# CHAPTER 11 Graph Pattern Mining

Graph data is becoming increasingly more ubiquitous in today's networked world. Examples include social networks as well as cell phone networks and blogs. The Internet is another example of graph data, as is the hyperlinked structure of the World Wide Web (WWW). Bioinformatics, especially systems biology, deals with understanding interaction networks between various types of biomolecules, such as protein–protein interactions, metabolic networks, gene networks, and so on. Another prominent source of graph data is the Semantic Web, and linked open data, with graphs represented using the Resource Description Framework (RDF) data model.

The goal of graph mining is to extract interesting subgraphs from a single large graph (e.g., a social network), or from a database of many graphs. In different applications we may be interested in different kinds of subgraph patterns, such as subtrees, complete graphs or cliques, bipartite cliques, dense subgraphs, and so on. These may represent, for example, communities in a social network, hub and authority pages on the WWW, cluster of proteins involved in similar biochemical functions, and so on. In this chapter we outline methods to mine all the frequent subgraphs that appear in a database of graphs.

#### 11.1 ISOMORPHISM AND SUPPORT

A graph is a pair G = (V, E) where V is a set of vertices, and  $E \subseteq V \times V$  is a set of edges. We assume that edges are unordered, so that the graph is undirected. If (u, v) is an edge, we say that u and v are *adjacent* and that v is a *neighbor* of u, and vice versa. The set of all neighbors of u in G is given as  $N(u) = \{v \in V \mid (u, v) \in E\}$ . A *labeled graph* has labels associated with its vertices as well as edges. We use L(u) to denote the label of the vertex u, and L(u, v) to denote the label of the edge (u, v), with the set of vertex labels denoted as  $\Sigma_V$  and the set of edge labels as  $\Sigma_E$ . Given an edge  $(u, v) \in G$ , the tuple  $\langle u, v, L(u), L(v), L(u, v) \rangle$  that augments the edge with the node and edge labels is called an *extended edge*.

**Example 11.1.** Figure 11.1a shows an example of an unlabeled graph, whereas Figure 11.1b shows the same graph, with labels on the vertices, taken from the vertex

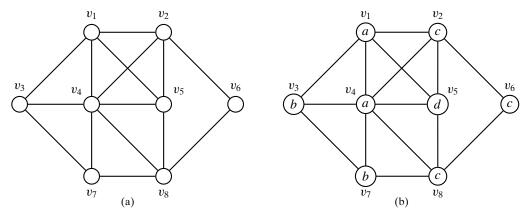


Figure 11.1. An unlabeled (a) and labeled (b) graph with eight vertices.

label set  $\Sigma_V = \{a, b, c, d\}$ . In this example, edges are all assumed to be unlabeled, and are therefore edge labels are not shown. Considering Figure 11.1b, the label of vertex  $v_4$  is  $L(v_4) = a$ , and its neighbors are  $N(v_4) = \{v_1, v_2, v_3, v_5, v_7, v_8\}$ . The edge  $(v_4, v_1)$  leads to the extended edge  $\langle v_4, v_1, a, a \rangle$ , where we omit the edge label  $L(v_4, v_1)$  because it is empty.

## **Subgraphs**

A graph G' = (V', E') is said to be a *subgraph* of G if  $V' \subseteq V$  and  $E' \subseteq E$ . Note that this definition allows for disconnected subgraphs. However, typically data mining applications call for *connected subgraphs*, defined as a subgraph G' such that  $V' \subseteq V$ ,  $E' \subseteq E$ , and for any two nodes  $u, v \in V'$ , there exists a *path* from u to v in G'.

**Example 11.2.** The graph defined by the bold edges in Figure 11.2a is a subgraph of the larger graph; it has vertex set  $V' = \{v_1, v_2, v_4, v_5, v_6, v_8\}$ . However, it is a disconnected subgraph. Figure 11.2b shows an example of a connected subgraph on the same vertex set V'.

#### **Graph and Subgraph Isomorphism**

A graph G' = (V', E') is said to be *isomorphic* to another graph G = (V, E) if there exists a bijective function  $\phi : V' \to V$ , i.e., both injective (into) and surjective (onto), such that

- 1.  $(u, v) \in E' \iff (\phi(u), \phi(v)) \in E$
- 2.  $\forall u \in V', L(u) = L(\phi(u))$
- 3.  $\forall (u, v) \in E', L(u, v) = L(\phi(u), \phi(v))$

In other words, the *isomorphism*  $\phi$  preserves the edge adjacencies as well as the vertex and edge labels. Put differently, the extended tuple  $\langle u, v, L(u), L(v), L(u, v) \rangle \in G'$  if and only if  $\langle \phi(u), \phi(v), L(\phi(u)), L(\phi(v)), L(\phi(u), \phi(v)) \rangle \in G$ .

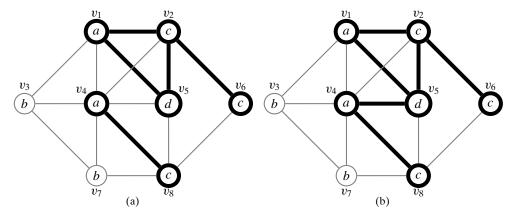


Figure 11.2. A subgraph (a) and connected subgraph (b).

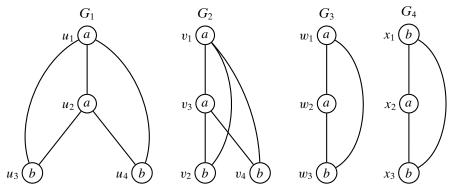


Figure 11.3. Graph and subgraph isomorphism.

If the function  $\phi$  is only injective but not surjective, we say that the mapping  $\phi$  is a *subgraph isomorphism* from G' to G. In this case, we say that G' is isomorphic to a subgraph of G, that is, G' is *subgraph isomorphic* to G, denoted  $G' \subseteq G$ ; we also say that G contains G'.

**Example 11.3.** In Figure 11.3,  $G_1 = (V_1, E_1)$  and  $G_2 = (V_2, E_2)$  are isomorphic graphs. There are several possible isomorphisms between  $G_1$  and  $G_2$ . An example of an isomorphism  $\phi: V_2 \to V_1$  is

$$\phi(v_1) = u_1$$
  $\phi(v_2) = u_3$   $\phi(v_3) = u_2$   $\phi(v_4) = u_4$ 

The inverse mapping  $\phi^{-1}$  specifies the isomorphism from  $G_1$  to  $G_2$ . For example,  $\phi^{-1}(u_1) = v_1$ ,  $\phi^{-1}(u_2) = v_3$ , and so on. The set of all possible isomorphisms from  $G_2$  to  $G_1$  are as follows:

The graph  $G_3$  is subgraph isomorphic to both  $G_1$  and  $G_2$ . The set of all possible subgraph isomorphisms from  $G_3$  to  $G_1$  are as follows:

$$\begin{array}{c|ccccc} & w_1 & w_2 & w_3 \\ \hline \phi_1 & u_1 & u_2 & u_3 \\ \phi_2 & u_1 & u_2 & u_4 \\ \phi_3 & u_2 & u_1 & u_3 \\ \phi_4 & u_2 & u_1 & u_4 \\ \end{array}$$

The graph  $G_4$  is not subgraph isomorphic to either  $G_1$  or  $G_2$ , and it is also not isomorphic to  $G_3$  because the extended edge  $\langle x_1, x_3, b, b \rangle$  has no possible mappings in  $G_1, G_2$  or  $G_3$ .

## **Subgraph Support**

Given a database of graphs,  $\mathbf{D} = \{G_1, G_2, \dots, G_n\}$ , and given some graph G, the support of G in  $\mathbf{D}$  is defined as follows:

$$sup(G) = \left| \left\{ G_i \in \mathbf{D} \mid G \subseteq G_i \right\} \right|$$

The support is simply the number of graphs in the database that contain G. Given a *minsup* threshold, the goal of graph mining is to mine all frequent connected subgraphs with  $sup(G) \ge minsup$ .

To mine all the frequent subgraphs, one has to search over the space of all possible graph patterns, which is exponential in size. If we consider subgraphs with m vertices, then there are  $\binom{m}{2} = O(m^2)$  possible edges. The number of possible subgraphs with m nodes is then  $O(2^{m^2})$  because we may decide either to include or exclude each of the edges. Many of these subgraphs will not be connected, but  $O(2^{m^2})$  is a convenient upper bound. When we add labels to the vertices and edges, the number of labeled graphs will be even more. Assume that  $|\Sigma_V| = |\Sigma_E| = s$ , then there are  $s^m$  possible ways to label the vertices and there are  $s^{m^2}$  ways to label the edges. Thus, the number of possible labeled subgraphs with m vertices is  $2^{m^2}s^ms^{m^2} = O((2s)^{m^2})$ . This is the worst case bound, as many of these subgraphs will be isomorphic to each other, with the number of distinct subgraphs being much less. Nevertheless, the search space is still enormous because we typically have to search for all subgraphs ranging from a single vertex to some maximum number of vertices given by the largest frequent subgraph.

There are two main challenges in frequent subgraph mining. The first is to systematically generate candidate subgraphs. We use *edge-growth* as the basic mechanism for extending the candidates. The mining process proceeds in a breadth-first (level-wise) or a depth-first manner, starting with an empty subgraph (i.e., with no edge), and adding a new edge each time. Such an edge may either connect two existing vertices in the graph or it may introduce a new vertex as one end of a new edge. The key is to perform nonredundant subgraph enumeration, such that we do not generate the same graph candidate more than once. This means that we have to perform graph isomorphism checking to make sure that duplicate graphs are removed. The second challenge is to count the support of a graph in the database. This involves subgraph isomorphism checking, as we have to find the set of graphs that contain a given candidate.

An effective strategy to enumerate subgraph patterns is the so-called *rightmost path extension*. Given a graph G, we perform a depth-first search (DFS) over its vertices, and create a DFS spanning tree, that is, one that covers or spans all the vertices. Edges that are included in the DFS tree are called *forward* edges, and all other edges are called *backward* edges. Backward edges create cycles in the graph. Once we have a DFS tree, define the *rightmost* path as the path from the root to the rightmost leaf, that is, to the leaf with the highest index in the DFS order.

**Example 11.4.** Consider the graph shown in Figure 11.4a. One of the possible DFS spanning trees is shown in Figure 11.4b (illustrated via bold edges), obtained by starting at  $v_1$  and then choosing the vertex with the smallest index at each step. Figure 11.5 shows the same graph (ignoring the dashed edges), rearranged to emphasize the DFS tree structure. For instance, the edges  $(v_1, v_2)$  and  $(v_2, v_3)$  are examples of forward edges, whereas  $(v_3, v_1)$ ,  $(v_4, v_1)$ , and  $(v_6, v_1)$  are all backward edges. The bold edges  $(v_1, v_5)$ ,  $(v_5, v_7)$  and  $(v_7, v_8)$  comprise the rightmost path.

For generating new candidates from a given graph G, we extend it by adding a new edge to vertices only on the rightmost path. We can either extend G by adding backward edges from the *rightmost vertex* to some other vertex on the rightmost path (disallowing self-loops or multi-edges), or we can extend G by adding forward edges from any of the vertices on the rightmost path. A backward extension does not add a new vertex, whereas a forward extension adds a new vertex.

For systematic candidate generation we impose a total order on the extensions, as follows: First, we try all backward extensions from the rightmost vertex, and then we try forward extensions from vertices on the rightmost path. Among the backward edge extensions, if  $u_r$  is the rightmost vertex, the extension  $(u_r, v_i)$  is tried before  $(u_r, v_j)$  if i < j. In other words, backward extensions closer to the root are considered before those farther away from the root along the rightmost path. Among the forward edge extensions, if  $v_x$  is the new vertex to be added, the extension  $(v_i, v_x)$  is tried before

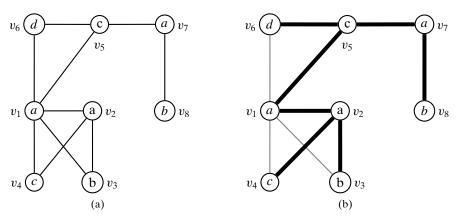


Figure 11.4. A graph (a) and a possible depth-first spanning tree (b).

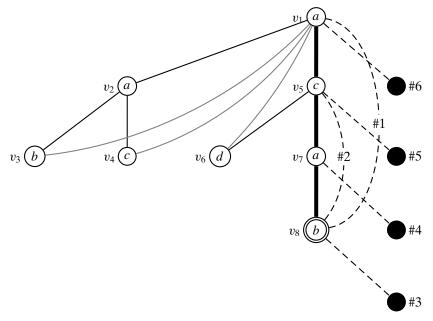


Figure 11.5. Rightmost path extensions. The bold path is the rightmost path in the DFS tree. The *rightmost* vertex is  $v_8$ , shown double circled. Solid black lines (thin and bold) indicate the *forward* edges, which are part of the DFS tree. The *backward* edges, which by definition are not part of the DFS tree, are shown in gray. The set of possible extensions on the rightmost path are shown with dashed lines. The precedence ordering of the extensions is also shown.

 $(v_j, v_x)$  if i > j. In other words, the vertices farther from the root (those at greater depth) are extended before those closer to the root. Also note that the new vertex will be numbered x = r + 1, as it will become the new rightmost vertex after the extension.

**Example 11.5.** Consider the order of extensions shown in Figure 11.5. Node  $v_8$  is the rightmost vertex; thus we try backward extensions only from  $v_8$ . The first extension, denoted #1 in Figure 11.5, is the backward edge  $(v_8, v_1)$  connecting  $v_8$  to the root, and the next extension is  $(v_8, v_5)$ , denoted #2, which is also backward. No other backward extensions are possible without introducing multiple edges between the same pair of vertices. The forward extensions are tried in reverse order, starting from the rightmost vertex  $v_8$  (extension denoted as #3) and ending at the root (extension denoted as #6). Thus, the forward extension  $(v_8, v_x)$ , denoted #3, comes before the forward extension  $(v_7, v_x)$ , denoted #4, and so on.

#### 11.2.1 Canonical Code

When generating candidates using rightmost path extensions, it is possible that duplicate, that is, isomorphic, graphs are generated via different extensions. Among the isomorphic candidates, we need to keep only one for further extension, whereas the others can be pruned to avoid redundant computation. The main idea is that if we can somehow sort or rank the isomorphic graphs, we can pick the *canonical representative*, say the one with the least rank, and extend only that graph.

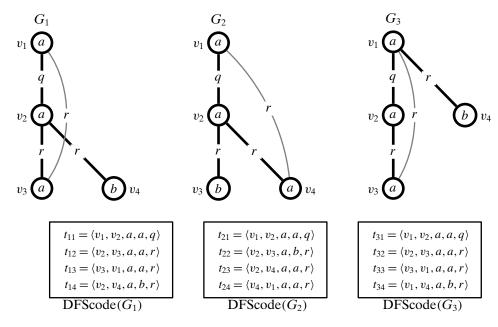


Figure 11.6. Canonical DFS code.  $G_1$  is canonical, whereas  $G_2$  and  $G_3$  are noncanonical. Vertex label set  $\Sigma_V = \{a, b\}$ , and edge label set  $\Sigma_E = \{q, r\}$ . The vertices are numbered in DFS order.

Let G be a graph and let  $T_G$  be a DFS spanning tree for G. The DFS tree  $T_G$  defines an ordering of both the nodes and edges in G. The DFS node ordering is obtained by numbering the nodes consecutively in the order they are visited in the DFS walk. We assume henceforth that for a pattern graph G the nodes are numbered according to their position in the DFS ordering, so that i < j implies that  $v_i$  comes before  $v_j$  in the DFS walk. The DFS edge ordering is obtained by following the edges between consecutive nodes in DFS order, with the condition that all the backward edges incident with vertex  $v_i$  are listed before any of the forward edges incident with it. The DFS code for a graph G, for a given DFS tree  $T_G$ , denoted DFScode(G), is defined as the sequence of extended edge tuples of the form  $\langle v_i, v_j, L(v_i), L(v_j), L(v_i, v_j) \rangle$  listed in the DFS edge order.

**Example 11.6.** Figure 11.6 shows the DFS codes for three graphs, which are all isomorphic to each other. The graphs have node and edge labels drawn from the label sets  $\Sigma_V = \{a,b\}$  and  $\Sigma_E = \{q,r\}$ . The edge labels are shown centered on the edges. The bold edges comprise the DFS tree for each graph. For  $G_1$ , the DFS node ordering is  $v_1, v_2, v_3, v_4$ , whereas the DFS edge ordering is  $(v_1, v_2), (v_2, v_3), (v_3, v_1),$  and  $(v_2, v_4)$ . Based on the DFS edge ordering, the first tuple in the DFS code for  $G_1$  is therefore  $\langle v_1, v_2, a, a, q \rangle$ . The next tuple is  $\langle v_2, v_3, a, a, r \rangle$  and so on. The DFS code for each graph is shown in the corresponding box below the graph.

#### Canonical DFS Code

A subgraph is *canonical* if it has the smallest DFS code among all possible isomorphic graphs, with the ordering between codes defined as follows. Let  $t_1$  and  $t_2$  be any two

DFS code tuples:

$$t_1 = \langle v_i, v_j, L(v_i), L(v_j), L(v_i, v_j) \rangle$$
  

$$t_2 = \langle v_x, v_y, L(v_x), L(v_y), L(v_x, v_y) \rangle$$

We say that  $t_1$  is smaller than  $t_2$ , written  $t_1 < t_2$ , iff

i) 
$$(v_i, v_j) <_e (v_x, v_y)$$
, or  
ii)  $(v_i, v_j) = (v_x, v_y)$  and  $(11.1)$   
 $\langle L(v_i), L(v_j), L(v_i, v_j) \rangle <_l \langle L(v_x), L(v_y), L(v_x, v_y) \rangle$ 

where  $<_e$  is an ordering on the edges and  $<_l$  is an ordering on the vertex and edge labels. The *label order*  $<_l$  is the standard lexicographic order on the vertex and edge labels. The *edge order*  $<_e$  is derived from the rules for rightmost path extension, namely that all of a node's backward extensions must be considered before any forward edge from that node, and deep DFS trees are preferred over bushy DFS trees. Formally, Let  $e_{ij} = (v_i, v_j)$  and  $e_{xy} = (v_x, v_y)$  be any two edges. We say that  $e_{ij} <_e e_{xy}$  iff

- Condition (1) If  $e_{ij}$  and  $e_{xy}$  are both forward edges, then (a) j < y, or (b) j = y and i > x. That is, (a) a forward extension to a node earlier in the DFS node order is smaller, or (b) if both the forward edges point to a node with the same DFS node order, then the forward extension from a node deeper in the tree is smaller.
- Condition (2) If  $e_{ij}$  and  $e_{xy}$  are both backward edges, then (a) i < x, or (b) i = x and j < y. That is, (a) a backward edge from a node earlier in the DFS node order is smaller, or (b) if both the backward edges originate from a node with the same DFS node order, then the backward edge to a node earlier in DFS node order (i.e., closer to the root along the rightmost path) is smaller.
- Condition (3) If  $e_{ij}$  is a forward and  $e_{xy}$  is a backward edge, then  $j \le x$ . That is, a forward edge to a node earlier in the DFS node order is smaller than a backward edge from that node or any node that comes after it in DFS node order.
- Condition (4) If  $e_{ij}$  is a backward and  $e_{xy}$  is a forward edge, then i < y. That is, a backward edge from a node earlier in DFS node order is smaller than a forward edge to any later node.

Given any two DFS codes, we can compare them tuple by tuple to check which is smaller. In particular, the *canonical DFS code* for a graph G is defined as follows:

$$C = \min_{G'} \left\{ \mathsf{DFScode}(G') \mid G' \text{ is isomorphic to } G \right\}$$

Given a candidate subgraph G, we can first determine whether its DFS code is canonical or not. Only canonical graphs need to be retained for extension, whereas noncanonical candidates can be removed from further consideration.

**Example 11.7.** Consider the DFS codes for the three graphs shown in Figure 11.6. Comparing  $G_1$  and  $G_2$ , we find that  $t_{11} = t_{21}$ , but  $t_{12} < t_{22}$  because  $\langle a, a, r \rangle <_l \langle a, b, r \rangle$ . Comparing the codes for  $G_1$  and  $G_3$ , we find that the first three tuples are equal for both the graphs, but  $t_{14} < t_{34}$  because

$$(v_i, v_i) = (v_2, v_4) <_e (v_1, v_4) = (v_x, v_y)$$

due to condition (1) above. That is, both are forward edges, and we have  $v_j = v_4 = v_y$  with  $v_i = v_2 > v_1 = v_x$ . In fact, it can be shown that the code for  $G_1$  is the canonical DFS code for all graphs isomorphic to  $G_1$ . Thus,  $G_1$  is the canonical candidate.

#### 11.3 THE GSPAN ALGORITHM

We describe the gSpan algorithm to mine all frequent subgraphs from a database of graphs. Given a database  $\mathbf{D} = \{G_1, G_2, ..., G_n\}$  comprising n graphs, and given a minimum support threshold *minsup*, the goal is to enumerate all (connected) subgraphs G that are frequent, that is,  $sup(G) \ge minsup$ . In gSpan, each graph is represented by its canonical DFS code, so that the task of enumerating frequent subgraphs is equivalent to the task of generating all canonical DFS codes for frequent subgraphs. Algorithm 11.1 shows the pseudo-code for gSpan.

gSpan enumerates patterns in a depth-first manner, starting with the empty code. Given a canonical and frequent code C, gSpan first determines the set of possible edge extensions along the rightmost path (line 1). The function RIGHTMOSTPATH-EXTENSIONS returns the set of edge extensions along with their support values,  $\mathcal{E}$ . Each extended edge t in  $\mathcal{E}$  leads to a new candidate DFS code  $C' = C \cup \{t\}$ , with support sup(C) = sup(t) (lines 3–4). For each new candidate code, gSpan checks whether it is frequent and canonical, and if so gSpan recursively extends C' (lines 5–6). The algorithm stops when there are no more frequent and canonical extensions possible.

## ALGORITHM 11.1. Algorithm GSPAN

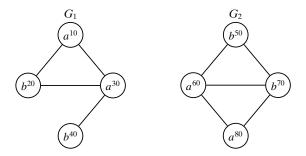


Figure 11.7. Example graph database.

**Example 11.8.** Consider the example graph database comprising  $G_1$  and  $G_2$  shown in Figure 11.7. Let minsup = 2, that is, assume that we are interested in mining subgraphs that appear in both the graphs in the database. For each graph the node labels and node numbers are both shown, for example, the node  $a^{10}$  in  $G_1$  means that node 10 has label a.

Figure 11.8 shows the candidate patterns enumerated by gSpan. For each candidate the nodes are numbered in the DFS tree order. The solid boxes show frequent subgraphs, whereas the dotted boxes show the infrequent ones. The dashed boxes represent noncanonical codes. Subgraphs that do not occur even once are not shown. The figure also shows the DFS codes and their corresponding graphs.

The mining process begins with the empty DFS code  $C_0$  corresponding to the empty subgraph. The set of possible 1-edge extensions comprises the new set of candidates. Among these,  $C_3$  is pruned because it is not canonical (it is isomorphic to  $C_2$ ), whereas  $C_4$  is pruned because it is not frequent. The remaining two candidates,  $C_1$  and  $C_2$ , are both frequent and canonical, and are thus considered for further extension. The depth-first search considers  $C_1$  before  $C_2$ , with the rightmost path extensions of  $C_1$  being  $C_5$  and  $C_6$ . However,  $C_6$  is not canonical; it is isomorphic to  $C_5$ , which has the canonical DFS code. Further extensions of  $C_5$  are processed recursively. Once the recursion from  $C_1$  completes, gSpan moves on to  $C_2$ , which will be recursively extended via rightmost edge extensions as illustrated by the subtree under  $C_2$ . After processing  $C_2$ , gSpan terminates because no other frequent and canonical extensions are found. In this example,  $C_{12}$  is a maximal frequent subgraph, that is, no supergraph of  $C_{12}$  is frequent.

This example also shows the importance of duplicate elimination via canonical checking. The groups of isomorphic subgraphs encountered during the execution of gSpan are as follows:  $\{C_2, C_3\}$ ,  $\{C_5, C_6, C_{17}\}$ ,  $\{C_7, C_{19}\}$ ,  $\{C_9, C_{25}\}$ ,  $\{C_{20}, C_{21}, C_{22}, C_{24}\}$ , and  $\{C_{12}, C_{13}, C_{14}\}$ . Within each group the first graph is canonical and thus the remaining codes are pruned.

For a complete description of gSpan we have to specify the algorithm for enumerating the rightmost path extensions and their support, so that infrequent patterns can be eliminated, and the procedure for checking whether a given DFS code is canonical, so that duplicate patterns can be pruned. These are detailed next.

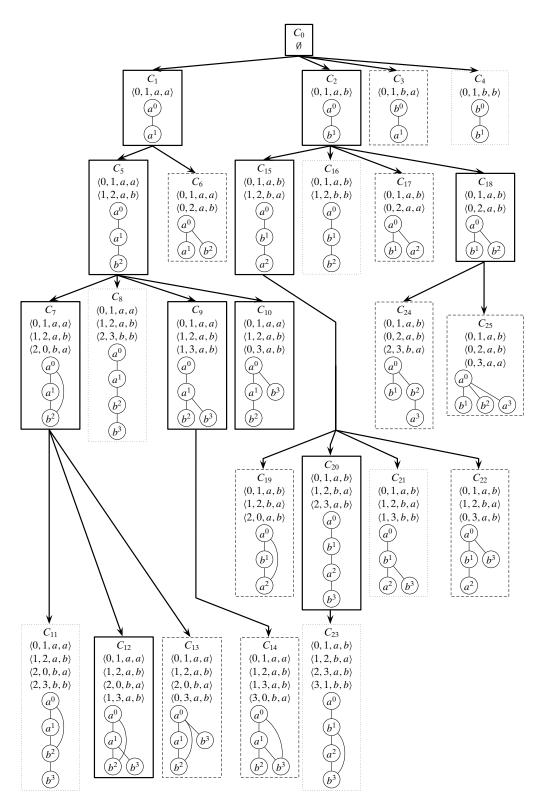


Figure 11.8. Frequent graph mining: minsup = 2. Solid boxes indicate the frequent subgraphs, dotted the infrequent, and dashed the noncanonical subgraphs.

## 11.3.1 Extension and Support Computation

The support computation task is to find the number of graphs in the database **D** that contain a candidate subgraph, which is very expensive because it involves subgraph isomorphism checks. gSpan combines the tasks of enumerating candidate extensions and support computation.

Assume that  $\mathbf{D} = \{G_1, G_2, \dots, G_n\}$  comprises n graphs. Let  $C = \{t_1, t_2, \dots, t_k\}$  denote a frequent canonical DFS code comprising k edges, and let G(C) denote the graph corresponding to code C. The task is to compute the set of possible rightmost path extensions from C, along with their support values, which is accomplished via the pseudo-code in Algorithm 11.2.

Given code C, gSpan first records the nodes on the rightmost path (R), and the rightmost child  $(u_r)$ . Next, gSpan considers each graph  $G_i \in \mathbf{D}$ . If  $C = \emptyset$ , then each distinct label tuple of the form  $\langle L(x), L(y), L(x,y) \rangle$  for adjacent nodes x and y in  $G_i$  contributes a forward extension  $\langle 0, 1, L(x), L(y), L(x,y) \rangle$  (lines 6-8). On the other hand, if C is not empty, then gSpan enumerates all possible subgraph isomorphisms  $\Phi_i$  between the code C and graph  $G_i$  via the function SUBGRAPHISOMORPHISMS (line 10). Given subgraph isomorphism  $\phi \in \Phi_i$ , gSpan finds all possible forward and backward edge extensions, and stores them in the extension set  $\mathcal{E}$ .

Backward extensions (lines 12–15) are allowed only from the rightmost child  $u_r$  in C to some other node on the rightmost path R. The method considers each neighbor x of  $\phi(u_r)$  in  $G_i$  and checks whether it is a mapping for some vertex  $v = \phi^{-1}(x)$  along the rightmost path R in C. If the edge  $(u_r, v)$  does not already exist in C, it is a new extension, and the extended tuple  $b = \langle u_r, v, L(u_r), L(v), L(u_r, v) \rangle$  is added to the set of extensions  $\mathcal{E}$ , along with the graph id i that contributed to that extension.

Forward extensions (lines 16–19) are allowed only from nodes on the rightmost path R to new nodes. For each node u in R, the algorithm finds a neighbor x in  $G_i$  that is not in a mapping from some node in C. For each such node x, the forward extension  $f = \langle u, u_r + 1, L(\phi(u)), L(x), L(\phi(u), x) \rangle$  is added to  $\mathcal{E}$ , along with the graph id i. Because a forward extension adds a new vertex to the graph G(C), the id of the new node in C must be  $u_r + 1$ , that is, one more than the highest numbered node in C, which by definition is the rightmost child  $u_r$ .

Once all the backward and forward extensions have been cataloged over all graphs  $G_i$  in the database **D**, we compute their support by counting the number of distinct graph ids that contribute to each extension. Finally, the method returns the set of all extensions and their supports in sorted order (increasing) based on the tuple comparison operator in Eq. (11.1).

**Example 11.9.** Consider the canonical code C and the corresponding graph G(C) shown in Figure 11.9a. For this code all the vertices are on the rightmost path, that is,  $R = \{0, 1, 2\}$ , and the rightmost child is  $u_r = 2$ .

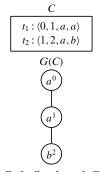
The sets of all possible isomorphisms from C to graphs  $G_1$  and  $G_2$  in the database (shown in Figure 11.7) are listed in Figure 11.9b as  $\Phi_1$  and  $\Phi_2$ . For example, the first isomorphism  $\phi_1: G(C) \to G_1$  is defined as

$$\phi_1(0) = 10$$
  $\phi_1(1) = 30$   $\phi_1(2) = 20$ 

## ALGORITHM 11.2. Rightmost Path Extensions and Their Support

```
RIGHTMOSTPATH-EXTENSIONS (C, D):
1 R \leftarrow nodes on the rightmost path in C
2 u_r \leftarrow \text{rightmost child in } C // \text{ dfs number}
3 \mathcal{E} \leftarrow \emptyset // set of extensions from C
4 foreach G_i \in \mathbf{D}, i = 1, \dots, n do
       if C = \emptyset then
            // add distinct label tuples in G_i as forward
                 extensions
            foreach distinct \langle L(x), L(y), L(x, y) \rangle \in G_i do
6
                 f = \langle 0, 1, L(x), L(y), L(x, y) \rangle
7
                Add tuple f to \mathcal{E} along with graph id i
8
       else
            \Phi_i = \text{SUBGRAPHISOMORPHISMS}(C, G_i)
10
            foreach isomorphism \phi \in \Phi_i do
11
                // backward extensions from rightmost child
                foreach x \in N_{G_i}(\phi(u_r)) such that \exists v \leftarrow \phi^{-1}(x) do
12
                     if v \in R and (u_r, v) \notin G(C) then
13
                         b = \langle u_r, v, L(u_r), L(v), L(u_r, v) \rangle
14
                         Add tuple b to \mathcal{E} along with graph id i
15
                 // forward extensions from nodes on rightmost path
                foreach u \in R do
16
                     foreach x \in N_{G_i}(\phi(u)) and \not\exists \phi^{-1}(x) do
17
                         f = \langle u, u_r + 1, L(\phi(u)), L(x), L(\phi(u), x) \rangle
18
                         Add tuple f to \mathcal{E} along with graph id i
19
   // Compute the support of each extension
20 foreach distinct extension s \in \mathcal{E} do
       sup(s) = number of distinct graph ids that support tuple s
22 return set of pairs \langle s, sup(s) \rangle for extensions s \in \mathcal{E}, in tuple sorted order
```

The list of possible backward and forward extensions for each isomorphism is shown in Figure 11.9c. For example, there are two possible edge extensions from the isomorphism  $\phi_1$ . The first is a backward edge extension  $\langle 2,0,b,a\rangle$ , as (20,10) is a valid backward edge in  $G_1$ . That is, the node x=10 is a neighbor of  $\phi(2)=20$  in  $G_1$ ,  $\phi^{-1}(10)=0=v$  is on the rightmost path, and the edge (2,0) is not already in G(C), which satisfy the backward extension steps in lines 12–15 in Algorithm 11.2. The second extension is a forward one  $\langle 1,3,a,b\rangle$ , as  $\langle 30,40,a,b\rangle$  is a valid extended edge in  $G_1$ . That is, x=40 is a neighbor of  $\phi(1)=30$  in  $G_1$ , and node 40 has not already been mapped to any node in G(C), that is,  $\phi_1^{-1}(40)$  does not exist. These conditions satisfy the forward extension steps in lines 16–19 in Algorithm 11.2.



(a) (	Code	C	and	graph	G(	C)
-------	------	---	-----	-------	----	----

Φ	φ	0	1	2
	$\phi_1$	10	30	20
$\Phi_1$	$\phi_2$	10	30	40
	$\phi_3$	30	10	20
	$\phi_4$	60	80	70
$\Phi_2$	$\phi_5$	80	60	50
	$\phi_6$	80	60	70

(b) Subgraph isomorphisms

Id	φ	Extensions
	$\phi_1$	$\{\langle 2,0,b,a\rangle,\langle 1,3,a,b\rangle\}$
$G_1$	$\phi_2$	$\{\langle 1,3,a,b\rangle,\langle 0,3,a,b\rangle\}$
	$\phi_3$	$ \{\langle 2, 0, b, a \rangle, \langle 1, 3, a, b \rangle \} $ $ \{\langle 1, 3, a, b \rangle, \langle 0, 3, a, b \rangle \} $ $ \{\langle 2, 0, b, a \rangle, \langle 0, 3, a, b \rangle \} $
$G_2$	$\phi_5$	$\{\langle 2,3,b,b\rangle,\langle 1,3,a,b\rangle\}$
	$\phi_6$	
•	-	(c) Edge extensions

	1.1
$\langle 2, 0, b, a \rangle$	2
$\langle 2, 3, b, b \rangle$	1
$\langle 1,3,a,b\rangle$	2
$\langle 0, 3, a, b \rangle$	2
	1) 1

Extension | Support

(d) Extensions (sorted) and supports

Figure 11.9. Rightmost path extensions.

Given the set of all the edge extensions, and the graph ids that contribute to them, we obtain support for each extension by counting how many graphs contribute to it. The final set of extensions, in sorted order, along with their support values is shown in Figure 11.9d. With minsup = 2, the only infrequent extension is  $\langle 2, 3, b, b \rangle$ .

#### **Subgraph Isomorphisms**

The key step in listing the edge extensions for a given code C is to enumerate all the possible isomorphisms from C to each graph  $G_i \in \mathbf{D}$ . The function SUBGRAPHI-SOMORPHISMS, shown in Algorithm 11.3, accepts a code C and a graph G, and returns the set of all isomorphisms between C and G. The set of isomorphisms  $\Phi$  is initialized by mapping vertex 0 in C to each vertex x in G that shares the same label as 0, that is, if L(x) = L(0) (line 1). The method considers each tuple  $t_i$  in C and extends the current set of partial isomorphisms. Let  $t_i = \langle u, v, L(u), L(v), L(u, v) \rangle$ . We have to check if each isomorphism  $\phi \in \Phi$  can be extended in G using the information from  $t_i$  (lines 5–12). If  $t_i$  is a forward edge, then we seek a neighbor x of  $\phi(u)$  in G such that x has not already been mapped to some vertex in C, that is,  $\phi^{-1}(x)$  should not exist, and the node and edge labels should match, that is, L(x) = L(v), and  $L(\phi(u), x) = L(u, v)$ . If so,  $\phi$  can be extended with the mapping  $\phi(v) \to x$ . The new extended isomorphism, denoted  $\phi'$ , is added to the initially empty set of isomorphisms  $\Phi'$ . If  $t_i$  is a backward edge, we have to check if  $\phi(v)$  is a neighbor of  $\phi(u)$  in G. If so, we add the current isomorphism  $\phi$  to  $\Phi'$ . Thus,

### **ALGORITHM 11.3. Enumerate Subgraph Isomorphisms**

```
SUBGRAPHISOMORPHISMS (C = \{t_1, t_2, \dots, t_k\}, G):
1 \Phi \leftarrow \{\phi(0) \rightarrow x \mid x \in G \text{ and } L(x) = L(0)\}
2 foreach t_i \in C, i = 1, \ldots, k do
        \langle u, v, L(u), L(v), L(u, v) \rangle \leftarrow t_i // \text{ expand extended edge } t_i
3
        \Phi' \leftarrow \emptyset // partial isomorphisms including t_i
4
        foreach partial isomorphism \phi \in \Phi do
5
             if v > u then
6
                  // forward edge
                  foreach x \in N_G(\phi(u)) do
7
                       if \not\exists \phi^{-1}(x) and L(x) = L(v) and L(\phi(u), x) = L(u, v) then
8
                            \phi' \leftarrow \phi \cup \{\phi(v) \rightarrow x\}
                            Add \phi' to \Phi'
10
             else
11
                  // backward edge
                  if \phi(v) \in N_{G_i}(\phi(u)) then \operatorname{Add} \phi to \Phi' // valid isomorphism
12
        \Phi \leftarrow \Phi' // update partial isomorphisms
13
14 return Φ
```

only those isomorphisms that can be extended in the forward case, or those that satisfy the backward edge, are retained for further checking. Once all the extended edges in C have been processed, the set  $\Phi$  contains all the valid isomorphisms from C to G.

**Example 11.10.** Figure 11.10 illustrates the subgraph isomorphism enumeration algorithm from the code C to each of the graphs  $G_1$  and  $G_2$  in the database shown in Figure 11.7.

For  $G_1$ , the set of isomorphisms  $\Phi$  is initialized by mapping the first node of C to all nodes labeled a in  $G_1$  because L(0)=a. Thus,  $\Phi=\{\phi_1(0)\to 10,\phi_2(0)\to 30\}$ . We next consider each tuple in C, and see which isomorphisms can be extended. The first tuple  $t_1=\langle 0,1,a,a\rangle$  is a forward edge, thus for  $\phi_1$ , we consider neighbors x of 10 that are labeled a and not included in the isomorphism yet. The only other vertex that satisfies this condition is 30; thus the isomorphism is extended by mapping  $\phi_1(1)\to 30$ . In a similar manner the second isomorphism  $\phi_2$  is extended by adding  $\phi_2(1)\to 10$ , as shown in Figure 11.10. For the second tuple  $t_2=\langle 1,2,a,b\rangle$ , the isomorphism  $\phi_1$  has two possible extensions, as 30 has two neighbors labeled b, namely 20 and 40. The extended mappings are denoted  $\phi_1'$  and  $\phi_1''$ . For  $\phi_2$  there is only one extension.

The isomorphisms of C in  $G_2$  can be found in a similar manner. The complete sets of isomorphisms in each database graph are shown in Figure 11.10.

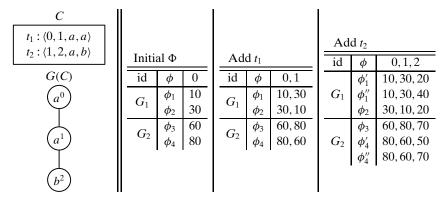


Figure 11.10. Subgraph isomorphisms.

## 11.3.2 Canonicality Checking

Given a DFS code  $C = \{t_1, t_2, \dots, t_k\}$  comprising k extended edge tuples and the corresponding graph G(C), the task is to check whether the code C is canonical. This can be accomplished by trying to reconstruct the canonical code  $C^*$  for G(C) in an iterative manner starting from the empty code and selecting the least rightmost path extension at each step, where the least edge extension is based on the extended tuple comparison operator in Eq. (11.1). If at any step the current (partial) canonical DFS code  $C^*$  is smaller than C, then we know that C cannot be canonical and can thus be pruned. On the other hand, if no smaller code is found after k extensions then C must be canonical. The pseudo-code for canonicality checking is given in Algorithm 11.4. The method can be considered as a restricted version of gSpan in that the graph G(C) plays the role of a graph in the database, and  $C^*$  plays the role of a candidate extension. The key difference is that we consider only the smallest rightmost path edge extension among all the possible candidate extensions.

## ALGORITHM 11.4. Canonicality Checking: Algorithm IsCanonical

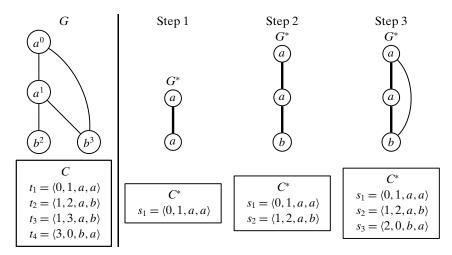


Figure 11.11. Canonicality checking.

**Example 11.11.** Consider the subgraph candidate  $C_{14}$  from Figure 11.8, which is replicated as graph G in Figure 11.11, along with its DFS code C. From an initial canonical code  $C^* = \emptyset$ , the smallest rightmost edge extension  $s_1$  is added in Step 1. Because  $s_1 = t_1$ , we proceed to the next step, which finds the smallest edge extension  $s_2$ . Once again  $s_2 = t_2$ , so we proceed to the third step. The least possible edge extension for  $G^*$  is the extended edge  $s_3$ . However, we find that  $s_3 < t_3$ , which means that C cannot be canonical, and there is no need to try further edge extensions.

#### 11.4 FURTHER READING

The gSpan algorithm was described in Yan and Han (2002), along with the notion of canonical DFS code. A different notion of canonical graphs using canonical adjacency matrices was described in Huan, Wang, and Prins (2003). Level-wise algorithms to mine frequent subgraphs appear in Kuramochi and Karypis (2001) and Inokuchi, Washio, and Motoda (2000). Markov chain Monte Carlo methods to sample a set of representative graph patterns were proposed in Al Hasan and Zaki (2009). For an efficient algorithm to mine frequent tree patterns see Zaki (2002).

Al Hasan, M. and Zaki, M. J. (2009). "Output space sampling for graph patterns." *Proceedings of the VLDB Endowment*, 2(1): 730–741.

Huan, J., Wang, W., and Prins, J. (2003). "Efficient mining of frequent subgraphs in the presence of isomorphism." *In Proceedings of the IEEE International Conference on Data Mining*. IEEE, pp. 549–552.

Inokuchi, A., Washio, T., and Motoda, H. (2000). "An apriori-based algorithm for mining frequent substructures from graph data." *In Proceedings of the European Conference on Principles of Data Mining and Knowledge Discovery.* Springer, pp. 13–23.

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Kuramochi, M. and Karypis, G. (2001). "Frequent subgraph discovery." *In Proceedings of the IEEE International Conference on Data Mining*. IEEE, pp. 313–320.

- Yan, X. and Han, J. (2002). "gSpan: Graph-based substructure pattern mining." *In Proceedings of the IEEE International Conference on Data Mining*. IEEE, pp. 721–724.
- Zaki, M. J. (2002). "Efficiently mining frequent trees in a forest." In Proceedings of the 8th ACM SIGKDD International Conference on Knowledge Discovery and Data Mining. ACM, pp. 71–80.

#### 11.5 EXERCISES

**Q1.** Find the canonical DFS code for the graph in Figure 11.12. Try to eliminate some codes without generating the complete search tree. For example, you can eliminate a code if you can show that it will have a larger code than some other code.

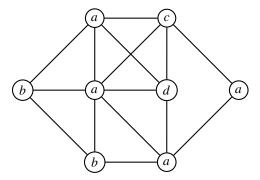


Figure 11.12. Graph for Q1.

**Q2.** Given the graph in Figure 11.13. Mine all the frequent subgraphs with minsup = 1. For each frequent subgraph, also show its canonical code.

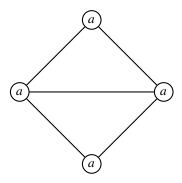


Figure 11.13. Graph for Q2.

**Q3.** Consider the graph shown in Figure 11.14. Show all its isomorphic graphs and their DFS codes, and find the canonical representative (you may omit isomorphic graphs that can definitely not have canonical codes).

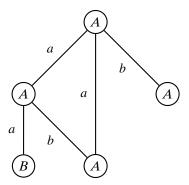


Figure 11.14. Graph for Q3.

**Q4.** Given the graphs in Figure 11.15, separate them into isomorphic groups.

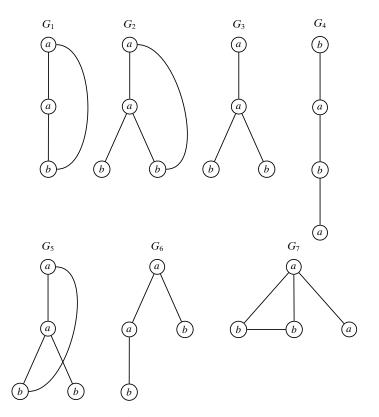


Figure 11.15. Data for Q4.

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**Q5.** Given the graph in Figure 11.16. Find the *maximum* DFS code for the graph, subject to the constraint that all extensions (whether forward or backward) are done only from the right most path.

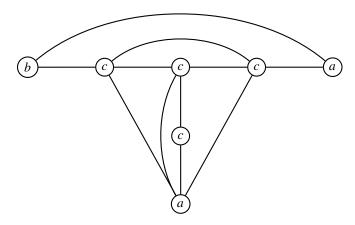


Figure 11.16. Graph for Q5.

**Q6.** For an edge labeled undirected graph G = (V, E), define its labeled adjacency matrix **A** as follows:

$$\mathbf{A}(i,j) = \begin{cases} L(v_i) & \text{if } i = j \\ L(v_i, v_j) & \text{if } (v_i, v_j) \in E \\ 0 & \text{Otherwise} \end{cases}$$

where  $L(v_i)$  is the label for vertex  $v_i$  and  $L(v_i, v_j)$  is the label for edge  $(v_i, v_j)$ . In other words, the labeled adjacency matrix has the node labels on the main diagonal, and it has the label of the edge  $(v_i, v_j)$  in cell  $\mathbf{A}(i, j)$ . Finally, a 0 in cell  $\mathbf{A}(i, j)$  means that there is no edge between  $v_i$  and  $v_i$ .

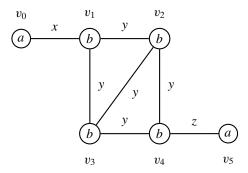


Figure 11.17. Graph for Q6.

Given a particular permutation of the vertices, a *matrix code* for the graph is obtained by concatenating the lower triangular submatrix of **A** row-by-row. For

example, one possible matrix corresponding to the default vertex permutation  $v_0v_1v_2v_3v_4v_5$  for the graph in Figure 11.17 is given as

а					
х	b				
0	у	b			
0	у	у	b		
0	0	у	у	b	
0	0	0	0	z	а

The code for the matrix above is axb0yb0yyb000yzb0000za. Given the total ordering on the labels

find the maximum matrix code for the graph in Figure 11.17. That is, among all possible vertex permutations and the corresponding matrix codes, you have to choose the lexicographically largest code.