure 1 shows an example of a reference encoding scheme. A first sends B the alphabet in lexicographical order. When each item of the stream arrives A sends B the codeword of the gap to its most recent occurrence. For instance, A sends E(3) to encode the first occurrence of b and sends E(1) to encode the next occurrence of b. The complete reference stream is E(3)E(1)E(6)E(5)E(2)E(4).

Let O be a reference encoding; denote $L^{O}(S_t)$ as the length of the data including the length of the alphabet. The average number of bits per character is calculated as $\frac{L^{O}(S_t)}{t}$. The following theorem shows that when the data stream is generated by an *i.i.d* source, i.e. the same assumption guaranteeing the optimality of the Huffman code, the reference encoding scheme approximates the optimal solution by a constant factor with probability 1.

THEOREM 1 (NEAR OPTIMALITY). Given an i.i.d data stream S_t , let H(P) denote the entropy of the distribution of the characters in the stream. If the Elias Delta code is used to encode natural numbers then:

$$Pr\left(\lim_{t\to\infty}\frac{L^O(S_t)}{t} \le H(P) + \log_2(H(P)+1) + 1\right) = 1$$

PROOF. Due to space limit the complete proof of this theorem is available in an extension version¹. \Box

It has been shown that in expectation the lower bound of the average number of bits per character of any encoding scheme is H(P) [9]. Therefore, a corollary of Theorem 1 is that the *reference encoding* approximates the optimal solution by a constant factor $\alpha = 2$ plus one extra bit.

In the proof of Theorem 1 we can also notice that the gaps between two consecutive occurrences of a character represent the usage of the character in the offline encoding because in expectation the gap is proportional to the entropy of the character, i.e. $-\log p_i$. This property is very important because it provides us with a lot of conveniences in designing an effective algorithm to find compressing patterns in a data stream. In particular, since gaps can be calculated instantly without the knowledge about the whole data stream, using reference encoding we can solve all the aforementioned issues of the offline encodings.

3.2 Online encoding with non-singletons

The reference encoding can be extended to the case using singleton together with non-singleton patterns to encode a data stream. Let $\mathfrak{S} = S_1 S_2 \cdots S_t$ denote a stream of sequences where each S_i is a sequence of events. Let D be a dictionary containing all characters of the alphabet and some non-singletons. Reference encodings compress a data stream \mathfrak{S} by replacing instances of words in the dictionary by references to the most recent occurrences of the words. If the words are non-singletons, beside the references, gaps between characters of the encoded words must be stored together with the references. Therefore, in a reference encoding, beside the reference stream we also have a gap stream.

Similar to the case with non-singleton, first we need to encode the dictionary D (now contains both singleton and non-singleton). We add a special symbol \sharp to the alphabet. The binary representation of the dictionary starts with the codeword of the size of the dictionary. It is followed by the codewords of all the characters in the alphabet each with length $\lceil \log_2 |D| \rceil$. The representations of every non-singleton follow right after that. The binary representation of a non-singleton contains codewords of the characters of the non-singleton. Non-singletons are separated from each other by the special character \sharp .

EXAMPLE 2 (DICTIONARY REPRESENTATION). The dictionary $D = \{a, b, c, \sharp, ab, abc\}$ can be represented as follows $E(6)C(a)C(b)C(c)C(\sharp)C(a)C(b)C(\sharp)C(a)C(b)C(c)$. The representation starts with E(6) indicating the size of D. It follows by the codewords of all the characters and the binary representation of the non-singletons separated by \sharp .

Having the binary representation of the dictionary, A first sends that representation to B. After that A send the encoding of the actual stream with the reference stream and the gap stream. The following example show how to encode a sequence with a reference encoding.

¹http://www.win.tue.nl/~lamthuy/projects/zips.pdf



Figure 2: An example of encoding the sequence S = abbcacbacbacbacbacbacbacbac with a reference encoding. The arrows represent the references between two consecutive encoded occurrences of a word which represent as a reference stream E(1)E(7)E(5)E(8)E(2)E(2)E(2)E(2)E(2)E(2)E(8). Having the reference stream we can reconstruct the stream but not in the same order of event occurrence. Therefore, we need a gap stream indicating the gaps between consecutive characters in each encoded non-singletons. For example, the gap stream is E(1)E(2)E(2)E(2)E(2)E(2)E(1)E(1).

dictionary, E(7) is the reference of the following b to the position of b in the dictionary and so on. The gap stream is E(1)E(2)E(2)E(2)E(2)E(1)E(1), where for instance, the first codewords E(1)E(2) indicate the gaps between a and b, b and c in the first occurrence of abc at position (1, 2, 4) in the stream. For non-singleton, there is no gap information representing in the gap stream.

EXAMPLE 4 (DECODING). In Figure 2, the sequence can be decoded as follows. Reading the first codeword of the reference stream, i.e. E(1), the decoder refers one step back to get abc. There will be two gaps (between a, b and b, c) so the decoder reads the next two codewords from the gap stream, i.e. E(1), and E(2). Knowing the gaps it can infer the positions of abc. In this case, the positions are 1,2 and 4. Subsequently, the decoder reads the next codeword from the reference stream, i.e. E(7), it refers seven steps back and decode the current reference as b. There is no gap because the word is a singleton, the position of b in the stream corresponds to the earliest position that has not been occupied by any decoded character, i.e. 3. The decoder continues decode the other references of the stream in the same way.

Different from the singleton case, there might be a lot of different reference encodings for a stream given a dictionary. Each reference encoding incurs different description lengths. Finding an optimal dictionary and an optimal reference encoding is the main problem we solve in this paper.

4. PROBLEM DEFINITION

Given a data stream \mathfrak{S} and a dictionary D denote $L_D^C(\mathfrak{S})$ as the description length of the data (including the cost to store the dictionary) in the encoding C. The problem of mining compressing sequential patterns in data stream can be formulated as follows:

DEFINITION 2 (COMPRESSING PATTERNS MINING). Given a stream of sequences \mathfrak{S} , find a dictionary D and an encoding C such that $L_D^C(\mathfrak{S})$ is minimized.

Generally, the problem of finding the optimal lossless compressed form of a sequence is NP-complete [11]. In this work, Algorithm 1 $\operatorname{Zips}(S)$

- 1: Input: Event stream $\mathfrak{S} = S_1 S_2 \cdots$
- 2: **Output**: Dictionary D
- $3: D \longleftarrow \emptyset$
- 4: for t = 1 to ∞ do
- 5: while $S_t \neq \epsilon$ do
- $6: \qquad w = encode(S_t)$
- 7: $w^* = extend(w)$
- 8: $update(w^*)$
- 9: end while
- 10: end for
- 11: Return D

Problem 2 is similar to the data compression problem but with additional constraint on the number of passes through data. Therefore, in next section we discuss a heuristic algorithm inspired by the idea of the *Lempel-Ziv*'s data compression algorithm [9] to solve this problem.

5. ALGORITHMS

In this section, we discuss an algorithm for finding a good set of compressing patterns from a data stream. Our algorithm is single-pass, memory-efficient and scalable. We call our algorithm Zips as for *Zip a stream*. There are three important subproblems that Zips will solve. The first problem concerns how to grow a dictionary of promising candidate patterns for encoding the stream. Since the memory is limited, the second problem is how to keep a small set of important candidates and evict from the dictionary unpromising candidates. The last problem is that having a dictionary how to encode the next sequence effectively with existing words in the dictionary.

The pseudo-code depicted in Algorithm 1 shows how Zips work. It has three subroutines each of them solves one of the three subproblems:

1. Compress a sequence given a dictionary: for every new sequence S_t in the stream, Zips uses the subroutine $encode(S_t)$ to find the word w in the dictionary which gives the most compression benefit when it is used to



Figure 3: An illustration of how compression benefit is calculated: (a) The sequence S in the uncompressed form. (b) two instances of w = abc: $S^1(w)$ and $S^2(w)$ and their references to the most recent encoded instance of w highlighted by the green color.

encode the uncompressed part of the sequence. Subsequently, the instance corresponds to the best chosen word is removed from S_t . Detail about how to choose w is discussed in subsection 5.1.

- 2. Grow a dictionary: Zips uses extend(w) to extend the word w chosen by the procedure $encode(S_t)$. Word extensions are discussed in subsection 5.2.
- 3. Dictionary update: the new extension is added to the dictionary. When the dictionary size exceeds the memory limit, a space-saving algorithm is used to evict unpromising words from the dictionary (subsection 5.3).

These steps are iteratively repeated as long as S_t is not encoded completely. When compression of the sequence finishes, Zips continues to compress the next in a similar way.

5.1 Compress a sequence:

Let S be a sequence, consider a dictionary word $w = a_1a_2\cdots a_k$, let S(w) denote an instance of w in S. Let g_2, g_3, \cdots, g_k be the gaps between the consecutive characters in S(w). We denote the gap between the current occurrence and the most recent encoded occurrence of w by g. Let $\overline{g_i}$ $i = 1, \cdots, k$ be the gap between the current and the most recent occurrence of a_i . Therefore, to calculate the compression benefit we subtract the size of encoding S(w) and the cost of encoding the gap to the last encoded occurrence of S(w) from the size of encoding each singleton:

$$B(S(w)) = \sum_{i=1}^{k} |E(\overline{g_i})| - |E(g)| - \sum_{i=2}^{k} |E(g_i)|$$
(1)

EXAMPLE 5 (COMPRESSION BENEFIT). Figure 3.a shows a sequence S in the uncompressed form and Figure 3.b shows the current form of S. Assume that the instance of w = abcat positions 1, 2, 4 is already compressed. Consider two instances of abc in the uncompressed part of S:

1. $S^1(w) = (a,3)(b,5)(c,7)$: the cost to replace this instance by a pointer is equal to the sum of the cost to encode the reference to the previous encoded instance of abc |E(1)| plus the cost of gaps |E(2)| + |E(2)|. The cost of representing this instance in an uncompressed form is |E(2)| + |E(3)| + |E(3)|. Therefore the compression benefit of using this instance to encode the sequence is $B(S^1(w)) = |E(2)| + |E(3)| + |E(3)| - |E(1)| - |E(2)| - |E(2)| = 3$ bits.

2. $S^{2}(w) = (a, 3)(b, 6)(c, 8)$: the compression benefit of using $S^{2}(w)$ to encode the sequence is calculated in a similar way: $B(S^{2}(w)) = |E(2)| + |E(1)| + |E(1)| - |E(1)| - |E(3)| - |E(2)| = -3$ bits.

In order to ensure that every symbol of the sequence is encoded, the next instance considered for encoding has to start at the first non-encoded symbol of the sequence. There maybe many instances of w in S that start with the first nonencoded symbol of S, denote $S^*(w) = argmax_{S(w)}B(S(w))$ as the one that results in the maximum compression benefit. We call $S^*(w)$ the best match of w in S. Given a dictionary, the encoding function depicted in Algorithm 2 first goes through the dictionary and finds the best match starting at the next uncompressed character of every dictionary word in the sequence S (line 4). Among all the best matches, it greedily chooses the one that results in the maximum compression benefit (line 6).

For any given dictionary word $w = a_1 a_2 \cdots a_k$, the most important subroutine of Algorithm 2 is to find the best match $S^*(w)$. This problem can be solved by creating a directed acyclic graph G(V, E) as follows:

- 1. Initially, V contains a start node s and an end node e
- 2. For every occurrence of a_i at position p in S, add a vertex (a_i, p) to V
- 3. Connect s with the node (a_1, p) by a directed edge and add to that edge a weight value equal to $|E(\overline{g}_1)| - |E(g)|$.
- 4. Connect every vertex (a_k, p) with e by a directed edge with weight 0.
- 5. For all q > p connect (a_i, p) to (a_{i+1}, q) by a directed edge with weight value $|E(\overline{g}_{i+1})| |E(q-p)|$

Algorithm 2 encode(S)

- 1: **Input**: a sequence S and dictionary $D = w_1 w_2 \cdots w_N$
- 2: **Output**: the word *w* that starts at the first non-encoded symbol gives the most additional compression benefit
- 3: for i = 1 to N do
- 4: $S^*(w_i) = bestmatch(S, w_i)$
- 5: **end for**
- 6: $max = argmax_i B(S^*(w_i))$
- 7: Return w_{max}



Figure 4: A directed acyclic graph created from the instances of abc in the uncompressed part of S shown in Figure 3.b

THEOREM 2. The best match $S^*(w)$ corresponds to the directed path from s to e with the maximum sum of the weight values along the path.

The proof of theorem 2 is trivial since any instance of win S corresponds to a directed path in the directed acyclic graph and vice versa. The sum of the weights along a directed path is equal to the benefit of using the corresponding instance of w to encode the sequence. Finding the directed path with maximum weight sum in a directed graph is a well-known problem in graph theory. That problem can be solved by a simple dynamic programming algorithm in linear time of the size of the graph, i.e. $O(|S|^2)[12]$.

EXAMPLE 6 (THE BEST MATCH IN A GRAPH). Figure 4 shows the directed acyclic graph created from the instances of abc in the uncompressed part of S shown in Figure 3.b. The best match of abc corresponds to the path s(a,3)(b,5)(c,7)e with the maximum sum of weights equal to 3 bits.

It is important to notice that in order to evaluate the compression benefit of a dictionary word, Equation 1 only requires bookkeeping the position of the most recent encoded instance of the word. This is in contrast to the offline encodings used in recent work [2, 3, 7] in which the bookkeeping of the word usage and the gaps cost is a must. When a new instance of a word is replaced by a pointer, the relative usage and the codewords of all dictionary words also change. As a result, the compression benefit needs to be recalculated by a pass through the dictionary. This operation is an expensive task when the dictionary size is unbounded.

5.2 Dictionary extension:

Initially, the dictionary contains all singletons; it is iteratively expanded with the locally best words. In each step, when the best match of a dictionary word has been found, Zips extends the best match with one extra character and adds this extension to the dictionary. There are different options to choose the character for extension. In this work, Zips chooses the next uncompressed character right after the word. The choice is inspired by the same extension method suggested by the *Lempel-Ziv* compression algorithms.

Moreover, there is another reason behind our choice. When w is encoded for the first time, the reference to the previous encoded instance of w is undefined although the word has been already added to the dictionary. Under that circumstance, we have to differentiate between two cases: either a reference to an extension or to an encoded word. To achieve this goal one extra flag bit is added to every reference. When the flag bit is equal to 1, the reference refers to an extension of an encoded word. Otherwise, it refers to an encoded



Figure 5: An example of word is extended and encoded the first time and the second time. One extra flag bit is added to every reference to differentiate two cases.

word. When the former case happens by extending the word with the character right after it, the decoder always knows where to find the last character of the extension. When a word is added to a dictionary all of its prefixes have been already added to the dictionary. This property enables us to store the dictionary by using a prefix tree.

EXAMPLE 7. Figure 5 shows the first moment t_0 when the word w = abc is added to the dictionary and two other moments t_1 and t_2 when it is used to encode the stream. At t_1 , the flag bit 1 is used to notify the decoder that the gap g_1 is a reference to an extension of an encoded word, while at t_2 the decoder understands that the gap g_2 is a reference to the previous encoded instance of w.

References to extensions may cause ambiguous references. For instance, a potential case of ambiguous reference called *reference loop* is discussed in the following example:

EXAMPLE 8 (REFERENCE LOOPS). At time point t_0 the word w = ab at positions p_1p_2 is extended by c at position p_3 . Later on at another time point $t_1 > t_0$, another instance of abc at $q_1q_2q_3$ ($q_1 > p_1$) refers back to the instance abc at $p_1p_2p_3$. At time point $t_2 > t_1$, another instance of abc at $r_1r_2p_3$ ($r_1 > q_1$) refers back to the instance abc at $q_1q_2q_3$. In this case, c at p_3 and c at q_3 refers to each other forming a reference loop.

Reference loops result in ambiguity when we decode the sequence. Therefore, when we look for the next best matches, in order to avoid reference loops, we always check if the new match incurs a loop. The match that incurs a loop is not considered for encoding the sequence. Checks for ambiguity can be done efficiently by creating paths of references. Vertices of a reference path are events in the sequence. Edge between two consecutive vertices of the path corresponds to a reference between the associated events. Since the total sizes of all the paths is at most the sequence length, its is cheap to store the paths for a bounded size sequence. Moreover, updating and checking if a path is a loop can be done in O(1) if vertices of the path are stored in a hashmap.

EXAMPLE 9 (REFERENCE PATHS). The references paths of the encoding in Example 8 are: $(a, r_1) \mapsto (a, q_1) \mapsto (a, p_1)$, $(b, r_2) \mapsto (b, q_2) \mapsto (b, p_2)$ and $(c, p_3) \mapsto (c, q_3) \mapsto (c, p_3)$. The last path is a loop.

5.3 Dictionary update:

New extensions are added to the dictionary until the dictionary exceeds memory limit. When it happens the *spacesaving* algorithm is used to evict unpromising words from the dictionary. The space-saving algorithm [10] is a well-known Algorithm 3 update (w^*)

word 1: Input: \mathbf{a} w^* and dictionary D_ $\{w_1, w_2, \cdots, w_N\}$ 2: **Output**: the dictionary D 3: $m \leftarrow |i:w_i|$ is a non-singleton 4: $v = argmin_i w_i [1]$ and v is non-singleton at a leaf of the prefix-tree 5: if m > M and $w^* \notin D$ then $D = D \setminus \{v\}$ 6: $w^*[1] = w^*[2] = v[1]$ 7: $D = D \bigcup \{w^*\}$ 8: 9: else if $w^* \notin D$ then $w^*[1] = w^*[2] = 0$ 10: $D = D \bigcup \{w^*\}$ 11:12: else 13: add additional compression benefit to $w^*[1]$ 14: end if 15: Return D



Figure 6: The running time and the average update time per sequence of the Zips algorithm in the Tweets dataset when the stream size increases. Zips scales linearly with the size of the stream.

method proposed for finding the most frequent items in a stream of items given a budget on the maximum memory. In this work, we propose a similar space-saving algorithm to keep the number of non-singleton words in the dictionary at below a predefined number M while it can be able to return the set of compressing patterns with high accuracy.

The algorithm works as follows, for every non-singleton word w it maintains a counter with two fields. The first field denoted as w[1] contains an over-estimate of the compression benefit of w. The second field denoted as w[2] contains the compression benefit of the word with least compression benefit in the dictionary at the moment that w is inserted into the dictionary.

Every time when a word w is chosen by Algorithm 2 to encode its best match in the sequence S_t , the compression benefit of the word is updated. The word w is then extended to w^* with an extra character by the extension subroutine. In its turn, Algorithm 3 checks if the dictionary already contains w^* . If the dictionary does not contains w^* and it is full with M non-singleton words, the least compressing word v resident at a leaf of the dictionary prefix-tree is removed from the tree. Subsequently, the word w^* is inserted into the tree and its compression benefit can be over-estimated as w[1] = w[2] = v[1]. The first counter of every word is always greater than the true compression benefit of the



Figure 7: Running time (x-axis) against the data size (y-axis) of three algorithms in log-log scales. The Zips algorithm scales linearly with the data size while the GoKrimp and the SQS algorithm scales quadratically with the data size.

words. This property ensures that the new emerging word is not removed very quickly because its accumulated compression benefit is dominated by long lasting words in the dictionary. For any word w, the difference between w[2]and w[1] is the actual compression benefit of w since the moment that w is inserted into the dictionary. At anytime point when we need to find the most compressing patterns, we compare the value of w[2] - w[1] and select those with highest w[2] - w[1]. In section 6 we show empirical results with different datasets that this algorithm is very effective in finding the most compressing patterns with high accuracy even with limited memory.

6. EXPERIMENTS

We perform experiments with one synthetic dataset with known ground truth and two real-world datasets. Our implementation of the Zips algorithm in C++ together with the datasets are available for download at our project website². All the experiments were carried out on a machine

²www.win.tue.nl/~lamthuy/zips.html

with 16 processor cores, 2 Ghz, 12 GB memory, 194 GB local disk, Fedora 14 / 64-bit. As baseline algorithms, we choose the GoKrimp algorithm [2, 3] and the SQS algorithm [7] for comparison in terms of running time, scalability, and interpretability of the set of patterns.

6.1 Data

Three different datasets are:

- 1. JMLR: contains 787 abstracts of articles in the Journal of Machine Learning Research. English words are stemmed and stop words are removed. JMLR is small but it is considered as a benchmark dataset in the recent work [2, 3, 7]. The dataset is chosen also because the set of extracted patterns can be easily interpreted.
- 2. **Tweets**: contains over 1270000 tweets from 1250 different twitter accounts³. All tweets are ordered ascending by timestamp, English words are stemmed and stop words are removed. After preprocessing, the dataset contains over 900000 tweets. Similar to the JMLR dataset, this dataset is chosen because the set of extracted patterns can be easily interpreted.
- 3. **Plant**: is a synthetic dataset generated in the same way as the generation of the plant10 and plant50 dataset used in [7]. The plant10 and plant50 are small so we generate a larger one with ten patterns each with 5 events occurs 100 times at random positions in a sequence with length 100000 generated by 1000 independent noise event types.

6.2 **Running time and Scalability**

Figure 6 plots the running time and the average update time per sequence of the Zips algorithm in the Tweets dataset when the data stream size (the number of sequences) increases. Three different lines correspond to different maximum dictionary size settings M = 1000, M = 5000 and M = 10000 respectively. The results show that the Zips algorithm scales linearly with the size of the stream. The average update time per sequence is constant given a maximum dictionary size setting. For example, when M = 10000, Zips can handle one update in about 20-100 milliseconds.

Figure 7 shows the running time in y-axis of the Zips algorithm against the stream size in x-axis in the Tweets dataset when the maximum dictionary size is set to 1000. In the same figure, the running time of the baseline algorithms GoKrimp and SQS are also shown. There are some missing points in the results corresponding to the SQS algorithm because we set a deadline of ten hours to get the results corresponding to a point. The missing points corresponding to the cases when the SQS program did not finish in time.

In the log-log scale, running time lines resemble straight lines. This result shows that the running time of Zips, GoKrimp and SQS is the power of data size, i.e. $T \sim \alpha |S|^{\beta}$. Using linear fitting functions in log-log scale we found that with the Zips algorithm $\beta = 1.01$, i.e. Zips scales linearly with the data size. Meanwhile, for the SQS algorithm the exponent is $\beta = 2.2$ and for the GoKrimp algorithm the exponent is $\beta = 2.01$. Therefore, both GoKrimp and SQS do not scale linearly with the data size and hence they are not suitable for data stream applications.

6.3 JMLR

In Figure 8, we show the first 20 patterns extracted by two baseline algorithms GoKrimp and SQS and the Zips algorithm from the JMLR dataset. Three lists are slightly different but the important patterns such as "support vector machine", "data set", "machine learn", "bayesian network" or "state art" were discovered by all of the three algorithms. This experiment confirms that the Zips algorithm was able to find important patterns that are consistent with the results of state-of-the-art algorithms.

6.4 Tweets

Since the tweet dataset is large, we schedule the programs so that they terminate their running after two weeks. The SQS algorithm was not able to finish its running before the deadline while GoKrimp finished running after three days and Zips finished running after 35 hours. The set of patterns extracted by the Zips algorithm and the GoKrimp algorithm are shown in Figure 9. Patterns are visualized by the wordcloud tool in R such that more important patterns are represented as larger words. In both algorithms, the sets of patterns are very similar. The result shows the daily discussions of the 1250 twitter accounts about the topics regarding "social media", "Blog post', about "youtube video", about "iphone apps", about greetings such as "happy birthday", "good morning" and "good night", about custom service complaint etc.

6.5 Plants

It has been shown [7] that SQS successfully returned all ten patterns. We obtain the same result with GoKrimp and Zips. All three algorithm ranked 10 true patterns with highest scores.

7. CONCLUSIONS AND FUTURE WORK

In this paper we studied the problem of mining compressing patterns from a data stream. A new encoding scheme for sequence is proposed. The new encoding is convenient for streaming applications because it allows encoding the data in an online manner. Because the problem of mining the best set of patterns with respect to the given encoding is shown to be unsolvable under the streaming context, we propose a heuristic solution that solves the mining compressing problem effectively. In the experiments with one synthetic dataset with known ground-truths Zips was able to extract the most compressing patterns with high accuracy. Meanwhile, in the experiments with two real-world datasets it can find patterns that are similar to the-state-of-the-art algorithms extract from these datasets. More importantly, the proposed algorithm was able to scale linearly with the size of the stream while the state-of-the-art algorithms were not. There are several options to extend the current work. One of the most promising future work is to study the problem of mining compressing patterns for different kinds of data stream such as a stream of graphs.

8. **REFERENCES**

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³http://user.informatik.uni-goettingen.de/~txu/ cuckoo/dataset.html

Method	Patterns			
SQS	support vector machin	larg scale	featur select	sampl size
	machin learn	nearest neighbor	graphic model	learn algorithm
	state art	decis tree	real world	princip compon analysi
	data set	neural network	high dimension	logist regress
	bayesian network	cross valid	mutual inform	model select
GOKRIMP	support vector machin	state art	neural network	well known
	real world	high dimension	experiment result	special case
	machin learn	reproduc hilbert space	sampl size	solv problem
	data set	larg scale	supervis learn	signific improv
	bayesian network	independ compon analysi	support vector	object function
Zips	support vector machin	featur select	high dimension	cross valid
	data set	machine learn	paper propose	decis tree
	real world	bayesian network	graphic model	neutral network
	learn algorithm	model select	larg scale	well known
	state art	optim problem	result show	hilbert space

Figure 8: The first 20 patterns extracted from the JMLR dataset by two baseline algorithms GoKrimp and SQS and the Zips algorithm. Common patterns discovered by all the three algorithms are bold.



Figure 9: The top 20 most compressing patterns extracted by Zips (left) and by GoKrimp (right) from the Tweets dataset.

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