

A Survey of Pattern Mining in Dynamic Graphs

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Abstract

Graph data is found in numerous domains **such as for the analysis of social networks, sensor networks**, bioinformatics, industrial systems, and chemistry. Analyzing graphs to **identify** useful and interesting patterns is an important **research area**. It helps understanding graphs, and hence support decision making. Since two decades, many graph mining **algorithms** have been proposed to identify patterns such as frequent subgraphs, paths, cliques and trees. But **most of them** assume that graphs are static. This simplifying assumption makes it easy to design algorithms but discard information about how graphs evolve. This paper provides a detailed survey of techniques for mining interesting patterns in dynamic graphs, which can serve both as an introduction and as a guide to recent advances and opportunities in this research area. The main tasks related to mining patterns in dynamic graphs are reviewed such as discovering frequent subgraphs, evolution rules, motifs, subgraph sequences, recurrent and triggering patterns, and trend sequences. In addition, an overview of strategies and approaches to solve dynamic graph mining problems is presented, and their advantages and limitations are highlighted. Various extensions are also discussed such as to discover patterns in data **streams** and big data. Lastly, the article mentions several research opportunities.

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INTRODUCTION

Pattern mining is a key research area in data mining, which consists of applying algorithms to identify interesting patterns in data. Generally, a pattern can be considered interesting if it reveals some novel information that is useful to understand the past or predict the future. Over the years, techniques have been designed to extract patterns from several types of data such as transactional data^{9,10}, time series^{14–16}, business process logs¹¹, trajectories¹², spatial data¹⁷, sequences^{1,5,8,13,19–21} and graphs^{2–4}. Depending on the applications, what is an interesting pattern differs. Thus, algorithms have been proposed that apply various criteria to **identify** patterns such as finding those having a high occurrence frequency and confidence^{10,18}, rarity^{22,23}, profitability²⁴, or a low-cost^{6,7}. Pattern mining tasks can be very challenging because an algorithm must consider a potentially huge number of possible patterns to discover the desired ones. Consequently, efficient algorithms have been designed based on efficient data structures and search space pruning strategies^{1,10}.

Among the various types of data studied in this research area, graphs are one of the most important **ones** as they are found in numerous domains such as social networks⁹⁵, chemistry²⁷, vehicular networks⁴⁵, computer networks²⁵, bioinformatics⁴¹, XML data⁴¹, and geographical data¹⁰². Algorithms have been proposed to find various types of patterns in graphs such as subgraphs^{2,27,34,35}, trees^{41,42} and traversal paths^{56–58}. Moreover, various types of graphs have been studied such as weighted graphs⁵⁶, directed graphs^{32,42}, attributed graphs^{50,89,97,99,101,102}, and graph databases^{2,3}. A graph type that has attracted the interest of many researchers is *dynamic graphs*. Those are graphs that change over time in terms of attributes (labels) or structures (edges, vertices). Considering the time dimension when mining graphs allows understanding how they evolve and is key to many applications such as social network analysis. However, mining patterns in dynamic graphs is also more challenging because the search space is larger when taking time into account.

Graph mining is a very active research field. Although some surveys have been published on discovering patterns in graphs³, there is none on mining patterns in dynamic graphs. This paper addresses this issue by providing an up-to-date and detailed survey that is not only an introduction to the field but also reviews recent advances and opportunities.

It is to be noted that this survey focuses on analyzing graphs to find *patterns*, i.e. interesting sets of value that are appearing several times in data, are correlated or meet some other interestingness criteria set by the user. Other useful graph analysis tasks are considered to be outside the scope of this survey such as identifying prestigious nodes⁵², detecting communities^{52,53}, calculating descriptive measures⁵⁴, clustering nodes⁵⁴, graph summarization⁵⁵, automatic node labeling (relational classification)⁵¹, and graph visualization⁵⁴.

This survey is organized as follows. It first presents important concepts and challenges related to mining patterns in static graphs. Then, the following sections discuss techniques for discovering patterns in a single dynamic graph, common to several dynamic graphs (a graph database), and in attributed graphs. In these sections, an overview of techniques employed for discovering different kinds of patterns is presented. Then, the paper discusses research opportunities. Finally, a conclusion is drawn.

MINING PATTERNS IN STATIC GRAPH(S)

Before discussing techniques for mining patterns in dynamic graphs, this section presents a brief overview of techniques for mining patterns in static graphs, as they are related. Then, the next section reviews techniques for mining patterns in dynamic graphs.

Note that this section is not as detailed as the following one about dynamic graphs because in-depth surveys dedicated to mining patterns in static graphs have been published³, and the focus of this paper is dynamic graphs.

PRELIMINARY DEFINITIONS

In its most simple form, a *graph* is a tuple $G = (V, E)$ such that V is a vertex set and E is an edge set. The vertex set of a graph is a non empty finite set of elements called *vertices*. The edge set contains zero or more *edges*, defined as two-elements subsets of V . In other words, $E \subseteq V \times V$. Let there be two vertices $u, v \in V$. If there exists an edge $\{u, v\} \in E$, then u and v are said to be *adjacent*, *connected* and *reachable* from each other, and $\{u, v\}$ is said to *connect* or *join* u and v . Moreover, it is said that $\{u, v\}$ is *incident* to u and to v . Two edges e_1 and e_2 are *adjacent* if they are incident to a same vertice w , i.e. $\exists w \in e_1 \cap e_2$.

Some common vocabulary to describe graphs is presented below. Note that in the above definition of graph, self-loops (edges connected to themselves) are not allowed.

A graph is *connected* if there exists a walk from any vertex u to any other vertex v . A *walk* in a graph is a sequence of vertices $\langle v_1, v_2, \dots, v_n \rangle$ such that v_1 is adjacent to v_2 , v_2 is adjacent to v_3 , \dots , v_{n-1} is adjacent to v_n . If such walk exists then v_n is said to be *reachable* from v_1 . A graph that is not connected is said to be *disconnected*.

A *directed graph* (also called *digraph*) is a graph $G = (V, E)$, where V is a non empty finite vertex set and E is a set of directed edges. A *directed edge* is an ordered vertex pair (u, v) where $u, v \in V$, which indicates that v can be *reached* from u , or equivalently that v is a *successor* of u , and that u is *connected* and *adjacent* to v . A directed edge (u, v) is often represented visually as an arrow from u to v , and is different from the reverse edge (v, u) . Note that a directed graph may contain self-loops.

A graph is *weighted* if weights are assigned to edges and/or vertices. This is done by adding two functions mapping vertices and/or edges to positive real numbers, that is $VW : V \mapsto \mathbb{R}^+$ and $EW : E \mapsto \mathbb{R}^+$, respectively. A weight can be interpreted as indicating the relative importance of an edge or vertex, or have other application specific meanings.

A graph is said to be *simple* if it is not weighted, undirected, has no self-loop and has **at most one** edge between any pair of vertices.

The above graph types are useful to model how some elements (objects) are connected to each other. To encode semantic information in a graph, labels may be added. A *labeled graph* is a tuple $G = (V, E, L_V, L_E, \phi_V, \phi_E)$ where V is an edge set, $E \subseteq V \times V$ is an edge set, L_V is a set of vertex labels, L_E is a set of edge labels, ϕ_V is a function mapping vertex to labels ($\phi_V:V \mapsto L_V$), and ϕ_E is a function mapping edges to labels ($\phi_E:E \mapsto L_E$).

An *attributed graph* is a tuple $G = (V, A, E, \lambda)$ where V is a vertex set, A is an attribute set, $E \subseteq V \times V$ is an edge set, and $\lambda : V \times A \mapsto \mathbb{R}$ is a function that maps each vertex-attribute pair to a number, which may represents a value or label. Note that it is also possible to define attributed graphs in a more general way such that edges may also have multiple attributes. That definition would then be similar to that of *multi-relational graphs* (graphs where multiple edges of different types may connect a same pair of nodes) and that of *multi-labeled graphs* (graphs where nodes and edges can have multiple labels). But a

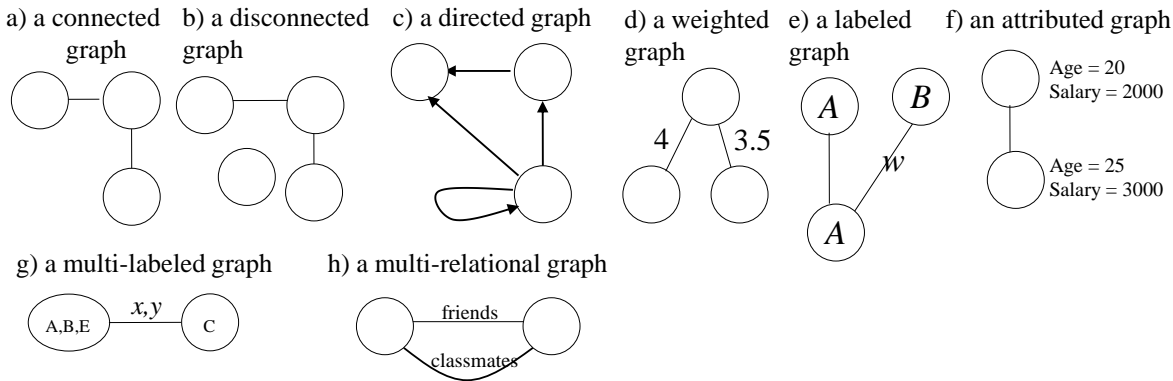


Figure 1: Different types of static graphs

difference with these latter definitions is that attributed graphs explicitly group labels by attributes, while multi-labeled graphs and multi-relational graphs do not explicitly represent attributes. Various other types of graphs have also been studied. Fig. 1 shows examples of the main types of graphs, discussed above.

MINING PATTERNS IN A STATIC GRAPH DATABASE

Several tasks have been proposed to find patterns in a database of static graphs.

Frequent subgraph mining. It is one of the most popular graph mining task. It aims at finding all subgraphs that appear frequently in a database of simple connected graphs^{2-4,25}. Given a parameter called the minimum support threshold (*minsup*), a graph is frequent if it appears in no less than *minsup* input graphs. The assumption of frequent subgraph mining is that a subgraph is interesting if it appears many times in a set of graphs. For example, this task can be useful to find an association between elements that is common to several chemical molecules.

Formally, frequent subgraph mining is defined as follows. Let there be a *graph database* $GD = \{G_1, G_2 \dots G_n\}$ consisting of n simple labeled graphs. Consider two labeled graphs $G_x = (V_x, E_x, L_{V_x}, L_{E_x}, \phi_{V_x}, \phi_{E_x})$ and $G_y = (V_y, E_y, L_{V_y}, L_{E_y}, \phi_{V_y}, \phi_{E_y})$. The graph G_x is said to be *isomorphic to* G_y if and only if there exists a bijective function $f : V_x \rightarrow V_y$ such that (1) $\forall v \in V_x, L_{V_x}(v) = L_{V_y}(f(v))$ and (2) $\forall \{u, v\} \in E_x, \{f(u), f(v)\} \in E_y$ and $L_{E_x}(u, v) = L_{E_y}(f(u), f(v))$. A graph G_x is said to be a *subgraph isomorphism* of (to appear

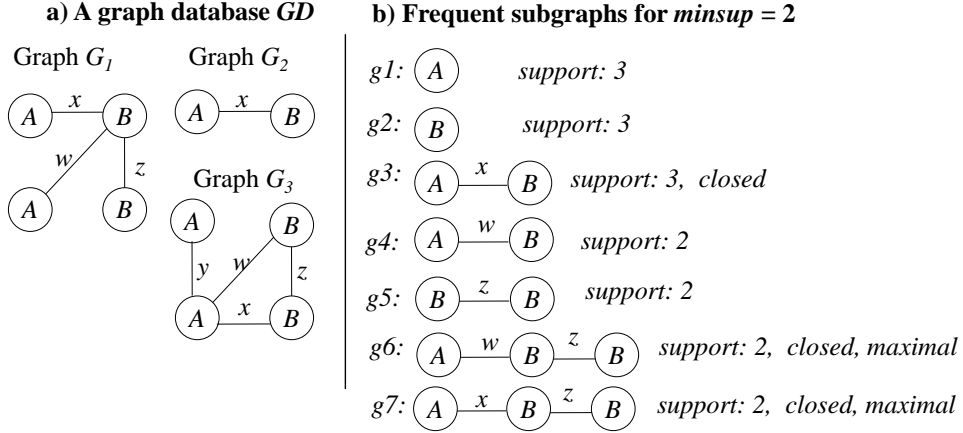


Figure 2: A graph database and frequent subgraphs found for $minsup = 2$

in) a graph G_z , denoted as $G_x \sqsubseteq G_z$, if there exists a subgraph $G_y \subseteq G_z$ such that G_x is isomorphic to G_y . A subgraph isomorphism is also called an *embedding*. The *support* of a graph G_x in a graph database GD is defined as $sup(G_x) = |\{g | g \in GD \wedge G_x \sqsubseteq g\}|$. Given a graph database GD and a *minsup* threshold ($minsup > 0$), the task of frequent subgraph mining is to enumerate all frequent subgraphs (graphs having a support no less than *minsup*).

For example, consider the graph database of Fig. 2 a) containing three graphs, G_1, G_2 and G_3 . If $minsup = 2$, seven frequent subgraphs are found, represented on Fig. 2 b) with their support. For example, the subgraph g_6 has a support of 2 because it appears in G_1 and G_3 , while subgraph g_1 has a support of 3 because it appears in G_1, G_2 and G_3 . It is to be noted that support calculations ignore that a graph may have multiple occurrences in an input graph (e.g. g_1 appears multiple times in G_1).

The problem of frequent subgraph mining is difficult because a potentially very large number of subgraphs must be considered, and their support must be calculated, to find the frequent subgraphs. Several efficient algorithms have been proposed to find frequent subgraphs efficiently. Generally, they start **searching** from graphs each having a single vertex or edge, and recursively append edges to these graphs to obtain larger graphs. Algorithms such as FSG²⁶ perform a *breadth-first search* to explore the search space of all subgraphs. They first find all subgraphs having one edge. Then, the algorithms grow these subgraphs to find those having two edges. Then, those having three edges are considered and so on,

until no patterns can be generated. Other algorithms adopt a depth-first search. They also start from patterns **each having a single edge** but recursively grow a pattern before growing others. Such algorithms are MoFa²⁷, gSpan², FFSM³⁴, Gaston³⁵ and FPGraphMiner²⁵. These algorithms have the same input and output but use different search strategies and data structures **in their search for patterns**.

Generally, **two** key challenges for designing an efficient frequent subgraph mining algorithm **are** how to explore the search space and how to perform support counting. To avoid exploring the whole search space, a key property is that the support measure is anti-monotonic, i.e. the support of a subgraph is always greater or equal to those of its supergraphs^{2,26,34,35}. Thus, if a subgraph is infrequent, all its supergraphs can be eliminated from the search space as they cannot be frequent subgraphs. This property, often called *downward closure property* or *Apriori* property is used by most frequent subgraph mining algorithms, and has been used for several other pattern mining problems such as itemset mining^{10,18,120,121} and sequential pattern mining¹. Another problem related to search space exploration is that an algorithm may generate candidate subgraphs that are isomorphic (equivalent) to subgraphs that it has previously considered. To avoid considering these subgraphs again several algorithms perform *isomorphism checking*. Though both efficient exact and approximate linear time isomorphism checking algorithms have been proposed³⁶, it remains a computationally expensive task.

A few key strategies adopted by some popular frequent subgraph mining algorithms are described in more details. The FSG algorithm²⁶ performs a breadth-first search. It generates candidate subgraphs having $k + 1$ edges by *joining* pairs of frequent subgraphs having k edges that have $k - 1$ edges in common. This join strategy can avoid considering many infrequent subgraphs. However, a drawback is that many **k edge subgraphs** may need to be kept in memory to generate the **$k + 1$ edge subgraphs**. Besides, this approach can produce some subgraphs that do not exist in the database. To calculate the support of a generated subgraph, FSG utilizes a vertical database representation which consists of annotating each subgraph with the list of input graphs and nodes where the subgraph appears. When joining two subgraphs, their lists are then combined to obtain that of the generated subgraph, and hence its support. To facilitate graph comparison, FSG calculates a unique code for

each subgraph. Such unique code, called a *canonical labelling* or *canonical representation*, facilitates isomorphism checking since all isomorphic graphs have the same code. The gSpan algorithm² is arguably the most popular frequent subgraph mining algorithm. It performs a depth-first search, which starts from single edge subgraphs and *extends* each one recursively by adding one edge at a time. To avoid generating graphs that do not exist, gSpan utilizes a *pattern-growth approach*. That is, it searches for extensions of a subgraph by scanning the input graphs where the subgraph appears. For each identified frequent subgraph, gSpan calculates a canonical code called DFS code based on the depth-first search traversal order. If the DFS code is not minimal, the subgraph is a duplicate of another subgraph and can be ignored. This technique is an efficient solution for identifying duplicate subgraphs. gSpan performs isomorphism checking only to check if a subgraph appears in an input graph. An advantage of the depth-first search approach of gSpan over the breadth-first search is that only subgraphs of the current recursive call are kept in memory at any time. The FFSM³⁴ algorithm also performs a depth-first search, but a different scheme is used for the canonical labelling of subgraphs, which is based on a canonical adjacency matrix representation of graphs. Besides, FFSM adopts both the concept of join between subgraphs to generate larger subgraphs and that of extension, and was shown to be competitive with gSpan. The FPGraphMiner²⁵ algorithm first creates a BitCode structure (a bit vector) for each single edge subgraph that indicates in which input graphs the subgraph appears. Then, FPGraphMiner stores all subgraphs in a special graph called FP-graph, where subgraphs having a same bitcode are stored in a same node, and nodes having the same support are grouped in clusters. Then, frequent subgraphs are mined directly from the FP-graph structure using a depth-first search without back-tracking. This approach was shown to be very efficient when compared with several previous algorithms.

Although frequent subgraph mining is useful, a problem is how to set the *minsup* threshold²⁸⁻³¹. If it is set too high, few patterns are found, while if it is set too low, too many patterns may be found and algorithms may have very long runtimes or run out of memory. To address this issue, the problem of *top-k frequent subgraph mining* was proposed, where the user can directly set the number k of patterns to find rather than using the *minsup* threshold. The first algorithm for this problem, named TGP²⁹, initially reads an input

graph database to obtain the DFS codes of all subgraphs of each input graph. Thereafter, TGP stores all these DFS code in a large structure named Lexicographical Pattern Net. Then, TGP searches for the top-k subgraphs using that structure, while gradually raising an internal minsup threshold initially set to 0. A major issue with TGP is that it explicitly generates all patterns to then find the top-k patterns, which makes it inefficient even for moderately large graph databases^{28,29}. An alternative algorithm named FS³ was designed to trade result completeness and accuracy for efficiency³⁰. To apply FS³, a user must set k , the size s for subgraphs to be found, and a number of iterations. During each iteration, FS³ samples a graph from the database, and then (2) samples a s -size subgraph biased toward frequent subgraphs in the whole database using the Markov Chain Monte Carlo method. A reason for FS³'s efficiency is that it does not perform subgraph isomorphism checking but a drawback is that it may incorrectly calculate the support of patterns. Hence, the algorithm may return infrequent patterns and miss frequent patterns. Another approximate algorithm named kSIM³¹ was proposed to process a restricted type of graphs called induced subgraphs, and was shown to outperform FS³ in runtime and accuracy. Recently, an exact algorithm for top-k frequent subgraph mining was presented, named TKG²⁸, which extends gSpan and has similar performance.

Besides top-k frequent subgraph mining, many other extensions of the frequent subgraph mining problems have been proposed to find subgraphs using other measures to select patterns such as density, edge connectivity and vertex connectivity³⁷, and subgraphs having a high correlation⁴⁰.

To reduce the number of patterns that are presented to the user, algorithms have also been proposed to mine concise representations of frequent subgraphs such as closed and maximal subgraphs. A *frequent closed subgraph* is a frequent subgraph that is not a subgraph of any other frequent subgraph having the same support^{29,32}. A *frequent maximal subgraph* is a frequent subgraph that is not a subgraph of any other frequent subgraph^{38,39}. Hence, closed subgraphs are a subset of maximal **subgraphs**. For example, in Fig. 2 (right), closed and maximal frequent subgraphs are indicated. An interesting property is that frequent maximal subgraphs allow recovering all frequent subgraphs without scanning the database, while frequent closed subgraphs also allow deriving their support values. Moreover, mining

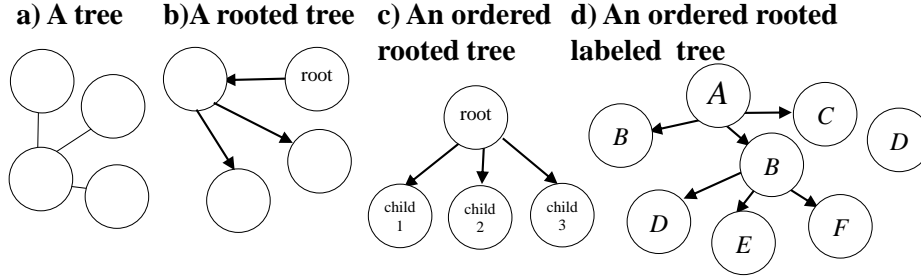


Figure 3: Different types of trees

maximal and closed subgraphs is often faster than mining all frequent subgraphs. Another notable concise representation of frequent subgraphs that is useful for classification tasks is *frequent generator subgraphs*³³. They are the frequent subgraphs that have no proper subgraphs having the same support³².

Some frequent subgraph mining algorithms can also be extended with small modifications to mine partially labeled graphs, directed graphs, graphs with self-loops, and multiple edges between vertices³².

Frequent subtree mining. Another popular task for mining patterns in static graphs is *frequent subtree mining*^{41,42}. The problem consists of identifying a set of subtrees appearing frequently in a database of trees. Several types of trees are considered in the literature⁴² (which are graphs). A *tree* $T = (V, E)$ is an undirected connected graph that is acyclic (there exists no vertex $v \in V$ such that there is a walk $\langle v, \dots, v \rangle$ that starts from v and leads to v). A *rooted tree* $T = (V, E, r)$ is a directed acyclic graph where a vertex (node) $r \in V$ is called the *root*. Moreover, there exists a walk from the root to every other node, and no walk leading to the root. Note that a tree that has no node designated as root is sometimes called a free tree. An *ordered rooted tree* is a rooted tree where each vertex's children are ordered as a list from first to last.<https://www.overleaf.com/project/5cd2f5574ae8d80fa8148eb2> A *labeled tree* is a tree such that each vertex has a label. Fig. 3 shows examples of these main types of trees. Trees can represent many types of data. For example, the tag structures of XML documents and hierarchies of DNS servers can be viewed as ordered rooted trees.

Three main types of subtrees can be extracted from a tree database^{41,42}. They are defined as follows. Let there be two ordered labeled trees $T_x = (V_x, E_x, L_{V_x}, L_{E_x}, \phi_{V_x}, \phi_{E_x})$ and $T_y = (V_y, E_y, L_{V_y}, L_{E_y}, \phi_{V_y}, \phi_{E_y})$. The tree T_x is a *bottom-up subtree* of T_y if $E_x \subseteq E_y$, $V_x \subseteq V_y$,

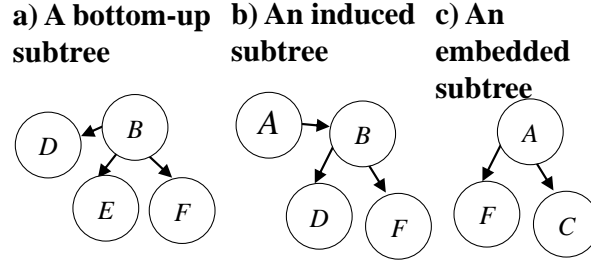


Figure 4: Different types of subtrees of the tree of Fig. 3 d)

labels of nodes/edges and the ordering of child nodes are preserved in T_x , and all descendants of each node $v \in V_y$ are also in T_x . The tree T_x is an *induced subtree* of T_y if $E_x \subseteq E_y$, $V_x \subseteq V_y$, labels of nodes/edges are preserved in T_x , and parent-child relationships between nodes are preserved. The tree T_x is an *embedded subtree* of T_y if $E_x \subseteq E_y$, $V_x \subseteq V_y$, labels of nodes/edges are preserved in T_x , and ancestor/descendant relationships between nodes are preserved. Thus, all bottom-up subtrees are induced subtrees, and all induced subtrees are embedded subtrees ($BottomUpSubtrees \subseteq InducedSubtrees \subseteq EmbeddedSubtrees$). Three examples of subtrees of the tree of Fig. 3 d) are shown in Fig. 4. The goal of frequent subtree mining is to find all subtrees that are bottom-up, induced or embedded subtrees of at least *minsup* trees of a tree database. Some classic algorithms are TreeMiner⁴¹ for mining frequent ordered embedded subtrees in a database of rooted ordered trees, and FREQT⁴³ to discover all frequent induced subtrees in a database of rooted ordered trees. Tree mining has applications in bioinformatics such as identifying common phylogenetic subtrees and RNA structures⁴¹. Subtree mining can be viewed as a special case of subgraph mining. But the former is a tractable problem while the latter is intractable. Recently, the problem of frequent tree mining has been generalized as *frequent attributed tree mining* to consider trees and subtrees where each node may have multiple labels (*attributed trees*)⁵⁰.

Algorithms such as TreeMiner can also be extended with minor modifications for similar tree mining problems such as discovering unlabeled subtrees, unordered subtrees and frequent sub-forests (disconnected subtrees)⁴¹.

Other tasks. Several variations of the above tasks have also been proposed such as to discover frequent sub-DAG in a database of DAG (directed acyclic graphs)⁵⁹, and to discover frequent subgraphs in a database of outter-planar graphs (a type of **graphs** that can be drawn

on a plane without any crossing edges)⁶⁰.

MINING PATTERNS IN A SINGLE STATIC GRAPH

The previous subsection has reviewed techniques for mining patterns in a database of static graphs. This section reviews the main tasks for discovering patterns in a single static graph.

Frequent subgraph mining in a single graph. Some frequent subgraph mining algorithms can be adapted to mine frequent subgraphs appearing frequently in a single graph. To do this, it is necessary to define an appropriate support counting function. A simple solution is to define the support of a pattern as its number of occurrences in the graph. For example, consider the graph G of Fig. 5 (left). Fig. 5 (right) shows frequent subgraphs found in G when the minimum threshold is set to 2 occurrences (right). However, this simple definition raises two important problems. First, it allows subgraph occurrences to be overlapping, which may be undesirable for some applications. Second, this support measure is not anti-monotonic (the support of a subgraph may be greater, smaller or equal to the support of its supergraphs), and thus the powerful downward closure property cannot be directly used to reduce the search space^{47,48}. This is illustrated with a simple example⁴⁸. In a graph $\textcircled{X}-\textcircled{Y}-\textcircled{X}$, the subgraph \textcircled{Y} has one occurrence: $\textcircled{X}-\textcircled{Y}-\textcircled{X}$. The subgraph $\textcircled{X}-\textcircled{Y}$ has two overlapping occurrences: $\textcircled{X}-\textcircled{Y}-\textcircled{X}$ and $\textcircled{X}-\textcircled{Y}-\textcircled{X}$. And, $\textcircled{X}-\textcircled{Y}-\textcircled{X}$ has a single occurrence: $\textcircled{X}-\textcircled{Y}-\textcircled{X}$. To address this issue, several alternative support counting functions that are anti-monotonic have been defined⁴⁷. An example of such frequency measure is the MNI (Minimum Node Image) support⁸⁸, defined as follows. Consider a subgraph $G_x = (V_x, E_x)$ that has m subgraph isomorphisms (embeddings) in a single connected labeled graph G . For each such subgraph isomorphism $G_i = (V_i, E_i)$, let $f_i : V_x \rightarrow V_i$ be the mapping between vertices in G_x and G_i . The MNI support of G_x in G is defined as $MNI(G_x, G) = \min_{v \in V_x} |\{f_i(v) | i = 1, 2 \dots m\}|$, i.e. the smallest number of vertices in G that match with a vertex in G_x as parts of its embeddings. For example, the MNI support of $\textcircled{X}-\textcircled{Y}$ in the graph $\textcircled{X}-\textcircled{Y}-\textcircled{X}$ is 1, while that of \textcircled{X} is 2, respecting the downward closure property. The MNI support has been used in several papers because it is anti-monotone and can be calculated efficiently^{49,66,88}. An up-to-date overview of other support measures for a single graph is presented in a paper by Meng and Tu⁴⁷.

Three representative algorithms to discover frequent subgraphs in a single graph are SIGRAM⁶⁵, GRAMI⁶⁶ and MuGram⁴⁹. SIGRAM first identifies all vertices that appear at least *minsup* times in the input graph and stores their embeddings. Then, SIGRAM uses these embeddings to extend these patterns to find larger subgraphs and in turn, calculate their embeddings and support. This process is then repeated to find all frequent **subgraphs** using either a breadth-first search or depth-first search. However, a drawback of that approach is that storing embeddings can consume a large amount of memory⁶⁶. To address this issue, GRAMI⁶⁶ does not store all embeddings and only tries to find enough embeddings of any subgraph to prove that it is frequent according to the MNI support. Support calculation are viewed as a constraint satisfaction problem. To avoid finding duplicate subgraphs, GRAMI uses the same canonical labeling as gSpan (called *DFS code*). GRAMI can support both directed and undirected graphs, some simple constraints (e.g. a label cannot appear more than a given number of times in a subgraph), and an approximate version was also designed. But the above algorithms do not support mining patterns in a multi-relational graph (where multiple edges of different types may connect a same pair of nodes such as in a social network)⁴⁹. The MuGram⁴⁹ algorithm was proposed to handle this case. MuGram performs a depth-first search starting from frequent edges to find all frequent subgraphs. When a frequent subgraph is found, its canonical representation is compared with those of previously found **subgraphs** to detect duplicate patterns. Similarly to GRAMI, MuGram applies heuristics to just find enough embeddings of a subgraph to prove that it is frequent. Besides the above studies, to improve efficiency of subgraph mining in a single graph, researchers have also proposed distributed algorithms⁶⁷.

Compressing patterns. Some algorithms have also been designed to find subgraphs that compress a graph⁶¹⁻⁶⁴. The most representative algorithm of this type is SUBDUE⁶¹. It finds subgraphs that compress an input graph well when replacing those subgraphs with single vertices. SUBDUE applies an heuristic beam search to reduce the search space, and **evaluates** compression based on the minimum description length principle. Finding compressing patterns is insightful for some domains but it is not guaranteed that those patterns are frequent⁶⁵. In recent years, various algorithms inspired by SUBDUE have been proposed⁶²⁻⁶⁴.

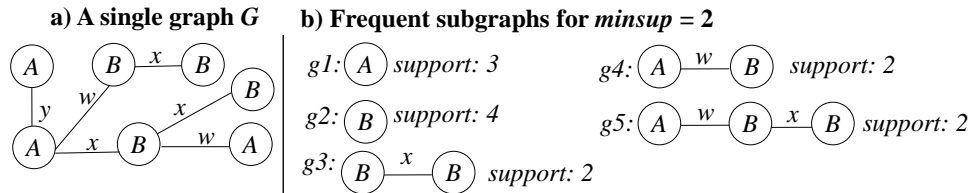


Figure 5: A single graph and frequent subgraphs found for $minsup = 2$

Motifs. A different perspective is adopted in the field of network sciences, where several studies have been done on identifying *network motifs* (subgraphs) that explain the underlying principles of a graph^{68,69}. The key idea in these studies is to compare motifs from an input graph with those in randomly generated graph(s) to identify motifs that are significantly different in the former compared to the latter with respect to a measure⁶⁹. For example, if the frequency measure is used, the aim is to find motifs that appear significantly more often in the input graph than in random graph(s). For such comparison, it is important that the randomly generated graph(s) have structural properties that are similar to those of the input graph (e.g. in terms of average number of edges per vertex). A detailed survey of a dozen measures for network motif discovery was published by Xia et al.⁶⁸. These measures can be generally categorized as statistical measures (e.g. frequency) and structural measures (e.g. number of edges per vertex)⁶⁸. Moreover, some measures are **designed** for using motifs for clustering and classification⁴⁷. Several algorithms have been **proposed** for identifying network motifs⁶⁹. While some of them guarantee an exact solution⁷¹⁻⁷³, others apply a sampling approach and find an approximate solution⁷⁰. The algorithms use several ideas **also utilized in** frequent subgraph mining such as relying on canonical labelings to identify duplicate motifs⁶⁹, performing a pattern-growth exploration to avoid generating non existing candidates^{70,72,73}, and applying symmetry breaking techniques to reduce **the number of** calculations^{70,71}. Studies on network motifs have several important applications for analyzing biological networks and have also been applied in other fields^{68,70,71,73}.

Frequent subtree mining in a single tree. Finding subtrees in a tree database is an important data mining task, which was discussed in the previous subsection. It was shown that some algorithms for mining subtrees in a tree database such as TreeMiner can also be easily adapted to mine frequent trees from a single large tree⁴¹.

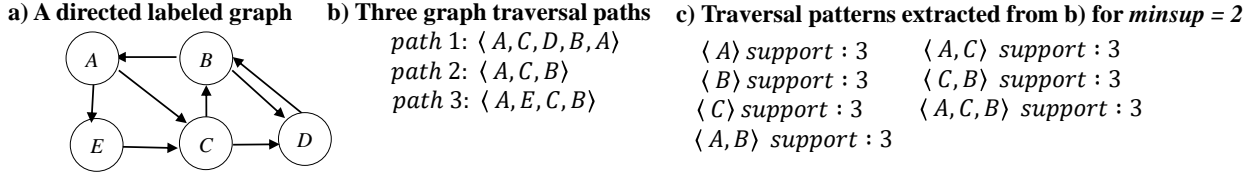


Figure 6: A directed labeled graph, three traversal paths, and some traversal patterns found for $minsup = 2$

Traversal pattern mining. This task consists of identifying all frequent sub-paths (*traversal patterns*) in a database of paths over a single graph. This task has applications such as analyzing webpage access patterns on a website (where pages are nodes and links are edges) and finding common car trajectories (where nodes are road intersections and edges are road segments connecting them)^{56–58}. The basic problem is to find all sub-paths appearing in at least $minsup$ paths, where $minsup$ is a **threshold** set by the users⁵⁷. For example, consider the graph of Fig. 6 a) and three trajectories over this graph depicted in Fig. 6 b). By setting $minsup = 2$, frequent subpaths of Fig. 6 c) are obtained. Sub-paths must preserve the visiting order of nodes but may skip some nodes. To obtain these patterns, a sequential pattern mining (SPM) algorithm¹ can be applied by considering each path as a sequence of symbols. The task of SPM consists of finding all subsequences that frequently occur in a set of sequences. In the context of traversal pattern mining, a SPM algorithm starts by finding frequent paths containing a single node and then recursively extends these frequent paths to find larger paths using a breadth-first or depth-first search, while pruning the search space using the anti-monotonicity of the support. However, a drawback of SPM algorithms is that they ignore the graph structure.

To exploit the graph structure, Nanopoulos and Manolopoulos⁵⁷ proposed three algorithms relying on different search strategies. Then, various extensions of traversal pattern mining have been proposed. The WTPMiner algorithm⁵⁶ mines frequent sub-paths from paths over a graph where weights are associated to edges or vertices to indicate their relative importance. A framework was also proposed to find traversal paths in a directed graph for specific time periods (e.g. to find the most frequent paths to reach a destination during lunch time)⁵⁸.

MINING PATTERNS IN DYNAMIC GRAPHS

Pattern mining algorithms reviewed in the previous section have many applications but do not consider the time dimension. Designing algorithms that consider time is desirable for many applications but is challenging for three main reasons. First, problem definitions and interestingness measures for selecting patterns in a static graph must often be redefined to handle time. But considering novel or adapted measures requires to study their properties and design novel strategies to handle them during the mining process^{79–81,102}. Second, while some algorithms can be extended in a relatively simple way to consider time^{74,75}, other algorithms are not easily adapted because optimizations, data structures and search strategies for mining patterns in static graphs are sometimes too specialized, and thus hard to extend. In such case, novel techniques must be designed. Third, adding the time dimension to an existing pattern mining problem generally makes the problem much more difficult. For instance, increasing the number of timestamps for some problems can greatly increase the size of the search space^{74,75,102}.

Several algorithms have been designed to efficiently discover patterns revealing how graph(s) evolve over time. These studies often draw inspiration from those on static graphs as some challenges remain the same but also include many novel ideas. The next three subsections review studies on discovering patterns in three main types of data: (1) a dynamic graph, (2) a database of dynamic graphs, and (3) a dynamic attributed graph. Then, the last subsection discusses other extensions. For the convenience of the reader, Fig. 7 shows a tree indicating the main data types and pattern types discussed in this section. Relationships between studies on dynamic graphs and those on static graphs will be discussed through the section.

MINING PATTERNS IN A SINGLE DYNAMIC GRAPH

Several studies have been done on finding patterns in a single dynamic graph. Researchers that have contributed to these studies are from various research fields such as data mining, statistics and network science. Several names have been used with a more or less similar meaning to refer to a graph that changes over time such as dynamic graph, dynamic

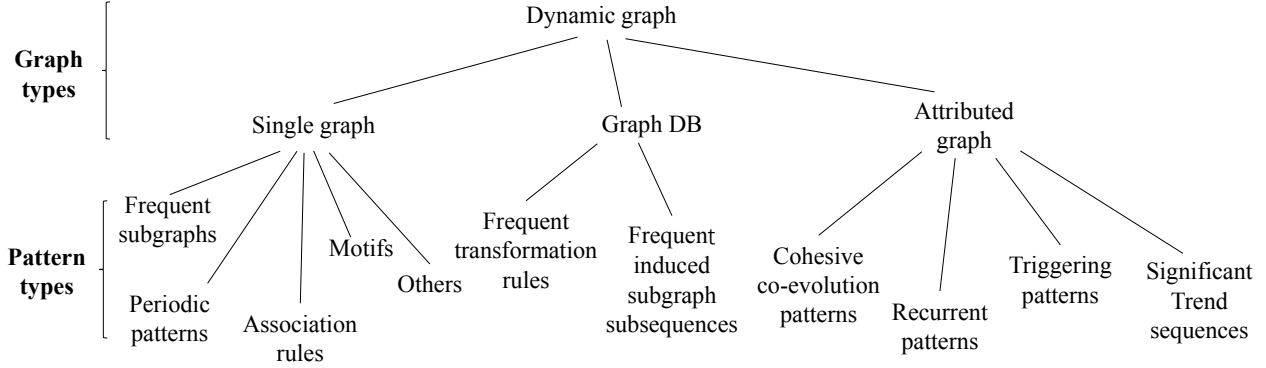


Figure 7: Main types of patterns discovered in dynamic graphs

network, evolving graph, temporal graph, graph sequence, time-series of graph, and time-varying graph. There are two main research sub-areas: (1) techniques to analyze a social network with a focus on community detection^{52,53}, and (2) pattern mining techniques to identify interesting patterns^{74,79,82,89}. As explained in the introduction, this survey focuses on techniques for mining patterns in dynamic graphs. To know more about community detection techniques, the interested reader may refer to recent surveys on this topic^{52,53}.

This subsection first introduces a formal definition of a dynamic graph and important related terms. Then, a taxonomy of the main types of dynamic graphs is presented based on what is changing in a graph. Lastly, key studies on pattern mining in a dynamic graph are discussed.

Single dynamic graph. Formally, a *single dynamic graph* is a sequence of labeled graphs $G = \langle G_1, G_2, \dots, G_T \rangle$ in which a graph G_t represents the state of the dynamic graph at time t . The graph $G_t = (V_t, E_t, \lambda_t)$ consists of a set of **vertices** V_t at time t , a set of edges $E_t \subseteq V_t \times V_t$ at time t , and a labelling function $\lambda_t : V_t \cup E_t \rightarrow \mathbb{R}$ mapping edges and vertices of G_t to labels (represented by numbers or literals) at time t . A graph at a time t is also called a *snapshot* of the dynamic graph G . For the sake of brevity, this subsection will refer to a single dynamic graph as a dynamic graph.

A taxonomy of dynamic graphs. The above definition is a generic definition of dynamic graph. Different variations of this definition are considered in the literature for the needs of different applications. They can be categorized based on their structure as discussed in the previous section (e.g. directed/undirected graphs, weighted graphs, trees,

and attributed graphs), but more importantly they can be described in terms of what is changing in a graph over time. Two main types of evolutions (changes) have been mainly considered. A *topological evolution* refers to changes in a graph topology such as adding and removing vertices and edges. A *label evolution* refers to changes of labels associated to edges and/or vertices.

Based on these concepts, three types of dynamic graphs can be identified: (1) dynamic graphs with only topological evolution^{74,76,79,80,82–84}, (2) dynamic graphs with only label evolution⁸⁹, and (3) dynamic graphs with both topological and label evolution^{75,86,87,90,91}. Studies on the third type of dynamic graphs mostly focus on a type of graph called *dynamic attributed graphs*, which will be discussed in a following subsection. Studies on such graphs mainly aim to analyze the evolution of multiple attributes and their relationships over time. Fig. 8 provides examples of these different types of dynamic graphs, which evolve over three timestamps t_1 , t_2 and t_3 . Although only vertices are labelled in this example, edge labels can also be considered. Note that besides these main types of graphs, some authors also use the terms *evolving graph* or *streaming graph* to refer to a graph that is continuously updated with new snapshots^{76,113,114}. Mining patterns in a streaming graph is more challenging as results must often be updated in real-time as new data arrives, and if the stream is infinite, data can only be read once.

The following paragraphs discuss the main types of patterns that are discovered in a dynamic graph.

Mining frequent subgraphs in a dynamic graph. FSM is the most well-studied problem for mining patterns in a static graph. Several studies have been done on mining frequent subgraphs in a single large graph^{47,48} (see previous section). And in general, FSM has inspired most **studies** on pattern mining in graphs. Nowadays, with the **increasing amount** of data, more and more data has rich temporal information that can be modeled as dynamic graphs. Hence, FSM was naturally extended to analyze dynamic graphs⁷⁴.

The first formal definition of FSM in a dynamic graph was proposed by Borgwardt et al.⁷⁴. They designed an algorithm, which internally represents a dynamic graph (a sequence of snapshots) as a single *union graph*. In this graph, each edge is labeled with a bit string (a sequence of 0s and 1s), which describes the edge’s status over time. For instance, an edge

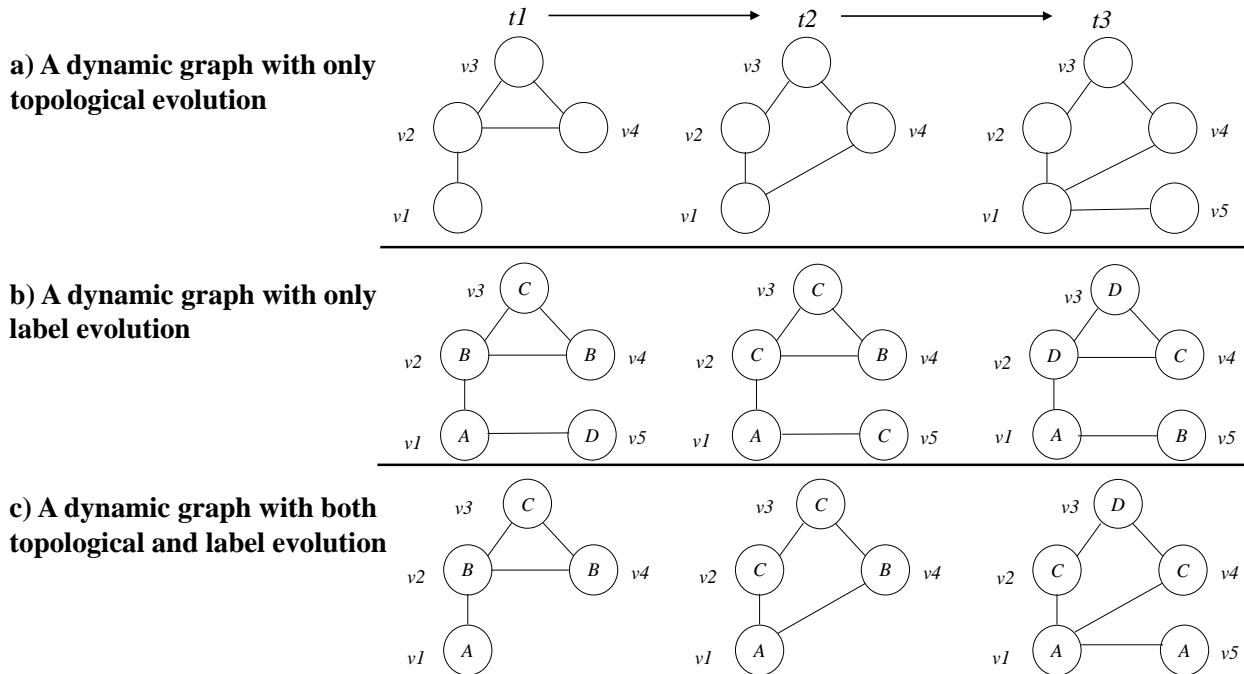


Figure 8: Three types of dynamic graphs

labeled as "010" means that this edge did not exist at time 1, appeared at time 2, and did not exist at time 3. This graph representation is very similar to that used for FSM in a single large static graph^{47,48}. The difference is that strings are added to edges to store information about edge evolution. To find frequent subgraphs, Borgwardt et al. extended the GREW algorithm⁷⁸, which is designed for FSM in a single graph. The modified algorithm is called Dynamic GREW⁷⁴. It is a **breadth-first** search algorithm, which generates large dynamic subgraphs by joining smaller frequent dynamic subgraphs previously found. To find frequent subgraphs, Dynamic GREW generates more candidates than GREW because many frequent substrings may need to be considered for a same subgraph. Moreover, isomorphism checking is more complex in Dynamic GREW because substring checking must be performed when comparing two subgraphs. The output of Dynamic GREW is a series of subgraphs with bit strings representing temporal behaviors over consecutive timestamps that frequently occurred in the input dynamic graph. The algorithm can output either synchronous subgraphs (each occurrence must start at the same time) or asynchronous subgraphs (occurrences of a subgraph are not required to start at the same time). Patterns found using Dynamic GREW can help understanding how graph edges evolve (appear or disappear)⁷⁸. It was shown to

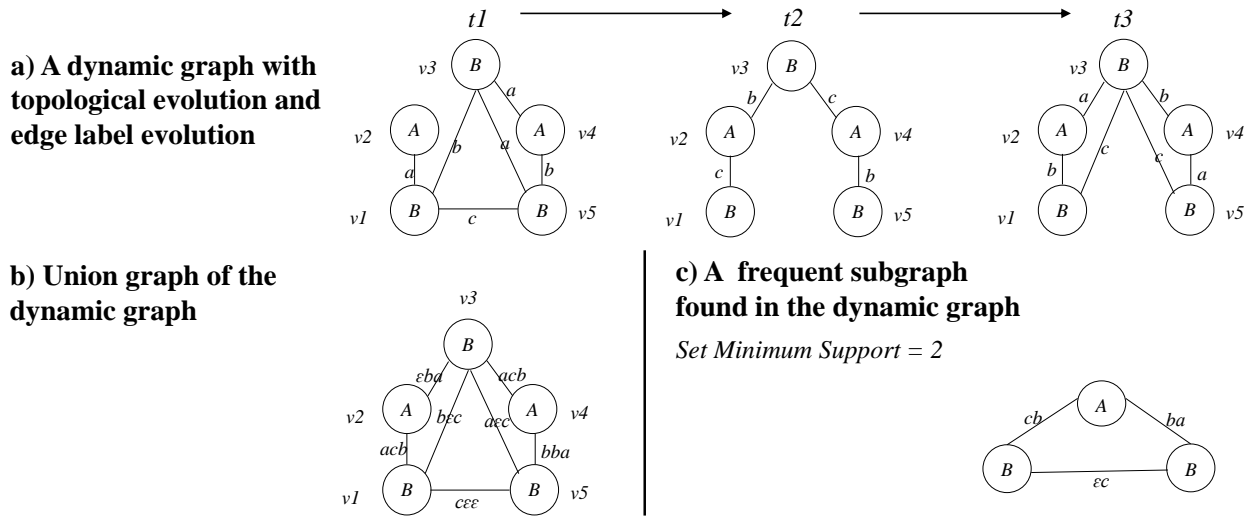


Figure 9: A dynamic graph, the corresponding union graph, and a frequent subgraph

reveal interesting patterns in e-mail interactions between employees of a company, which were modelled as a dynamic graph. But Dynamic GREW has three key limitations: it uses a greedy heuristic approach like GREW that trades completeness of results for speed, it considers that edges are unlabelled, and it can have long runtimes⁷⁵.

Wackerseuther et al. improved upon that work in several ways⁷⁵. They considered a richer graph model where each graph edge is annotated with a string that contains edge labels for each timestamp, rather than only 0s and 1s. An edge label is either a symbol (e.g. a , b and c) to describe an edge, or a special ϵ label indicating that an edge did not exist at the corresponding timestamp. Before mining patterns, a union graph is created by combining the information of all timestamps. For instance, Fig. 9 a) shows a dynamic graph and Fig. 9 b) shows the corresponding union graph representation.

Based on that graph representation, Wackerseuther et al. designed a framework to discover dynamic frequent subgraphs where edge strings indicate label evolution. For instance, Fig. 9 c) shows a frequent subgraph found for a minimum support of 2. To discover such patterns, Wackerseuther et al. designed a generic framework that first applies a traditional FSM algorithm to find all frequent subgraphs in the static input graph, while ignoring edge strings. Then, the framework searches in each subgraph to find dynamic frequent subgraphs with edge strings. This is done by checking all embeddings (occurrences) of a subgraph

to find frequently appearing edge substrings. To efficiently compare graph occurrences, a canonical edge order similar to the one used by gSpan² was defined, and a suffix-tree is used to find the frequent longest substrings in linear time. Advantages of this approach are that it is efficient, it guarantees finding all the desired patterns, it relies on existing FSM algorithms, and it considers label evolution. Wackerseuther et al. applied their approach to find interesting patterns in a single graph representing protein-protein interactions from yeast, where gene expression levels vary over time⁷⁵.

Contrary to the above two algorithms, which focus on finding frequent subgraphs representing dynamic behaviors, Abedlhamid et al.⁷⁶ proposed to identify subgraphs that are frequent in each snapshot of a dynamic graph. To perform this task, a simple solution is to apply a traditional FSM algorithm to each snapshot and then to combine the results. However, this is inefficient because it does not take advantage of the fact that consecutive snapshots are generally similar. To address this problem, Abedlhamid et al. developed an efficient incremental algorithm named IncGM+. Unlike the previous two algorithms, it does not model a dynamic graph as a single graph. It instead processes each new graph one by one using an approach inspired by the Moment algorithm for frequent itemset mining in a stream⁷⁷. This approach consists of mining frequent subgraphs in the first snapshot, and then to only update the "fringe" patterns (those who are at the boundary between frequent and infrequent patterns) when a new transaction (graph) is processed. This allows to reduce the computational cost of the mining task. IncGM+ also introduces three other performance optimizations. First, instead of storing all embeddings of each fringe subgraph, IncGM+ keeps a minimal number of embeddings for each fringe subgraph which can significantly reduce runtimes and memory usage. Second, a data structure was proposed to efficiently maintain embedding lists to perform fast support calculations. Third, graphs can be processed by batches to further improve efficiency. IncGM+ was applied to mine patterns in (1) Twitter data where a node is a person and a link indicates that a person follows another, (2) a graph where nodes are US patents and links are citations, (3) a similar graph for citations between research papers, and (4) a dynamic graph representing communications between users of **an instant messaging software**.

Mining periodic patterns in a dynamic graph. The second main type of patterns

that has been studied in a single dynamic graph is periodic patterns^{79–81}. Lahiri et al.⁷⁹ introduced this concept to find repeating interactions in a dynamic graph. For example, one could study a dataset about cellphone calls between people to find interactions that are repeating over time. That study considered finding patterns in a dynamic graph, where snapshots are equally spaced in time (a time-series of graphs), edges representing interactions between vertex are directed or undirected, and vertex labels are unique. This latter assumption greatly simplifies subgraph mining problems as a graph can be represented as a set of integers (each representing the presence of an edge or vertex), which allows to efficiently perform isomorphism checking using the subset operator \subseteq . In that study, a subgraph is considered to be *periodic* in a time interval if it appears some minimum number of times in that interval, every consecutive occurrences of the subgraph in that interval is separated by the same amount of time, and the time interval cannot be extended while preserving the previous properties. Furthermore, to eliminate redundancy, it was proposed to only discover closed subgraphs³², and to find a minimal set of patterns that covers all periodic occurrences of all periodic subgraphs. An example of periodic pattern in the time interval $[t_1, t_5]$ is shown in Fig. 10 c), which appears at timestamps t_1 , t_3 , and t_5 of the dynamic graph of Fig. 10 a). Fig. 10 b) shows two frequent subgraphs that are found for a minimum support of 3 where the second one is filtered because it is not periodic. To assess the periodic behavior of a pattern, the *Purity* measure is used, which filters out periodic patterns that occur too frequently in a time sub-interval. Moreover, to allow finding periodic patterns that are not eqally spaced in time, it was proposed to apply smoothing to the snapshots as a preprocessing step. An algorithm named PSEMiner was designed to find the desired patterns in polynomial time. PSEMiner was applied to discover interesting repeating interactions in four types of dynamic graphs: (1) e-mail communications between employees of a business, (2) social interactions of plain zebras in Kenya, (3) cellphone interactions between university students, and (4) associations between celebrities that are photographed together in the press⁷⁹. Several other algorithms have then been proposed to improve the efficiency of mining periodic subgraphs in a dynamic graph^{80,81}. One of those studies applied periodic subgraph mining to analyze social interactions on the Youtube website to find patterns indicating how users join groups, and on data from the Facebook social network to study how users interact through wall

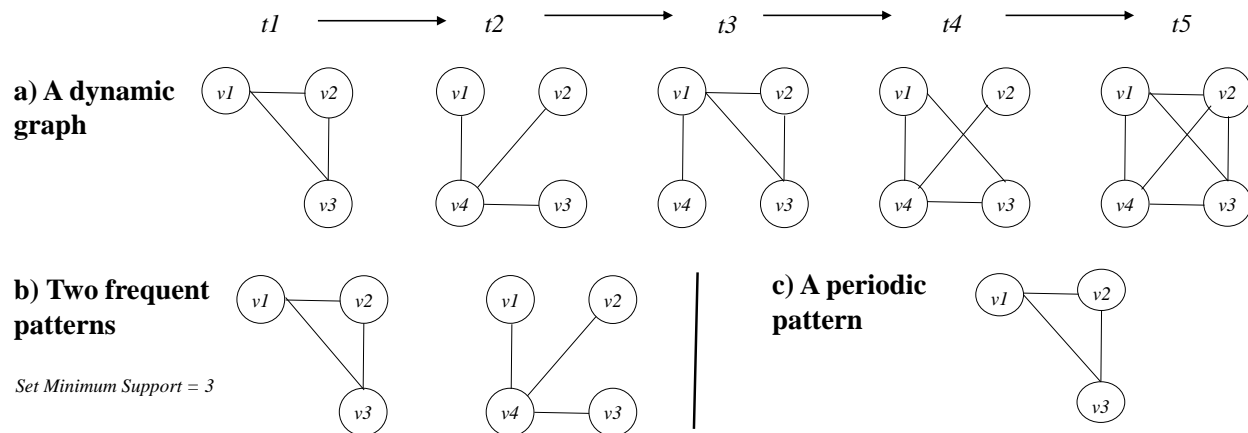


Figure 10: A dynamic graph, some frequent patterns and a periodic pattern

messages⁸⁰.

Mining association rules in a dynamic graph. The third main type of patterns found in a dynamic graph is rules^{82–84,86,87}. Such studies draw inspiration from the traditional data mining task of discovering association rules in transaction databases¹⁰.

Berlingerio et al.⁸² introduced an algorithm named GERW to discover a novel type of rules called graph evolution rules (GER) in a dynamic graph. Similar to previous **algorithms**, GERW first transforms a graph sequence into a single union graph to then mine patterns **in that graph**. In the union graph of GERW, a label on each edge indicates the timestamp of its first appearance. Then, a FSM algorithm for a large static graph can be applied on the union graph to find frequent subgraphs⁸⁸. However, the constraint that labels should be identical for two embeddings of a subgraph should not be enforced because edge labels represent timestamps instead of attribute values. Thus, the GERW algorithm first mines relative-time patterns, that is subgraphs having embeddings that are structurally isomorphic and where edge labels (timestamps) differ only by a constant. For instance, Fig. 11 a) shows a union graph, and Fig. 11 b) shows two embeddings of a subgraph where timestamps differ only by 1 time unit. These embeddings are said to belong to the same equivalence class though their edge labels are not the same. After finding all frequent patterns, the algorithm extracts graph evolution rules from these patterns. A rule has the form $body \rightarrow head$ where $head$ is a frequent pattern and $body$ is obtained by removing the edges having the largest timestamps such that the subgraph remains connected. A rule is output if the confidence between $body$

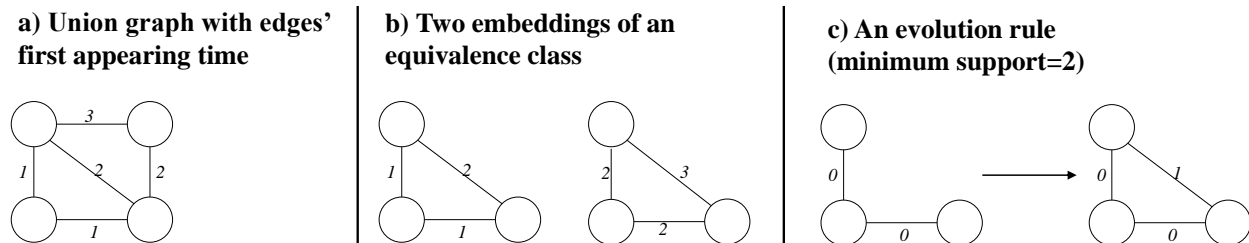


Figure 11: A graph evolution rule found in a dynamic graph

and *head* satisfies a minimum confidence threshold. GERW was applied to analyze (1) friendships links between users on the Flickr and Yahoo 360 websites, and (2) co-authorship of research papers on the Arxiv and DBLP websites⁸². **Thereafter**, other algorithms were proposed based on the same framework to mine rules in different kinds of dynamic graphs and/or to address limitations of GERW such that it does not consider edge **deletion** and edge relabeling.

Leung et al.⁸³ proposed to mine Link Formation Rules (LFR) in directed dynamic graphs with multiple edge labels. LFR are discovered by enforcing more strict constraints on rules than GERW does. A LFR indicates how a subgraph is extended by adding a new link from a start node to an end node, with the restriction that every node in the subgraph must have a direct link to the start node or end node. A GER is equivalent to a set of LFR having the same base patterns and where edge timestamps respect the GER's definition. However, these two studies focus on finding association rules in two different types of dynamic graphs. Ozaki et al.⁸⁴ then adapted the concept of LFR for simple dynamic graphs. In that work, a LFR represents the addition of an undirected edge to a base pattern (subgraph). Furthermore, two relationships between LFR are defined. First, two LFR are said to be *correlated* if their base pattern is the same and the two extra edges always appear together when the base pattern appears. Second, two LFR are said to be *contrast* when their base pattern is the same and always one or the other extra edge appears when the base pattern appears. To reduce redundancy among patterns and improve pattern mining efficiency, the study only consider δ -closed subgraphs⁸⁵ as base patterns. The above studies have been applied to analyze interactions between users of a product review website named Epinions and a mobile social network called MyGamma⁸³, as well as e-mail communications and student

interactions⁸⁴.

All the above rule mining algorithms are useful but have the limitation that they cannot handle dynamic graphs with edge or vertex deletion and relabeling. This is because their union graph structure only considers the timestamp of the first appearance of each edge. To address this limitation, two methods were proposed that keep more information about dynamic graphs by transforming them into a database of union graphs. The first method named DGRMiner was introduced by Vaculik⁸⁶. It can find rules either in a dynamic graph or in a database of dynamic graphs, where any type of changes may appear and where edges may be directed or undirected. To mine patterns in a dynamic graph containing n snapshots, DGRMiner first transforms the dynamic graph into a set of $n - 1$ union graphs that will then be treated as a set of static graphs for mining patterns. The k -th union graph contains the union of the first k snapshots and indicates changes between the k -th and the $(k + 1)$ -th snapshot using relative timestamps. After creating union graphs, a modified frequent subgraph mining algorithm inspired by gSpan is applied on them to mine rules that have a high support and confidence (an estimation of the rule’s conditional probability). The algorithm applies a depth-first search and detects duplicate subgraphs using a modified version of gSpan’s DFS code, which stores information about timestamps and edge directions. DGRMiner was **applied** to analyze e-mail interactions and student resolution proofs in propositional logics. The second method proposed by Scharwachter et al.⁸⁷ is named EvoMine and supports dynamic graphs with edge **deletion** and relabeling. EvoMine has many similarities with DGRMiner, the main difference being that EvoMine relies on bit strings to represent the status of edges and vertices for FSM. Moreover, EvoMine creates union graphs only from pairs of consecutive snapshots. The EvoMine implementation uses the gSpan algorithm as its core for finding frequent subgraphs. Scharwachter et al. applied EvoMine to analyze research paper co-authorship relationships on DBLP as well as interactions between users on the EPinion website.

Mining motifs in a dynamic graph. The fourth main type of patterns found in a dynamic graph are *motifs*, which are small patterns of interconnections that occur more frequently in a large graph than they would in a randomized or reference graph. Motifs thus characterize a graph’s structure and can reveal its design principles. The concept of *motifs*

was initially proposed by the network science community in the context of static graphs to analyze various types of data such as biological, ecological and web data^{111,112} (as discussed in the previous section). Then, it was extended for dynamic graphs.

Jin et al.⁸⁹ proposed to extract *trend motifs* from a labeled dynamic graph where the topology is fixed, vertices have weights, and only weights are changing over time. A *trend Motif* is defined as a connected subgraph consisting of nodes that show a trend (increasing or decreasing weights) over some time span. The assumption is that a change of a node’s weight is not an isolated event and it tends to be correlated with that of its neighbors. Jin et al. designed an algorithm to find all *frequent trend motifs* that are over-represented in a dynamic graph. The algorithm first finds maximal time intervals where trends appear for each vertex. Then, a **depth-first search** is applied to find induced subgraphs combining either increasing or decreasing trends for several vertices, and finally a breadth-first search is performed to generate frequent motifs combining several motif occurrences. As in many other studies, a canonical code is calculated for each pattern to detect duplicates. The algorithm revealed interesting patterns in stock market and micro-array data.

Ahmed and **Karypis then** proposed to mine coevolving relational motifs (CRM)⁹⁰ in a dynamic labeled graph where labels can change, and edges may be added or removed over time. A CRM is set of motifs (sets of vertices) that **change** in a consistent way over time. The biggest difference between *trend motifs* and CRM is that the latter can capture multiple trends over several timestamps (a consecutive sequence of motifs), while each node in a *trend motif* can only have a single trend (increase or decrease) over a time span. An algorithm named CRMminer was proposed to extract all frequent CRM. The algorithm first finds frequent motifs occurrences using a depth-first search, which relies on a modified version of gSpan’s DFS code to provide a canonical labelling for CRM. Then, a traditional sequential pattern mining algorithm¹ performing a depth-first search is applied to combine motifs occurrences into frequent sequences. CRMminer was applied to analyze real data including cell culture bioprocess data collected from bioreactors and years of sales data from retail stores. To further improve the efficiency and reduce redundancy, Ahmed and Karypis proposed to mine coevolving induced relational motifs (CIRM)⁹¹, **by adding** the constraint of finding only induced subgraphs for CRM. An algorithm efficient algorithm named CIRMiner

was designed for this task.

The above motif definitions can only capture temporal changes that appear between two consecutive snapshots, which is very restrictive and can miss some important changes. Paranjape et al. proposed to find another type of motifs in dynamic graphs with only topological changes⁹². The proposed method creates an union graph to represent dynamic graphs where edge labels store all appearing timestamp of edges. Considering the edge appearing order, a δ -temporal motif is defined as an ordered sequence of edges within a time window δ such that the induced static graph of its edges is connected. The authors developed algorithms to count the number of instances of δ -temporal motifs. Though the algorithms are fast and can scale to large graphs, they can only find 2-node motifs and 3-node, 3-edge star or triangle motifs. Moreover, the algorithms only count the number of instances of each motif but do not enumerate them, and do not use search space pruning strategies (differently from most algorithms reviewed in this survey). The algorithms were applied on various datasets of user interactions including cryptocurrency transactions, SMS and phone calls.

Other patterns. Besides the four main types of patterns described above, a few other types of patterns have been studied. The next paragraphs discuss four of them.

Robardet et al. proposed a constraint-based pattern mining approach⁹³ to find dense and isolated subgraphs. Finding such patterns is similar to the task of community detection. Dense and isolated subgraphs are detected in each snapshot and their evolution is studied over time in terms of formation, dissolution, growth, and stability. The approach was utilized to analyze the usage of public shared bikes and interactions among students.

Another type of evolution patterns was proposed by Ahmed and Karypis⁹⁴. That study focuses on the evolution of Induced Relational Subgraphs (IRS). An IRS is an induced subgraph where vertices, edges and labels remain unchanged for a long time (a parameter). They designed an algorithm to find *evolving induced relational states*, which are patterns having the form $S_1 \rightarrow S_2 \rightarrow S_3$ where the notation S_i denotes an IRS. The algorithm was applied to analyze e-mail communications, patent citations as well as the import and export relations of 192 countries.

Bogdanov et al.⁹⁵ designed an algorithm named MEEEDEN to find patterns in a spe-

cial type of dynamic graphs having binary edge labels $\{-1, 1\}$. The algorithm finds the highest-scoring temporal subgraphs, called *Heaviest Dynamic Subgraph* (HDS). The score of a subgraph is the sum of its edge weights. A HDS in a dynamic graph representing traffic movement can for example indicate the most congested connected streets during some periods of time. While finding high-scoring subgraphs in a static graph is easy, the problem is difficult in a dynamic graph because various time sub-intervals must be considered. To find patterns efficiently, MEEDEN utilizes pruning techniques to ignore unpromising time sub-intervals. The algorithm was applied to analyze transportation, social media and communication graphs.

In another study, Yang et al.⁹⁶ proposed to detect areas of an undirected dynamic graph that are frequently changing. This is useful for several applications such as analyzing traffic data as frequently changing areas may face frequent traffic jam. Yang et al. proposed to mine the most frequently changing components of a dynamic graph, where components are dense and connected subgraphs. Using that proposed approach, patterns were mined in internet traffic data, comments on a news website, and shopping data.

Summary. This subsection has reviewed algorithms for mining various types of patterns in a single dynamic graph. Several of those algorithms use a union graph-based representation, but encode time in different ways and utilize different measures, according to the task^{82,86,87,92}. While some studies directly apply a traditional FSM algorithm on a union graph with some preprocessing or post-processing techniques^{75,87}, others apply custom algorithms. Moreover, the reviewed algorithms generally adapt concepts used for static graph mining such as that of using a canonical labeling to detect duplicate patterns^{75,87,89,90}, that of using a breadth-first search by combining smaller patterns to generate larger patterns^{82,89}, or that of using a pattern-growth approach to avoid generating candidates that do not exist in the database⁸⁶. Some algorithms such as CRMminer also find subgraphs and then use a sequential pattern mining algorithm to combine subgraphs into sequences⁹⁰. Besides, some motif discovery algorithms utilize more of a brute-force approach by simply evaluating all patterns without search space pruning strategies⁹².

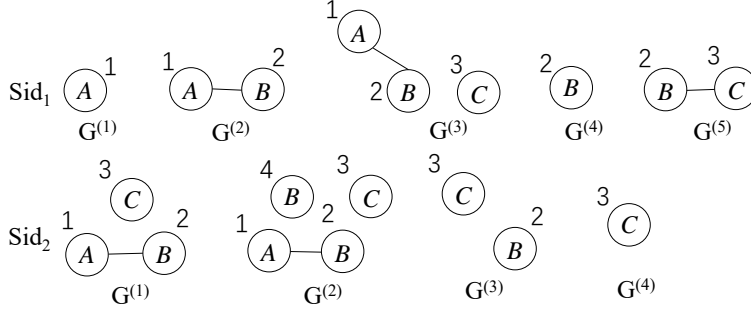


Figure 12: A graph sequence database containing two graph sequences

MINING PATTERNS IN A GRAPH SEQUENCE DATABASE

Studies reviewed in the previous section have considered mining patterns in a single dynamic graph (a graph sequence). This section reviews studies on mining patterns in a graph sequence database, that is patterns appearing in several graph sequences. The section first introduces important definitions, and then describes the main types of patterns that are found in a graph sequence database. Graph sequence databases are found in several domains such as in social networks and gene networks. For example, in a social network, each person may have a graph sequence representing the evolution of its relationships with others^{108–110}.

Formally, a *graph sequence database* GDB is a set of tuples $\langle sid, d \rangle$, where *sid* is a unique sequence **identifier (ID)** and *d* is a graph sequence. Recall that a graph sequence $d = \langle G^{(1)}, G^{(2)}, \dots, G^{(m)} \rangle$ is an ordered list of graphs (as defined in the previous subsection).

For example, Fig. 12 shows a graph sequence database that contains two graph sequences having the IDs *Sid1* and *Sid2*. In sequence *Sid1*, there are five elements (graphs), unique IDs (1, 2, 3 and 4) are used to refer to vertices, and each vertex has a label (*A*, *B* or *C*).

Real-life graph sequence databases are often large and sparse. Representing a graph sequence database as a list of graphs can require a considerable amount of space because many parts of a graph may remain unchanged over time. Storing these unmodified elements for consecutive timestamps results in storing redundant information. Using this simple representation can also make it time and memory consuming for mining knowledge in a large and sparse graph sequence database. To compactly represent a graph and facilitate its analysis, several methods have been proposed such as Graph Grammar¹⁰⁶. However, this latter was designed to be applied to a single graph. To compactly and efficiently represent a

graph sequence, Akihiro et al.¹⁰⁴ focused on the differences between each pair of successive graphs and proposed a novel graph grammatical framework which describes a graph sequence as the application of several transformation **rules to** an initial graph.

A transformation rule indicates a change (e.g. adding an edge or changing a label) that appeared between two consecutive snapshots of a graph sequence. In other words, a rule indicates how a snapshot has changed from a timestamp to the next. A graph sequence can then be described by the successive application of transformation rules to the initial snapshot. Such sequence of transformations is called a transformation rule sequence. A more formal definition is given below¹⁰⁴.

Intrastate sequence. Let there be a graph sequence $d = \langle G^{(1)}, G^{(2)}, \dots, G^{(m)} \rangle$ containing m snapshots. Consider two graphs $G^{(j)}$ and $G^{(j+1)}$ from consecutive timestamps that have m_j differences. The differences between these two graphs can be described by a sequence of intermediate graphs $s^{(j)} = \langle G^{(j,1)}, G^{(j,2)}, \dots, G^{(j,m_j)} \rangle$ called intrastate sequence such that each intermediate graph has only a single difference with the preceding intermediate graph, and where $G^{(j,1)} = G^{(j)}$ and $G^{(j,m_j)} = G^{(j+1)}$. Based on this idea, the sequence d can be represented as an *interstate sequence* defined as $\langle s^{(1)}, s^{(2)}, \dots, s^{(m-1)} \rangle$.

Transformation rule (TR). A transformation rule that transforms a graph $G^{(j,k)}$ into another $G^{(j,k+1)}$ of an intrastate sequence is denoted as $\langle tr_{[o_{jk}, l_{jk}]}^{(j,k)} \rangle$ where tr is a literal indicating the transformation type, o_{jk} is the unique ID of the vertex or edge to which the transformation is applied, and l_{jk} is a label that the transformation assigns to a vertex or an edge. Six types of transformation rules are considered, named and denoted as follows: vertex insertion $vi_{[u,l]}^{(j,k)}$, vertex deletion $vd_{[u,]}^{(j,k)}$, vertex relabeling $vr_{[u,l]}^{(j,k)}$, edge insertion $ei_{[(u_1, u_2), l]}^{(j,k)}$, edge deletion $ed_{[(u_1, u_2),]}^{(j,k)}$ and edge relabeling $er_{[(u_1, u_2), l]}^{(j,k)}$.

Transformation rule sequence. Based on the concept of transformation rule, an intrastate sequence $s^{(j)} = \langle G^{(j,1)}, G^{(j,2)}, \dots, G^{(j,m_j)} \rangle$ of a sequence d can be represented by a sequence of transformation rules $seqtr(s^{(j)}) = \langle tr_{[o,l]}^{(j,1)}, tr_{[o,l]}^{(j,2)}, \dots, tr_{[o,l]}^{(j,m_j-1)} \rangle$. And an interstate sequence d' can be represented as $seqtr(d) = \langle seqtr(s^{(1)}), seqtr(s^{(2)}), \dots, seqtr(s^{(m-1)}) \rangle$.

Inclusion relation. Let there be an intrastate sequence $s^{(j)}$ of a graph sequence d , and another $s^{(h)}$ of a graph sequence d' . Their transformation rule sequences are $seqtr(s^{(j)}) = \langle tr_{[o,l]}^{(j,1)}, tr_{[o,l]}^{(j,2)}, \dots, tr_{[o,l]}^{(j,m_j-1)} \rangle$ and $seqtr(s^{(h)}) = \langle tr_{[o,l]}^{(h,1)}, tr_{[o,l]}^{(h,2)}, \dots, tr_{[o,l]}^{(h,m_h-1)} \rangle$, respectively. Iff

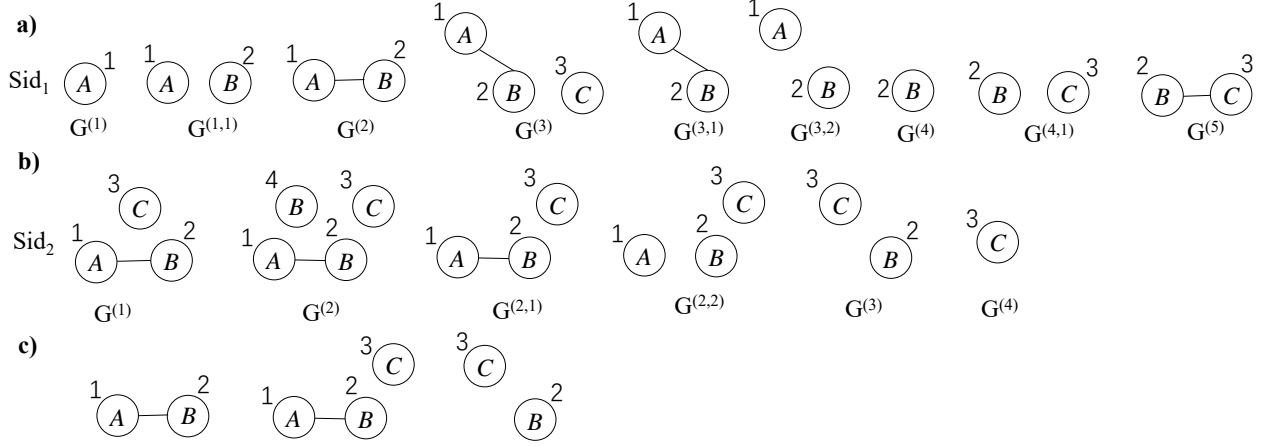


Figure 13: Two intrastate sequences and one of their subsequences

$\forall tr_{[o,l]}^{(h,r)} \in seqtr(s^{(h)}), \exists tr_{[o,l]}^{(j,k)} \in seqtr(s^{(j)})$ such that $tr_{[o,l]}^{(h,r)} = tr_{[o,l]}^{(j,k)}$, then $seqtr(s^{(h)}) \subseteq seqtr(s^{(j)})$. An inclusion relation is similarly defined for two graph sequences as $seqtr(d) = \langle seqtr(s^{(1)}) \text{ and } \dots, seqtr(s^{(m-1)}) \rangle$, $seqtr(d') = \langle seqtr(s^{(1)}) \dots, seqtr(s^{(m'-1)}) \rangle$. Iff there exist integers $1 \leq j_1 \leq \dots \leq j_{m-1} \leq m-1$ such that $seqtr(s^{(h)}) \subseteq seqtr(s^{(j_h)})$ for $h = 1, 2, \dots, m'-1$, then $seqtr(d') \subseteq seqtr(d)$.

The **support** (occurrence frequency) of a transformation rule sequence in a graph sequence database GDB is denoted and defined as $sup(seqtr(d)) = \frac{|\{d_i | d_i \in GDB, seqtr(d) \subseteq seqtr(d_i)\}|}{|GDB|}$.

For example, Fig 13 a) and b) show the intrastate sequences corresponding to sequence Sid_1 and Sid_2 of Fig 12, respectively, and Fig 13 c) shows one of their subsequences. Transformation rule sequences of Sid_1 and Sid_2 are $seqtr(Sid_1) = \langle vi_{[2,B]}^{(1,1)} ei_{[(1,2),\cdot]}^{(1,2)} vi_{[3,C]}^{(1,2)} vd_{[3,\cdot]}^{(3,1)} ed_{[(1,2),\cdot]}^{(3,2)} vd_{[1,\cdot]}^{(3,3)} vi_{[3,C]}^{(4,1)} ei_{[(2,2),\cdot]}^{(4,2)} \rangle$ and $seqtr(Sid_2) = \langle vi_{[4,B]}^{(1,1)} vd_{[4,\cdot]}^{(2,1)} ed_{[1,2,\cdot]}^{(2,2)} vd_{[1,A]}^{(2,3)} vd_{[3,C]}^{(3,1)} \rangle$, respectively. The subsequence c) is represented by the transformation rule sequence $\langle vi_{[3,C]}^{(1,1)} ed_{[1,2,\cdot]}^{(2,1)} vd_{[1,A]}^{(2,2)} \rangle$ and has a support of $2/2 = 1$.

Mining frequent transformation rules. Given a minimum support threshold, Akihiro et al.¹⁰⁴ proposed to mine the transformation rule sequences whose support is no less than a user-defined minimum support threshold, called Frequent Transformation Rule Sequence (FTRS). To efficiently find the FTRSs, Akihiro et al. designed a depth-first pattern growth algorithm named FTSMiner based on a sequential pattern mining algorithm named PrefixSpan¹⁰⁷. FTSMiner relies on the anti-monotonicity property of the support measure to reduce the search space, which states that if $seqtr(d_1) \subset seqtr(d_2)$ then

$sup(seqtr(d_1)) \geq sup(seqtr(d_2))$. The algorithm considers that each transformation rule is an item in a sequence to transform the problem of mining frequent transformation sequences into that of frequent sequential pattern mining¹. Akihiro et al.¹⁰⁴ also extended the above problem to mine FTRSs from connected graphs. Given a minimum support threshold, the connected graphs of all graph sequences in a GDB are **enumerated**. Then, all frequent connected subgraphs of those connected graphs representing graph sequences are generated by applying the traditional GASTON algorithm³⁵ for frequent subgraph mining. Finally, FTRSs are mined from each connected subgraph. FTRSs are interesting because they reveal the evolution sequences that frequently occur in a GDB. Thus, FTRSs may be used to understand a graph and do predictions in some applications. The algorithms were applied to analyze e-mail communications and interactions between persons.

Mining frequent, relevant induced subgraph subsequences. A limitation of transformation rule sequence mining is that changes in graphs have to be small or gradual and that there should not be too many vertices, otherwise the performance of the algorithm decreases. To address this problem and discover long sequences in large graphs, Akihiro and Takashi¹⁰⁵ proposed an algorithm named FRISSMiner to mine another type of patterns called *frequent relevant induced subgraph subsequence* (FRISS).

A graph $G'(V', E', L', l')$ is called a subgraph of $G(V, E, L, l)$ denoted as $G' \subseteq G$ if three conditions are satisfied, which are 1. $(\phi(v_1), \phi(v_2)) \in E$, if $(v_1, v_2) \in E'$, 2. $l'(v) = l(\phi(v))$, and 3. $l'((v_1, v_2)) = l((\phi(v_1), \phi(v_2)))$, where ϕ is a mapping function $\phi : V' \mapsto V$, and the reverse relation holds. *Induced* means that if two vertices in $V(G')$ are adjacent in G' , then they are also adjacent in G . Formally, an induced subgraph subsequence $b' = (G'^{(1)}, G'^{(2)}, \dots, G'^{(m)})$ of a graph sequence $b = (G^{(1)}, G^{(2)}, \dots, G^{(n)})$, where $G'^{(i)} \subseteq G^{(i)}$ and $\phi(L') \rightarrow L$, is denoted as $G' \subseteq_i G$. The support of b' is denoted and defined as $sup_i(b') = |\{sid | (\langle sid, d \rangle \in GDB) \wedge (b' \subseteq_i d)\}|$. The anti-monotocity property **holds** for ISSs.

Given a minimum support threshold, Akihiro et al.¹⁰⁵ proposed to mine frequent ISSs from connected graph sequences, which are called **frequent relevant** induced subgraph subsequences (FRISS). By defining a mapping function, the main focus is to mine subgraph subsequences that share the same structure with graph sequences in the GDB and where vertex labels match.

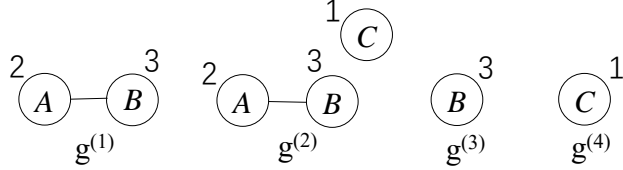


Figure 14: A frequent induced pattern

For example, a frequent ISS found in the GDB of Fig. 13 a) b), for a minimum support of 1, is shown in Fig. 14. It can be observed that the vertices that are adjacent in the frequent pattern are also adjacent in the GDB. Moreover, there is a correspondence relation between vertex labels in the graph sequence of the GDB and the subgraph sequence, where the mapping function is $\phi(1) = 2, \phi(2) = 3, \phi(3) = 1$.

The proposed FRISSMiner algorithm can be applied to directed or undirected graphs. FRISSMiner first **generates** a connected union graph for each graph sequence of the GDB. Then, all **frequent connected** induced subgraphs and their embeddings are found in each union graph by using a frequent subgraph mining algorithm. Then, **frequent connected** induced subgraphs are given as input to a modified version of the Prefixspan sequential pattern mining algorithm¹⁰⁷ to mine FRISS. Akihiro et al. have shown that FRISSMiner is useful to study e-mail interaction patterns.

Summary. Because dynamic graph sequence databases are often large and contain a large amount of information about vertices and edges from different graph sequences, it is difficult to mine knowledge (patterns) from such databases. The above algorithms mine frequent patterns that compactly represent graph sequences using transformation rules and by considering a mapping relation between vertex labels among different graph sequences. FRISS can reveal evolution patterns that are common to several sequences of a GDB, which is helpful to understand the distinct and representative features of a GDB. It is interesting to observe that the above algorithms break down **pattern** mining problems into two sub-problems that can be solved using traditional algorithms: (1) applying a traditional FSM algorithm on a union graph representation of graph sequences to find frequent subgraphs and their occurrences, and (2) then combining them using a modified sequential pattern mining algorithm.

It is challenging and useful to mine patterns in graph sequence databases and potential

applications are found in many domains. For example, besides the reported applications, algorithms could be used to analyze transportation data, where a graph sequence and its attributes may represent relationships between roads and indicate their states (e.g. light congestion or blocked) at different times. Furthermore, as there are much fewer algorithms for mining patterns in a graph sequence database than in a single dynamic graph, there are many research opportunities for developing novel algorithms.

MINING PATTERNS IN A DYNAMIC ATTRIBUTED GRAPH

Until now, this survey has mostly discussed studies for mining patterns in one or more dynamic labelled graphs, where a label may be associated to each vertex or edge. Although labelled graphs are used in many domains, it is desirable in some domains to consider more than one label per edge or vertex. For example, in social network analysis, a social graph may describe relationships (edges) between persons (vertices), where each person may be described using multiple attributes such as age, country and gender. Another example is research collaboration analysis where a research collaboration graph indicates the co-authorship relations (edges) between persons (vertices), and each person may be described using multiple attributes such as a publication count for different journals and conferences. Such data can be viewed as a dynamic graph where attribute values change over time, that is a *dynamic attributed graph*.

Formally, a *dynamic attributed graph* is a sequence of attributed graphs $\mathcal{G} = \langle G_1, G_2, \dots, G_{t_{max}} \rangle$ where $G_t = (\mathcal{V}_t, \mathcal{A}_t, E_t, \lambda_t)$, \mathcal{V}_t is a set of vertices, \mathcal{A}_t is a set of attributes, $E_t \subseteq \mathcal{V}_t \times \mathcal{V}_t$ is a set of edges, and $\lambda_t : \mathcal{V}_t \times \mathcal{A}_t \rightarrow \mathbb{R}$ is a function that associates a real value to each vertex-attribute pair, for the timestamp t . To be less influenced by noise when analyzing raw numerical values, a common practice is to convert numerical attribute values of a dynamic attributed graph into trends (attribute variations)^{89,102}. For example, Fig. 15 shows a dynamic attributed graph observed at six timestamps. In that figure, vertices are named 1, 2, 3, 4 and 5, attributes are named a_1 , a_2 and a_3 , and attribute values are numbers. Fig. 16 shows the result of converting that graph into a sequence of trend graphs. In particular, Fig. 16 (a) shows how attribute values have changed (trends) from the timestamp t_1 to t_2 either by increasing (+) or decreasing (-). Similarly, Fig. 16 (b), (c), (d) and (e)

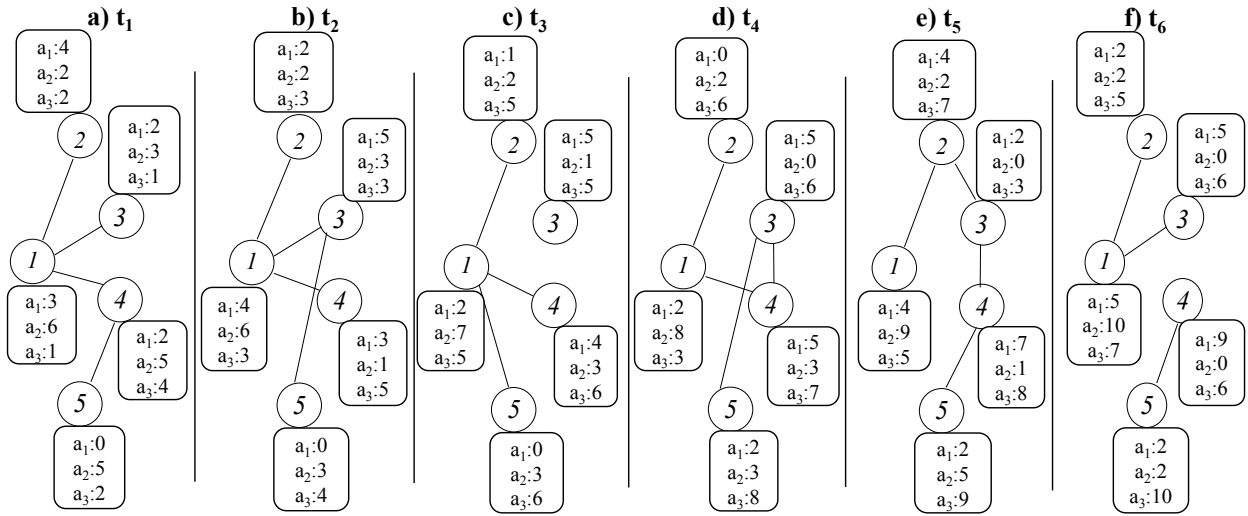


Figure 15: A dynamic attributed graph having six timestamps with raw numerical attribute values.

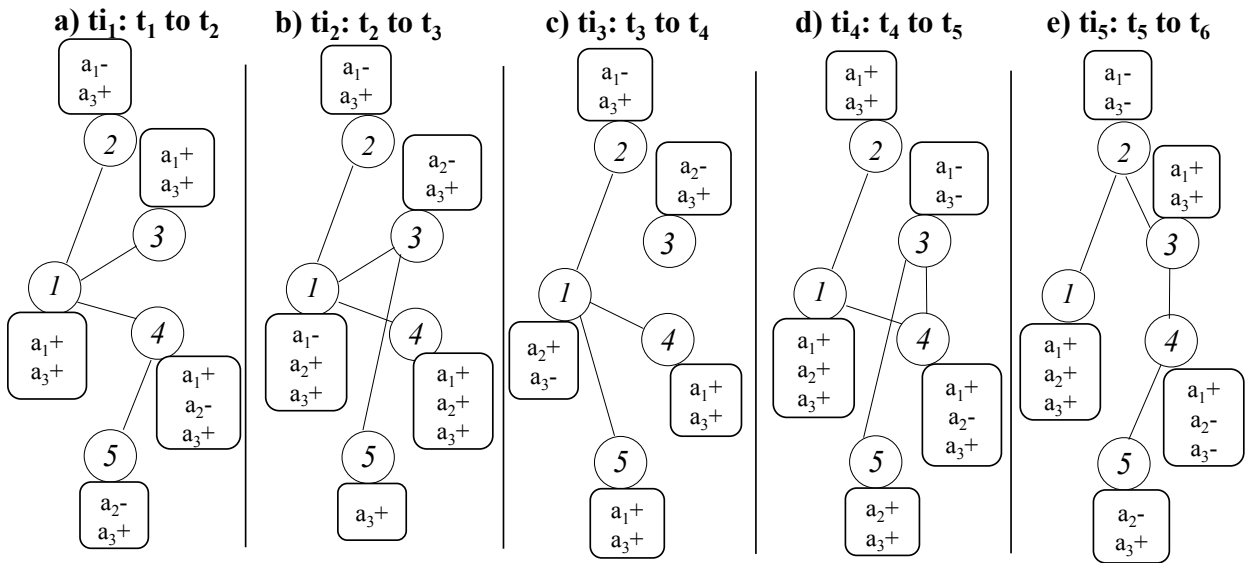


Figure 16: A sequence of trend graphs obtained by preprocessing the dynamic attributed graph of Fig. 15.

shows trends for timestamps t_2 to t_3 , t_3 to t_4 , t_4 to t_5 , and t_5 to t_6 . In the following these time intervals are called ti_1 , ti_2 , ti_3 , ti_4 and ti_5 , respectively.

A dynamic attributed graph is a generalization of the concept of dynamic graph, as defined in the previous subsection. Considering more than one attribute makes it possible to find more interesting patterns than when considering a single attribute. The reason is that patterns may **involve** one or more attributes, which provide information about how a graph evolves over time. Moreover, complex relationships between topological variations and attribute variations may be discovered. Using the concept of dynamic attributed graph also provides more flexibility to the user because rich information can be encoded using several attributes, such as local and global topological properties. The process of creating or selecting attributes for specific needs to then mine patterns is similar to that of feature engineering in machine learning. A user employing an algorithm for mining patterns in a dynamic attribute graph can change attributes and run the algorithm again, without having to change the algorithm. It is to be noted that to our best knowledge no studies on pattern mining in a dynamic attributed graph have considered assigning attributes to edges. The following paragraph describes the main pattern mining tasks for discovering patterns in dynamic attributed graphs.

Mining trend motifs. Trend motifs are a type of patterns found in dynamic attributed graphs having a single attribute⁸⁹. A trend motif is a connected subgraph whose vertices show the same trend (e.g. increase or decrease of an attribute value) during a time interval of two consecutive timestamps. Discovering trend motifs allows to find important changes in a dynamic system. A more detailed description of trend motifs was given in the subsection about mining patterns in a single dynamic graph.

Mining cohesive co-evolution patterns. The concept of cohesive co-evolution pattern was proposed by Desmier et al.⁹⁹. It is a set of vertices that are similar (based on a similarity measure) and display the same trends for some attribute(s) during a time interval. A co-evolution pattern may appear multiple times in a dynamic attributed graph, where each occurrence consists of time intervals formed by different pairs of timestamps that may or may not be consecutive. For example, Fig. 17 a) shows a cohesive co-evolution pattern found in the dynamic attributed graph of Fig. 16. This pattern indicates that during time intervals

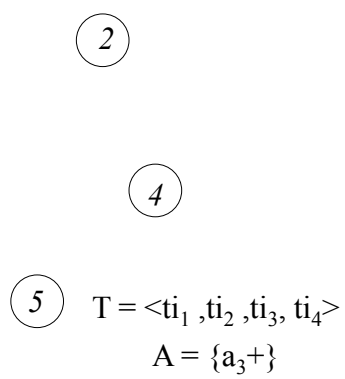
ti_1 , ti_2 , ti_3 and ti_4 , values of attributes a_3 of vertices 2, 4 and 5 have increased.

To filter uninteresting cohesive co-evolution patterns, several constraints have been considered such as a minimum similarity between vertices in a pattern and a volume constraint to ensure that patterns are less influenced by noise. Using these constraints can not only filter patterns but also improve the efficiency of pattern discovery. Because of the simple structure of co-evolution patterns, mining them does not require isomorphism checking or canonical labeling. Desmier et al. proposed an algorithm that directly enumerates cohesive co-evolution patterns by recursively appending vertices to patterns following a logical enumeration tree, and checking that constraints are satisfied. This approach ensures that no duplicates are generated.

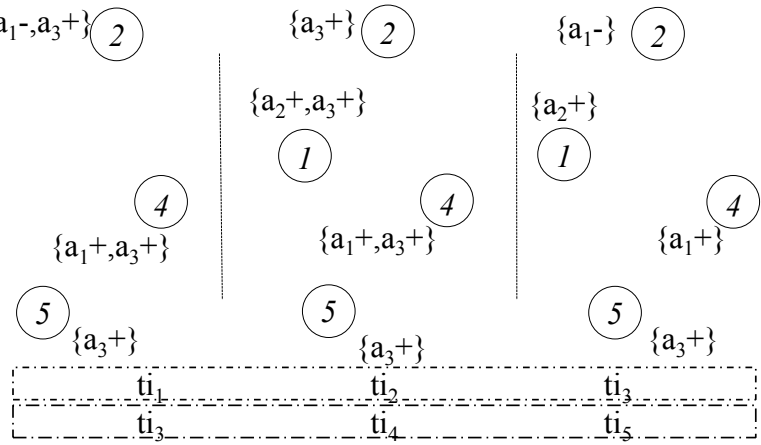
The algorithm was applied on a landslide dataset consisting of a series of satellite images where **vegetation declines** have been observed⁹⁹. These images were transformed in a dynamic attributed graph and patterns were extracted representing regions having experienced landslides. To make full use of the topological structure among vertices and provide more possibilities for the user, the authors also proposed other interestingness measures. **A user can utilize** these interestingness measures to more precisely express his preferences **to select** patterns, which can improve the efficiency of the algorithm (because constraints can help to reduce the search space).

Mining recurrent patterns. Cheng et al.⁹⁷ generalized the concept of cohesive co-evolution pattern by proposing a new type of patterns called *recurrent patterns*. These patterns capture how attribute values changed for a set of vertices over a sequence of time intervals. While a cohesive pattern is a subgraph, a recurrent pattern is a sequence of subgraphs (vertex sets), where each subgraph can be described using different attributes and trends. For example, Fig. 17 shows a recurrent pattern indicating changes appearing at three time intervals. This pattern has two instances, the first one is appearing at ti_1 , ti_2 and ti_3 of Fig. 16, and the second one at ti_3 , ti_4 and ti_5 . The algorithm proposed by Cheng et al., named RMiner, performs loops to consider the different combinations of time-intervals, to enumerate all patterns. This procedure does not require duplicate checking and canonical labeling. To select interesting recurrent patterns and improve the efficiency of pattern discovery, Cheng et al. also considered several constraints such that a pattern

a) A cohesive co-evolution pattern



b) A recurrent pattern



c) A significant trend sequence

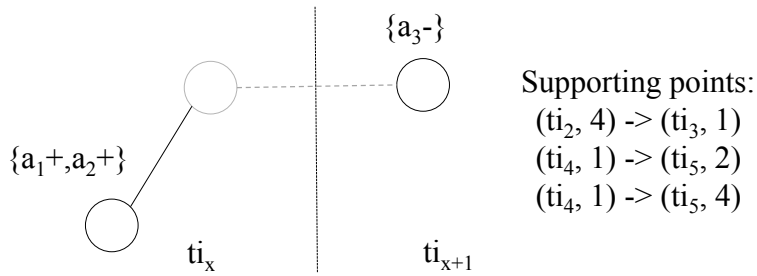


Figure 17: Examples of patterns found in the dynamic attributed graph of Fig. 16

must appear a minimum number of times, non-redundancy, and constraints on the volume of a vertex set and temporal continuity. Recurrent patterns allow to capture the frequent evolutions of trends for nodes in a dynamic attributed graph. RPFMiner was applied to extract patterns from a satellite image time series about aquaculture ponds to provide information to experts about how a set of connected ponds evolve together over time⁹⁸.

Mining triggering patterns of topological changes. Another type of patterns are *triggering patterns*¹⁰¹. A triggering pattern is a rule of the form $L \rightarrow R$ where L is a sequence of attribute variations followed by a single topological change, R . An example of triggering pattern is $\{a+, b+\}\{c-, d-, e-\} \rightarrow \{closeness-\}$, where a, b, c, d and e are node attributes or topological attributes, the symbol $+$ and $-$ indicate trends, and *closeness* is a topological attribute. This pattern indicates that trends on the left side of the rule triggered the topological change “closeness-” on the right side of rule. Furthermore, the growth rate of a pattern is measured to ensure that attribute variations triggered the topological change. The growth rate is defined in that paper as follows. Consider that vertices of the input graph are divided into two virtual databases. The first one consists of vertices whose attribute variation sequence contain R and the second one consists of the other vertices. The growth rate is the ratio of the frequency of L in the first database to its frequency in the second database. In the above example, a, b, c, d , and e can be topological attributes such as closeness, degree, number of cycles since such attributes can be encoded as node attributes.

To discover triggering patterns, it was proposed to convert a dynamic attributed graph into a sequence database and then to apply a frequent sequential pattern mining algorithm¹ to extract the desired patterns. In that sequence database representation, each sequence represents the set of attribute variations of a vertex over time. For example, a frequent sequence shared by several vertices may be $\langle \{closeness-, a+\}, \{numcycles+, b-\}, \{eigenvector+, c+\} \rangle$. The proposed approach was applied to analyze a trend graph of social bookmarking activity. Some discovered patterns revealed interesting insights such that if a user increases its number of bookmarks on some topics, it may trigger an increase of his number of followers.

Mining significant trend sequence. Although mining triggering patterns can reveal strong correlations between changes in a dynamic attributed graph, there are two main limitations. The first one is that to identify a triggering pattern, the correlation (growth rate)

is only calculated between the last attribute variation and the topological change. Hence, patterns may be found where attribute variations in L are weakly correlated with each other. Such patterns may be misleading for the user. The second limitation is that relationships between entities are only captured as topological properties, which cause considerable information loss and limit the information that can be expressed by the patterns.

To find more interesting strongly correlated patterns, Fournier-Viger et al.^{102,103} proposed a novel type of patterns called *significant trend sequences*. A significant trend sequence is an ordered list of attribute variations, where consecutive items are strongly correlated. For example, $\langle\{a_1+, a_2+\}, \{a_3-\}\rangle$ in Fig. 17 c) is a significant trend sequence that can be extracted from Fig. 16. To measure the correlation (significance), a novel significance measure is proposed, which is also based on the growth rate but calculated for all consecutive attribute variations. In the above example, $\{a_3-\}$ is strongly correlated with $\{a_1+, a_2+\}$ because $\{a_3-\}$ is not **globally frequent** but it very often follows $\{a_1+, a_2+\}$.

Mining trend sequences is not an easy problem as increasing the number of trends can exponentially increase the size of the search space. To efficiently mine the proposed patterns, two projection based algorithms named $TSeqMiner_{dfs}$ and $TSeqMiner_{bfs}$ were designed. They decompose the pattern mining task into two sub-tasks: (1) finding sets of frequent attribute variations for a time intervals using a modified itemset mining algorithm, (2) and extending patterns by combining sets from different time intervals using either a depth-first search or a breadth-first search. The two algorithms rely on several **search space** pruning strategies such as upper bound filtering to avoid exploring the whole search space **while ensuring that all patterns are found**. As most algorithms reviewed in this subsection, no duplicates are generated, and techniques commonly used in FSM such as canonical labeling and isomorphism checking are not required.

The two algorithms were first applied to the DBLP dataset, which is a dynamic attributed graph about co-authorship relationships between researchers in different conferences and journals over several years. Some interesting trend sequences were found such as $\langle\{VLDB+\}, \{ICDE+, VLDB =\}\rangle$, which indicates that an author publishing an increasing number of papers in VLDB is likely to publish more ICDE papers while having a stable number of publications in VLDB at the next timestamp (in terms of years). The two al-

gorithms were also applied on US flight data collected during several years. An interesting pattern found is $\{(NbCancellation--, NbDivertedFlights--, NbDelayedDepartures-), (NbDepartures-, NbCancellations-, NbDivertedFlights-, DelayedDepartures-, NbDelayedArrivals+)\}$, which indicates that after an airport recovered from a hurricane’s **damage**, the number of cancellations, diverted flights and delayed departures decreased, which then influenced airports that were not damaged by the hurricane (but are connected). In that pattern, the symbols $-$ and $+$ indicate a small increase or decrease, while $--$ and $++$ indicate a large increase or decrease, **respectively**.

Summary. This subsection has reviewed several tasks of mining patterns in a dynamic attributed graph. Different patterns are discovered using different measures, which can be suitable for different needs. It is interesting to observe that most algorithms in this subsection do not rely on traditional FSM techniques, except for the algorithm of Jin et al.⁸⁹. This is because the pattern types presented in this section are based on simple forms of subgraphs (e.g. a vertex set), though they include the time dimension and multiple attributes. Because of this, some algorithms instead rely on modified itemset mining or sequential pattern mining algorithms^{101,102}, or use custom pattern enumeration procedures^{97,99}. It is also interesting that pattern mining problems in a dynamic attributed graph can sometimes be decomposed into two sub-tasks corresponding to the **attribute dimension and the time dimension**¹⁰².

EXTENSIONS

The previous sections have described the main tasks for discovering patterns in dynamic graphs. There exists various extensions of these tasks that have not been discussed so far. This section gives a brief overview.

Several studies have been done on mining patterns in a data stream of graphs¹¹³, also sometimes called streaming graphs or evolving graphs⁷⁶. A stream is an infinite sequence of graphs arriving at high speed. To mine patterns in a stream, adapted algorithms need to be designed because unlike a database, a stream can only be read once and the full stream cannot be kept in memory. Moreover, data distribution may change over time in a stream. For instance, Nishioka et al.¹¹³ proposed two algorithms for mining frequent subgraphs in dense graph streams with limited memory using a disk-based structure. In

another study, Ray et al.¹¹⁴ proposed an algorithm to mine frequent subgraphs in a large attributed streaming graph. The algorithm is an approximate algorithm, which assumes that updates arrive as batches, and where each update is composed of adding nodes and edges. The algorithm timely reports the likely frequent subgraphs, which can help to monitor the network. The study was applied to social network and movie related data, among others.

Some papers have extended graph pattern mining tasks to consider uncertainty of the data. For example, Leung and Cuzzocrea¹¹⁵ proposed algorithms to mine frequent subgraphs in an uncertain data stream where edges are annotated with existential probabilities. Some papers have also considered finding rare patterns instead of frequent ones¹¹⁸ and weighted patterns¹¹⁹.

Other papers have also been proposed to solve pattern mining problems of specific applications. For example, Javel et al.¹¹⁶ designed a method for detecting sequences of changes in ontologies to reveal how it is edited over time. In that work, an ontology is represented as a dynamic attributed graph.

Another extension is about privacy-preserving data mining, where the goal is to hide sensitive patterns that may reveal important information. For example, Cheng et al.¹¹⁷ designed a two-phase algorithm for hiding sensitive subgraphs in the context of frequent subgraph mining from a graph database. Although this work is not on dynamic graphs, it is relevant as some dynamic subgraph mining algorithms rely on traditional subgraph mining algorithms.

RESEARCH OPPORTUNITIES

Mining patterns in dynamic graphs is an active research area. Although several papers have been published in this field, there are numerous research opportunities. Some of the key research opportunities are:

- **Design more efficient algorithms.** Since pattern mining is generally quite computationally expensive, it is important to design more efficient algorithms in terms of runtime and memory. This can be done by developing novel algorithms, search space pruning strategies and data structures. Moreover, GPU, multi-thread, and parallel

algorithms can be designed to scale to very large datasets. Moreover, additional constraints may be integrated in algorithms to select more interesting patterns and reduce the search space.

- **Discover patterns in more complex data.** A trend in recent years has been to consider more complex **data types** such as attributed graphs^{97,99,100,102}, streams^{113,114}, and graphs with uncertainty¹¹⁵. The reason is that complex data are found in many applications. Developing models to handle complex data is thus important to address real-life problems. Examples of **novel** problems that could be studied are to mine patterns in attributed graphs with attributes not only on vertices but also on edges and to mine patterns in a database of attributed graphs. Another interesting possibility is to consider other graph representations for representing dynamic graphs besides the snapshot-based model, which is used in all reviewed papers on pattern mining in dynamic graphs. Wehmuth et al.¹²² **provide** an interesting review of alternative models that could be used. For instance, one could consider a model with continuous time intervals.
- **Discover more complex pattern types.** Another important research direction is to develop algorithms to identify more complex patterns that provide more useful information to users. This can be done by extending pattern definitions or considering additional constraints or interestingness measures. A source of inspiration can be other pattern mining tasks such as itemset mining¹⁰ and sequential pattern mining¹, for which many extensions have been developed. **For example, various types of patterns involving time have been proposed such as peaks¹²³, trends¹²⁵ and patterns having a stable behavior over time¹²⁴.**
- **Novel applications.** Algorithms for mining patterns in dynamic graphs can be applied to novel applications where data can be represented as dynamic graphs. This is especially interesting for emerging applications such as the Internet of Things⁴⁶, edge computing⁴⁴, and Vehicular Ad-hoc NETWORKS (VANET)⁴⁵. This may not only provide solutions to applied problems but the applications may raise new challenges that may inspire further research.

CONCLUSION

Dynamic graphs are a type of data commonly found in numerous fields. Discovering patterns in such graphs can provide insights on data. This survey has provided an overview of algorithms for discovering patterns in dynamic graphs, including those for mining patterns in a dynamic graph, graph sequence database, and dynamic attributed graphs. Moreover, other extensions have been discussed such as graph mining in streams and uncertain data. Finally, research opportunities have been discussed.

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