



FINAL YEAR DISSERTATION

Market Basket Analysis with Graph Theory

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Declaration

I, Sahil Manojkumar Pattni, confirm that this work submitted for assessment is my own and is expressed in my own words. Any uses made within it of the works of other authors in any form (e.g., ideas, equations, figures, text, tables, programs) are properly acknowledged at any point of their use. A list of the references employed is included.

Date: April 14, 2021

Signed: Sahil Manojkumar Pattni

Abstract

In this digital age, data is being generated and collected at an unprecedented rate, with data analytics employed by corporations and small businesses alike to produce actionable insights, reduce costs, optimize operations and increase revenue. Association rules allow us to identify relationships between products that can provide insights into customer spending habits and product perception.

In this study, a minimum spanning tree (MST) will be generated from a transactional database such that only the strongest relationships between products remain. A clustering algorithm will be applied to this MST to identify high co-purchase segments, and association rules will then be extracted from these segments. **[ADD MORE HERE]**

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Chapter 1

Introduction

1.1 Motivation

For businesses such as groceries, hypermarkets, and retail outlets that deal with the trade of heterogeneous physical assets, operations such as inventory management and product placement play an instrumental role in determining the business' financial success. These involve asking questions such as:

- Which products should be placed at the entrance of the store? Which should be placed closer to the exit?
- Which products will benefit the most by being placed at eye-level?
- Which products should be placed next to each other to maximize the purchase volume?

One way to find optimal solutions to such queries is to employ the use of Association Rule Mining (also known as Market Basket Analysis). This set of techniques assess frequent itemsets (e.g. from sales data) and generate association rules between products. A prime example of the utility of association rules is the urban legend of "*Beers and Pampers*", where a company allegedly studied their point-of-sale data and found a strong association between the purchase of diapers and a particular brand of beer during a certain time. With *diapers* as the antecedent and *beer* as the consequent, this rule can be written as:

$$\{Diapers\} \rightarrow \{Beer\}$$

This is an example of a single-element rule, where both the antecedent and consequent are sets that contain only one element each. Several algorithms exist for association rule mining, most prominently the Apriori Algorithm (**Agrawal and Srikant 1994**) and FP-Growth (**Han et al. 2000**), however these algorithms tend to generate an overwhelming amount of rules, rendering it inconvenient for the end-user to extract actionable information from the ruleset. This paper proposes to improve upon an existing method (see: Section 2.4.5) that derives association rules from minimum spanning trees.

1.2 Aims

The aim of this paper is to improve upon the aforementioned method by introducing a method that allows for the generation of multi-element association rules (i.e. rules where either/both the antecedent and consequent of the rule contain more than one element).

1.3 Objectives

The research objectives for this paper are as described below:

1. Acquire a suitable dataset upon which the study can be conducted.
2. Construct an affinity graph from the chosen dataset.
3. Explore and evaluate MST extraction algorithms.
4. Identify and evaluate methodologies for generating multi-element association rules from an MST.
5. Extract MST from the graph and generate association-rules using the chosen methodologies.
6. Evaluate the generated association rules against the rules generated by the Apriori Algorithm.

Chapter 2

Background

2.1 Transaction and Association Representation

2.1.1 Binary Purchase Vectors

Let $I = I_1, I_2, \dots, I_m$ be a set of binary attributes (i.e. items), and let T be a database of transactions. As defined in (Agrawal et al. 1993), a binary purchase vector is a transaction t represented as a vector of length m , where:

$$t[k] = \begin{cases} 1 & \text{if } I_k \text{ purchased in } t \\ 0 & \text{otherwise} \end{cases}$$

For example, consider a grocer who only sells five items: milk, eggs, bread, apples and oranges. Consider the following transactions:

Transaction t_1 : Customer purchases bread and oranges.

Transaction t_2 : Customer purchases eggs, bread and apples.

Transaction t_3 : Customer purchases milk and oranges.

The binary purchase vectors for these transactions would be:

	milk	eggs	bread	apples	oranges
t_1	0	0	1	0	1
t_2	0	1	1	1	0
t_3	1	0	0	0	1

Table 2.1: Binary Purchase Vectors

2.2 Product Association

The association between products pairs across all transactions can be ascertained from the binary purchase vectors using the Pearson's Correlation Coefficient (**Pearson 1895**). Since the correlations are of two binary variables, the Pearson's Correlation Coefficient is equivalent to the Phi-Coefficient ϕ (**Ernest C and El-Sanhurry 1991**), where for n observations:

$$\phi = \sqrt{\frac{\chi^2}{n}}$$

Applying this correlation formula to the set of binary purchase vectors in Table 2.1, we get:

	milk	eggs	bread	apples	oranges
milk	1.0	-0.5	-1.0	-0.5	0.5
eggs	-0.5	1.0	0.5	1.0	-1.0
bread	-1.0	0.5	1.0	0.5	-0.5
apples	-0.5	1.0	0.5	1.0	-1.0
oranges	0.5	-1.0	-0.5	-1.0	1.0

Table 2.2: Correlation Matrix

Note that the correlation matrix is diagonally symmetrical. This is true of all correlation matrices where the items on the x-axis and y-axis are the same and in the same order. This correlation matrix can then be represented as a graph, where the nodes are the products, and the edges are the correlation values.

2.3 Graph Theory

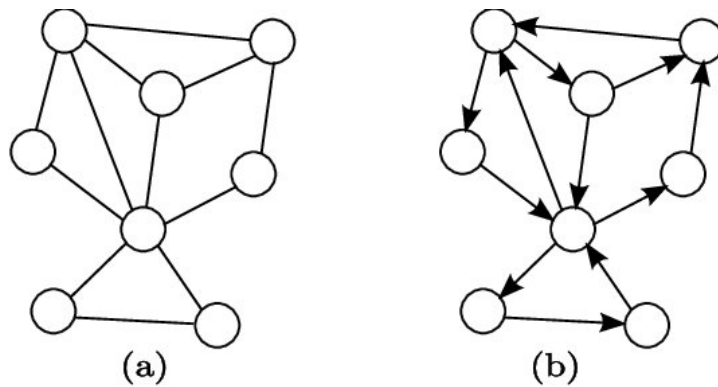


Figure 2.1: Undirected and Directed Graphs

In discrete mathematics and more specifically - graph theory, a graph is a data structure that contains a set of nodes (i.e. vertices) connected by lines (i.e. edges). These edges may be directed - such as in Figure 2.1:(a), or undirected - such as in Figure 2.1:(b). The edges may contain values (i.e. weights) between the two vertices it connects. A graph G with a set of vertices V and a set of edges E can be represented via the notation $G = (V, E)$. For the scope of this project, we will be building undirected graphs, where the weight between two vertices is the same in both directions.

2.3.1 Minimum Spanning Trees

Given an undirected $G = (V, E)$, a *spanning tree* can be described as a subgraph that is a tree which includes all the vertices V of G with the minimum number of edges required. A *minimum spanning tree* (MST) is the spanning tree with the smallest sum of edge weights. This means that if the graph has n vertices, each spanning tree - including the minimum spanning tree - will have $n - 1$ edges. Since a minimum spanning tree captures the lowest weights in a graph, with modifications it could be an excellent candidate to capture the opposite as well: the strongest associations between products. There are two widely used algorithms to extract the minimum spanning tree from a graph: Prim's algorithm and Kruskal's algorithm.

2.3.2 Prim's Algorithm

Independently discovered by three authors, Prim's algorithm (**Prim 1957**)(**Jarník 1930**)(**Dijkstra 1959**) is a greedy algorithm¹ to find the minimum spanning tree of an undirected, weighted graph G . To successfully implement the algorithm, three sets need to be maintained: a set of *discovered* edges, and two sets of vertices: a set of *undiscovered* vertices, and a set of *discovered* vertices. Figure 2.2 illustrates Prim's algorithm being applied to a graph. The algorithm is as follows:

Initialize an empty set of discovered edges: E , and two sets of vertices: an empty set D of the discovered vertices, and UD as the set of undiscovered vertices.

- Pick an arbitrary vertex as a starting point (in the case of Figure 2.2, the top right node). Add this vertex to D and remove it from UD .
- While UD is not empty:
 - Find the edge e_i with the smallest weight such that it connects together a vertex V_1 in D and V_2 in UD (to avoid forming cycles).

¹Selecting the locally optimal choice at each iteration of the solution

- Append V_2 to D and remove it from UD (i.e. V_2 is now discovered).
- Append e_i to E .

Once D contains all the vertices of G , the algorithm terminates, and the set D represents the minimum spanning tree, and $\sum_{i=1}^n e_i$ is the weight of the MST. The time complexity of this algorithm is $O(V^2)$.

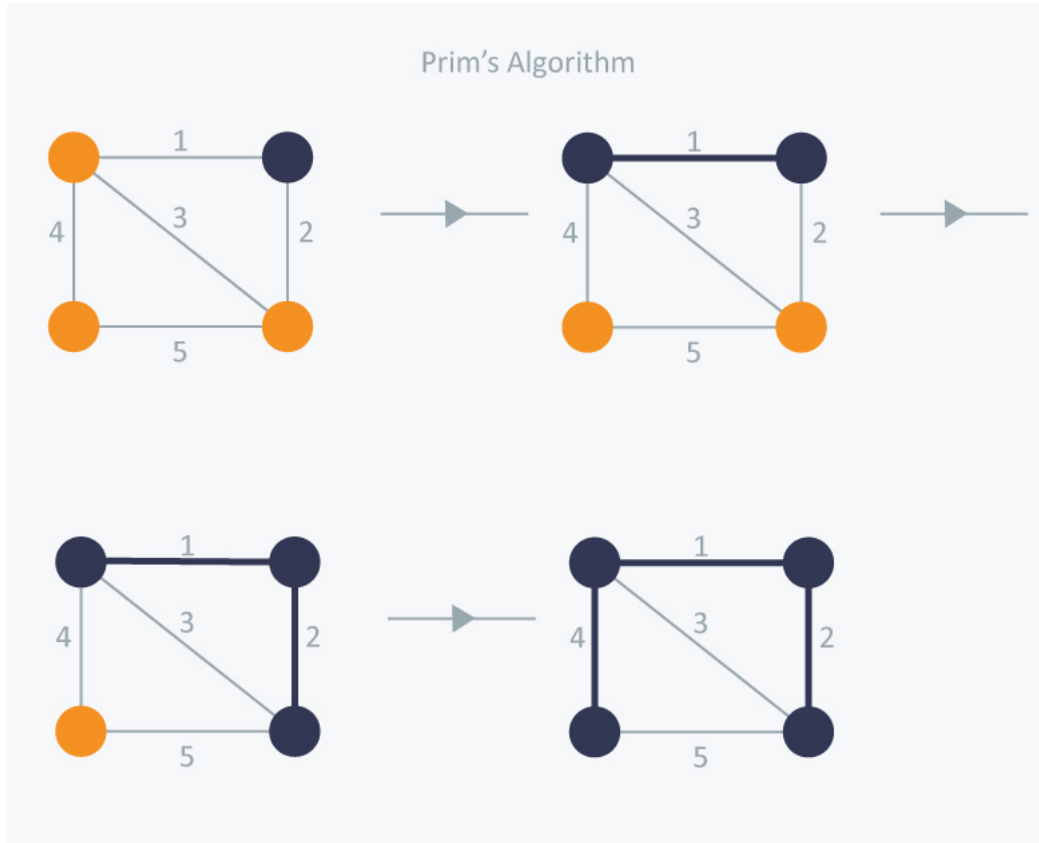


Figure 2.2: Prim's Algorithm applied to a graph (Abdelnabi 2018)

2.3.3 Kruskal's Algorithm

Another greedy algorithm, Kruskal's algorithm (Kruskal 1956) also extracts the MST from a graph. Unlike Prim's algorithm, Kruskal's doesn't select an edge that connects directly to the already built spanning tree, but rather picks the global optimal solution. The algorithm is as follows:

Maintaining a set of edges E , and an initially empty set of chosen edges C :

- Sort the set of edges E in ascending order.
- While the number of elements in C is not $n - 1$:

- Select the smallest edge e_i from E .
- If adding it does not form a cycle with the spanning tree formed so far, append e_i to C .
- Remove e_i from E .

The algorithm will terminate when $n - 1$ edges have been selected. The time complexity for this algorithm is $O(E \log E)$.

2.4 Related Work

2.4.1 Extracting Minimum Spanning Trees using K-Means

(Zhong et al. 2015) proposed a novel framework to extract the minimum spanning tree of a graph based on the K-Means clustering algorithm. Their methodology can be separated into two distinct phases. In the first phase, the data is partitioned into \sqrt{n} clusters via K-Means and the Kruskal's algorithm is applied to each of the clusters individually. Once the \sqrt{n} MSTs have been constructed, they are combined to form an approximate MST. In the second phase, new partitions are constructed based on the borders of the clusters identified in phase 1. Based on these new partitions, a second approximate MST is constructed. Finally, both graphs are merged such that the resulting graph has $2(n - 1)$ edges. The Kruskal's algorithm is run on this graph to get the final approximation of the MST.

Critical Analysis

The authors have proposed an efficient way to approximate a minimum spanning tree, with their methodology having a complexity of $O(N^{1.5})$, which is faster than the standard MST algorithm which has a complexity of $O(N^2)$. For clarity, we have illustrated the disparity between the author's algorithm and the standard algorithm on Figure 2.3.

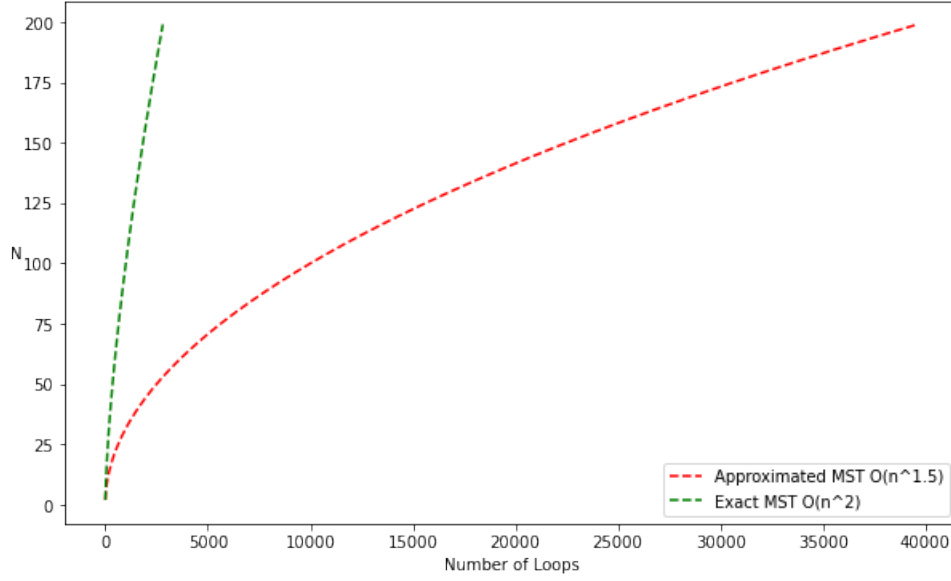


Figure 2.3: Efficiency of K-Means optimized MST vs. exact MST

The K-Means optimized algorithm is $\frac{\sqrt{n}-1}{\sqrt{n}}\%$ faster than the standard MST algorithm.

2.4.2 Markov Clustering

(Dongen 1969) introduced the novel Markov Clustering Algorithm (MCL), a graph clustering algorithm based on Markov Processes (Markov 2006) that favours sparse graphs (i.e. graphs where the average degree is smaller than the number of nodes). The reason for this follows the notion that dense regions in sparse graphs correspond with regions in which the number of k -length paths is relatively large - and therefore - paths whose beginnings and ends are in the same dense region have a higher probability for random walks² of length k as opposed to other paths. In other words, random walks originating from a node in a dense region have a high probability of ending in that same dense region. Given a non-negative weighted directional graph G , the MCL simulates flow in the graph. It does this by first mapping the graph G in a generic way onto a Markov matrix M_1 . Once completed, the set of transition probabilities (TP's) are iteratively recomputed via expansion and contraction, resulting in an array of Markov Graphs. In the expansion stage, higher TP's are calculated, whereas for the contraction phase, a new Markov Graph is created by rewarding high TP's and penalizing low TP's. The belief is that by doing so, the flow between dense regions that are sparsely connected will be removed, leaving only distinct and unconnected dense regions (i.e. clusters). The author tested the MCL

²A random walk is a stochastic process of of successively random steps along a space, in this case: a directed graph.

algorithm against randomly generated graphs which were known to possess a natural cluster structure. The authors noted that the MCL algorithm successfully managed to capture the segregation present in the graphs, and concluded that the algorithm was capable of handling graphs with large natural clusters.

Critical Analysis

The author proposed a novel way to apply the mathematics behind Markov processes to successfully segment and isolate densely connected segments of a graph. The results from this paper have inspired us to use the clustering algorithm in our own work.

2.4.3 Apriori Algorithm

(Agrawal et al. 1993) proposed a novel algorithm to generate all statistically significant association rules between items in a database, laying the foundation for association rule mining. Given a set of items $I = I_1, I_2, I_3, \dots, I_m$, the authors define an association rule to be of the form $X \rightarrow I_j$ where X is a set of items such that $X \in I, I_j \notin X$. The hypothetical database stated was a list of transactions, T , where each transaction t was a binary vector of length m , as described in Section 2.1.1. The authors define two constraints for assessing association rules:

Support

The support of an association rule is the proportion of transactions in which the itemsets in the rule are present. For a set of transactions T , where $T(i)$ denotes the set of transactions in which the set of items $i \in I$ was purchased:

$$\text{support}(i) = \frac{T(i)}{T}$$

Similarly, where $T(i_k, i_j)$ represents the set of transactions in which the itemsets $\{i_k \in I, i_j \in I\}$ were purchased, the support for the association rule $i_k \rightarrow i_j$ can be represented as:

$$\text{support}(i_k \rightarrow i_j) = \frac{T(i_k, i_j)}{T}$$

Support scores correspond with the statistical significance of a rule, and rules with low support scores may not occur frequently enough to draw reasonable conclusions from.

Confidence

Confidence is the conditional probability of an itemset $i_j \in I$ being present in a transaction given that itemset $i_k \in I$ is present in the same transaction. The confidence of the association rule $i_k \rightarrow i_j$ can be represented as:

$$\text{confidence}(i_k \rightarrow i_j) = \frac{T(i_k, i_j)}{T(i_k)} \equiv \frac{\text{support}(i_k \rightarrow i_j)}{\text{support}(i_k)}$$

The confidence of an association rule can be thought of as the rule's *strength*.

With these constraints defined, the authors state that their methodology for association rule mining can be split into two discrete steps:

1. The generation of candidate itemsets.
2. The generation of statistically significant association rules from the itemsets.

Candidate Itemset Generation

To generate candidate itemsets, the authors first generate all possible itemsets from the database, defining those whose support score was above a support constraint \min_{support} as *large itemsets*. The authors note that a brute-force check³ would be sub-optimal, taking up to 2^m passes of the database (where m is the number of items in the itemset I). Therefore, they introduced a methodology where they would only observe itemsets of length k on the k^{th} pass of the dataset, to see if the itemsets satisfied the support constraint. On the $(k+1)^{\text{th}}$ pass of the dataset, they need only check itemsets that are *1-extensions* (i.e. itemsets extended by only one item) or the *large itemsets* discovered in the previous pass. Their reasoning is now commonly known as *The Apriori Principle*, where they state that if an itemset i_k is *large* (i.e. satisfies the support constraint), then any subset $i_j \subset i_k$ will also be *large*. This reasoning also applies that if an itemset i_j is found to be *small* (i.e. did not satisfy the support constraint), then any superset $i_k; i_j \subset i_k$ will also be *small*. This allows the authors to prune the number of association rules whose support scores need to be computed, as if they find i_j to be *small*, any superset $i_k; i_j \subset i_k$ need not have its support score computed as it is known to be *small*.

However, if an itemset i_j is indeed found to be *large*, then another multiple passes over the dataset will be required to check the support scores for subsets of i_j . To avoid this, the authors devised a measure to calculate the expected support \bar{s} of an itemset. The expected support is used to estimate the support of $i_j = (i_p + i_q)$, not only when i_j is expected to be *large*, but also when i_p is expected to be *large* yet $(i_p + i_q)$ is expected to be *small*. This estimation further prunes the number of rules whose support scores need to be computed.

³checking every possible itemset iteratively.

Association Rule Generation

To generate association rules, the authors used the following technique:

for each *large* itemset $Y = i_1, i_2, \dots, i_k; k \geq 2$ from the set of non-pruned *large* itemsets, generate a set of association rules in the form $X \rightarrow i_j; X \subset Y, i_j \notin X$ such that X is of length $k - 1$. Therefore, each *large* itemset will produce k rules. From the generated rules, the authors discarded those rules whose confidence scores fell below the confidence constraint $min_{confidence}$.

Evaluation

The authors tested their methodology on a sales dataset with 46,783 transactions, with 63 distinct departments. Their configuration was composed of a support constraint of 1% (i.e. $min_{support} = 0.01$) and a confidence constraint of 50% (i.e. $min_{confidence} = 0.5$). The authors note that the rules produced follow what general intuition might suggest. For example:

$$\{\text{Auto Accessories, Tires}\} \rightarrow \{\text{Automotive Services}\}$$

Furthermore, the authors assessed the accuracy of their support estimation metric \bar{s} by observing the ratio of correctly estimated itemsets for both *small* and *large* against various values for the support constraint. They were able to conclude that their estimation accuracy was satisfactory, as their accuracy was 96% and above for support thresholds.

Critical Analysis

The authors have proposed a novel methodology that has been the bedrock of numerous research publications, including most of the papers in this literature review. Their estimation function performed with high accuracy, meaning it can reduce the number of passes through a database the algorithm has to take by a significant amount. Additionally, their pruning techniques allowed them to eliminate a large proportion of itemsets from the space. Even after the significant pruning of rules, a major drawback of this methodology is the large number of rules produced, although one could argue that only the highest performing rules need be observed in further detail.

2.4.4 Subjective Measurement of Association Rules

(Zekic-Susac and Has 2015) proposed a novel measure for the *interestingness* of association rules, identifying that a dominant, universally used measure did not exist. The authors' goal was to combine

objective measures such as the support, confidence and lift scores with more subjective measures. Instead of the Apriori approach, their methodology has them generate association rules via the *tree-building technique* - which compresses a large database into a Frequent-Pattern tree, citing that this technique was more efficient than the Apriori algorithm. The authors employed the heuristical unexpectedness measure⁴ and the heuristical actionability measure⁵ as their subjective measures, and a minimum confidence threshold of 51% as their objective measure. Since a subjective measure would require a human subject, the authors' used the estimations of a sales manager from a Croatian retail chain, and stored his responses in binary format for the subjective measures (i.e. 0 if a rule was unexpected else 1, 0 if a rule is not useful else 1). The dataset used for this paper was a real transactional dataset with 14,012 transactions and a set of 1,230 unique items (which was later pruned to 7,006 transactions and a set of 278 products) from the same Croatian retail chain their test subject worked at. The authors then generated association rules from the first-level hierarchical grouping of items from the dataset (items with a minimum support of 25%), of which 36 rules were identified as statistically significant. From this set of rules, only two rules satisfied both subjective measures and the confidence constraint, and therefore these two rules were identified as highly interesting. The authors then generated association rules from the second-level hierarchical grouping of items, where items that represented the same product (but had different a manufacturer, brand etc.) were grouped together. Of the rules generated, 15 satisfied their confidence constraints and had a support value able 10%. 5 rules from this set satisfied both their subjective and objective measures, more than the previous experiment The authors were able to conclude that the increase in accuracy and number of interesting rules resulted from the second level of grouping which generalized the products.

Critical Analysis

Whereas the original measure of statistical significance introduced by R. Aggarwal et al. was purely objective, the authors of this paper have presented a well thought out approach to combining the subjective metrics with objective ones to produce a human-verified association rule set. A few caveats to note, however: their study only involved one subject, which is rarely regarded to be statistically acceptable. An ideal study would require multiple, randomly chosen subjects to offset any bias that the singular subject would have had, and in addition, the larger their subject size, the closer their collective estimations will model the total population's. Another drawback of their approach is that by using human intuition as a

⁴How significantly a rule contradicts a user's prior beliefs.

⁵If the user believes they can use the information to their advantage (e.g. a promotion).

metric, they're promoting association rules that satisfy pre-existing notions about human behavior (e.g. if someone buys milk, they'll *probably* get eggs too), however these types of rules are usually regarded as common knowledge, whereas the beauty of association rule mining is in its ability to surface association rules that - while true - seem unintuitive, and therefore are less likely to be known by the management of such organizations.

2.4.5 Association Rules from Minimum Spanning Trees

(Valle et al. 2018) proposed a novel methodology to study the structure and behavior of consumer market baskets from the topology of a minimum spanning tree which represented the interdependencies between products, and use this information to complement the association rule generation process. The input to their proposed methodology was a correlation matrix between the set of all one-hot encoded purchase vectors such that each vector denoted the presence or non-presence of each product from the dataset in that vector. The dataset used for the MST construction was a list of 1,046,804 transactions containing a set of 3,240 unique products from a large supermarket chain branch in Santiago, Chile. When building this correlation matrix, the authors opted to use the Pearson's Coefficient (**Pearson 1895**) - which is equivalent to the coefficient ϕ for binary data (**Ernest C and El-Sanhurry 1991**) - over the traditionally used Jaccard distance to compute the similarity between the binary product vectors, as the former provides both a positive and negative association between products. Additionally, they used the distance function $d_{ij} = \sqrt{2(1 - \phi_{ij})}$ to transform the correlation matrix into a distance measurement (i.e. the weight of the edges)⁶. The authors constructed a MST for 220 product subcategories, and noted that there was a significant level of grouping between product sub-categories that belonged to the same parent category. To remove edges from the MST that were not statistically significant, the authors used the mutual information (**Cover and Thomas 2006**) measure $\sum_{x,y} \log_2 \frac{r(x,y)}{p(x)q(y)}$ between product subcategories p and q , and were able to prune 14 edges, all of which were connected to a terminal node, therefore effectively pruning 14 vertices from the MST too. To identify the most influential regions of the MST, the authors defined an influence zone of distances that were in the 10th percentile. To generate meaningful association rules, for each MST product i , the authors ran a search for the set of all association rules R_i such that $P_i \rightarrow P_j (i \neq j)$. Then from the resulting set of rules, they searched for rules that obeyed $P_i \rightarrow P_m$ where m a product node connected to the product i in the minimum spanning tree. For both resulting sets of rules for each product, the mean of their lift scores were observed, and the authors determined that the rules that were reinforced by the MST had a higher mean, and that a majority of

⁶ ϕ_{ij} is the correlation score between products at indexes i and j on the correlation matrix.

these rules had a lift score above the 90th percentile.

To identify the clusters each of the products should be identified under, the authors constructed a hierarchical tree using the average linkage clustering method, and by using an unspecified cut distance, they were able to produce 17 taxonomic groups (i.e. clusters). Cross-referencing their results with the actual parent categories of the products, they were able to conclude that the MST did indeed categorize the product sub-categories into clusters with a reasonable degree of accuracy. The authors then compared their MST to another methodology, namely the structured association map (SAM) (**Kim 2017**), using the Jaccard distance as a measure of similarity, and were able to generate interesting 2x2 rules (i.e. $\{A, B\} \rightarrow \{C, D\}$), all with lift scores above 1.0, with one rule even having a lift score of 106.46. They concluded that while both approaches provided different information, they both visually identify the strongest relationships between the products, and provide useful information to reduce the search space for association rules.

Critical Analysis

The authors' approach seems to be novel, thorough and well structured. Their methodology successfully employed the use of minimum spanning trees to complement the association rule generation process with sound results. One caveat of their approach is that they only used the MST to generate single-element rules (i.e. $\{A\} \rightarrow \{B\}$, where the antecedent and consequent contain only one element each). Using the distance score in conjunction with the importance function they defined (i.e. $\sum_{k \in K_u} \frac{1}{w_{uk}}$), they could have defined a system to produce multi-element rules (i.e. where either/both the antecedent and the consequent have more than one element), then rank them using their respective lifts. While single-element rules are easily understandable and tend to have high lift values when extracted from the MST, multi-element rules would provide an additional layer of insight as to how a range of products (perhaps a cluster) relate to another.

2.4.6 Summary

In conclusion, (**Agrawal et al. 1993**) introduced a formal system for association rules, and a method to generate them from a transactional database, which was further improved when (**Agrawal and Srikant 1994**) introduced the Apriori Algorithm. Instrumental to the success of this algorithm is its ability to prune a bulk of the rules such that their support scores need not be checked. (**Pennsylvania 2020**)

defines the equation for calculating the number of possible rules for an itemset d as:

$$\text{number of rules} = \sum_{k=1}^{d-1} \left(\binom{d}{k} \times \sum_{j=1}^{d-k} \binom{d-k}{j} \right)$$

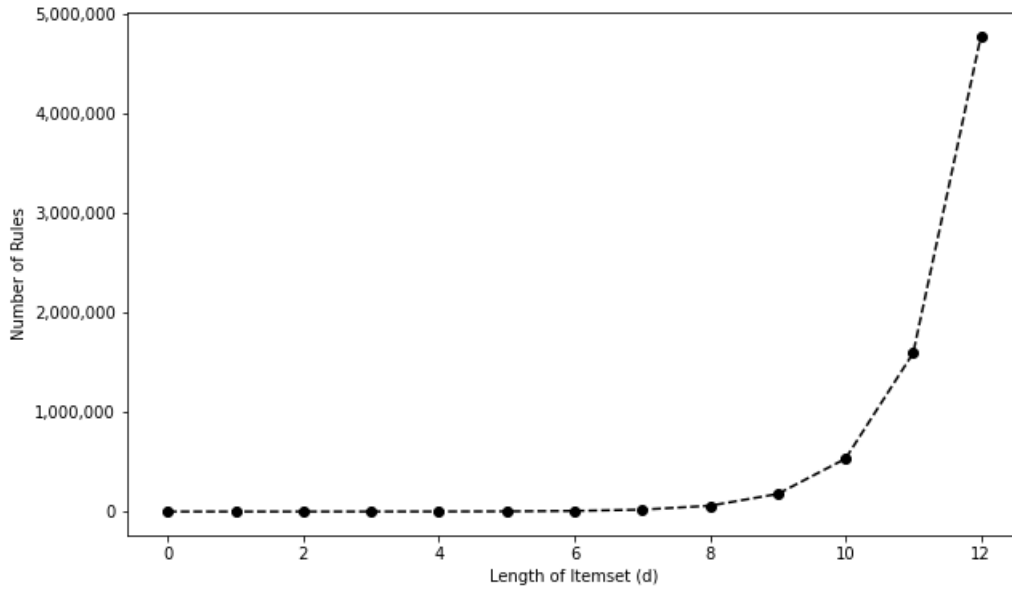


Figure 2.4: Number of Association Rules for an Itemset

We have plotted Figure 2.4 to illustrate the exponential growth of rules given an itemset's length d . This exponential growth reinforces the need for the rule pruning techniques introduced in (Agrawal et al. 1993) to be able to complete the computations within a reasonable time frame, and will therefore be incorporated into our methodology. However, while these papers only based the interestingness of an association rule on objective measures such as the support, confidence and lift scores of these rules, (Zekic-Susac and Has 2015) proposed a methodology where subjective human input was used to validate the interestingness of these rules. A drawback of the above methods, however, is the large number of rules produced. (Zhong et al. 2015) proposed a solution to the relatively slow computation time that the Prim's algorithm and Kruskal's algorithm offer, where the Kruskal's algorithm was optimized using K-Means, leading to a significant performance increase as illustrated in Figure 2.3. The methodology proposed in this paper was a framework, where Kruskal's Algorithm could be substituted with any MST algorithm, such as Prim's Algorithm. (Dongen 1969) introduced a novel way to identify and isolate

dense regions within a graph, and we have used their algorithm to cluster our own graphs. (**Valle et al. 2018**) introduced a methodology to extract high value association rules from a minimum spanning tree, used to complement the rules produced by the Apriori algorithm. A caveat of this approach, however, is that it can only produce rules such that the antecedent and consequent are sets with one element each. This paper has been the primary motivation for our work, and therefore we have incorporated into our work several techniques that the authors used and/or introduced. All the research conducted above has served as the inspiration for this project.

Chapter 3

Rule Generation via MSTs

3.1 Dataset Pre-Processing

The dataset used in this study is the transactional sales data from a chain of Brazilian gas-station stores (Kaggle 2020). Each row in the dataset represents the purchase of a specific product as part of a transaction - and as such, each row corresponds to the following columns:

- Company Code
- Order Number
- Employee
- Product
- Product Category
- Client
- Sale Date Time
- Product Cost
- Discount Amount

All personal and corporate names were exchanged for fictitious names by the author of the dataset in order to preserve the anonymity of those whose who could have otherwise been identified through the dataset. Before employing the dataset, sanitary procedures were carried out to ensure that the dataset was error-free and in a format suitable for graph generation. The steps have been detailed below.

1. Transaction Identifier

The *Order Number* field showed discrepancies, where a given order number could reference distinct transactions in different stores and cities, and at different dates and times. This could be due to

the stores maintaining their own order numbers, and also because the order numbers may reset after a predetermined limit. A unique transaction identifier - named `basket_id` - was created by concatenating the order number and the date, thereby mitigating the occurrence of a identifier that references multiple transactions.

2. Binary Purchase Vector transformation

The dataset was then transformed such that each transaction was represented by a binary purchase vector - as described in Section 2.1.1 - wherein each column represents a product category. The product categories were chosen for the graph representation over the products themselves as it would give a more generalized view on the associations between them, and the categories themselves were deemed specific enough that they would not be parent to children of significant variance.

3.2 MST Generation

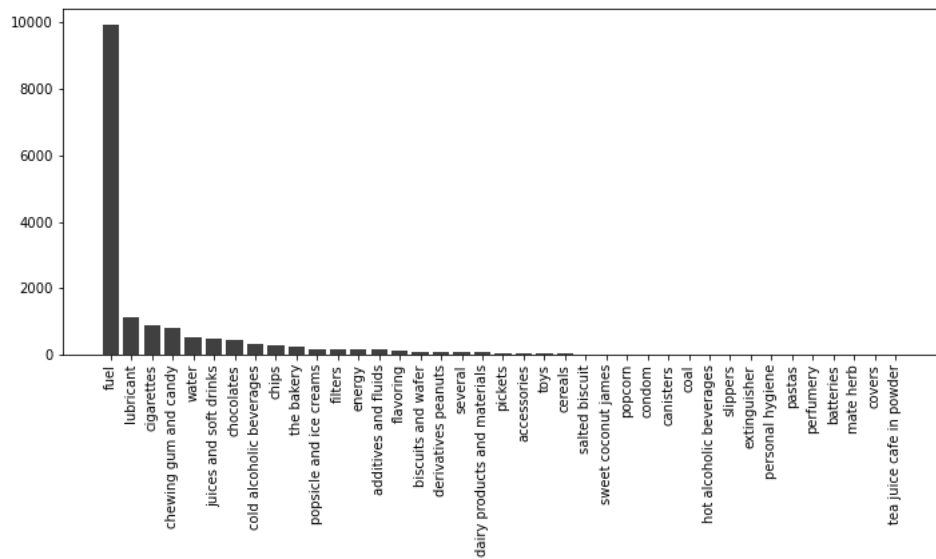


Figure 3.1: Category Distribution

The metrics used to assess the association rules - support, lift and confidence - are based on the proportional presence of a given itemset in the transactions. Since our dataset is from a gas-station store chain, fuel products dominate the transactional presence by a significant factor. Figure 3.1 highlights the disparity between the presence of *fuel* products and the others, with *fuel* being present in 99.28% of all transactions. To avoid the association rules being dominated by the *fuel* category - which should inherently understood

to be a key product for gas stations - the fuel category was purged from the dataset.

Following the methodology described in Section 2.1.1, a correlation matrix was computed from the binary purchase vectors using the Pearson's Correlation Coefficient. Since the correlations are of binary variables again, the correlation score is equivalent to the Phi-Coefficient ϕ . Figure 3.2 illustrates the correlation matrix. Only the values below the diagonal have been illustrated as the matrix is diagonally symmetrical.

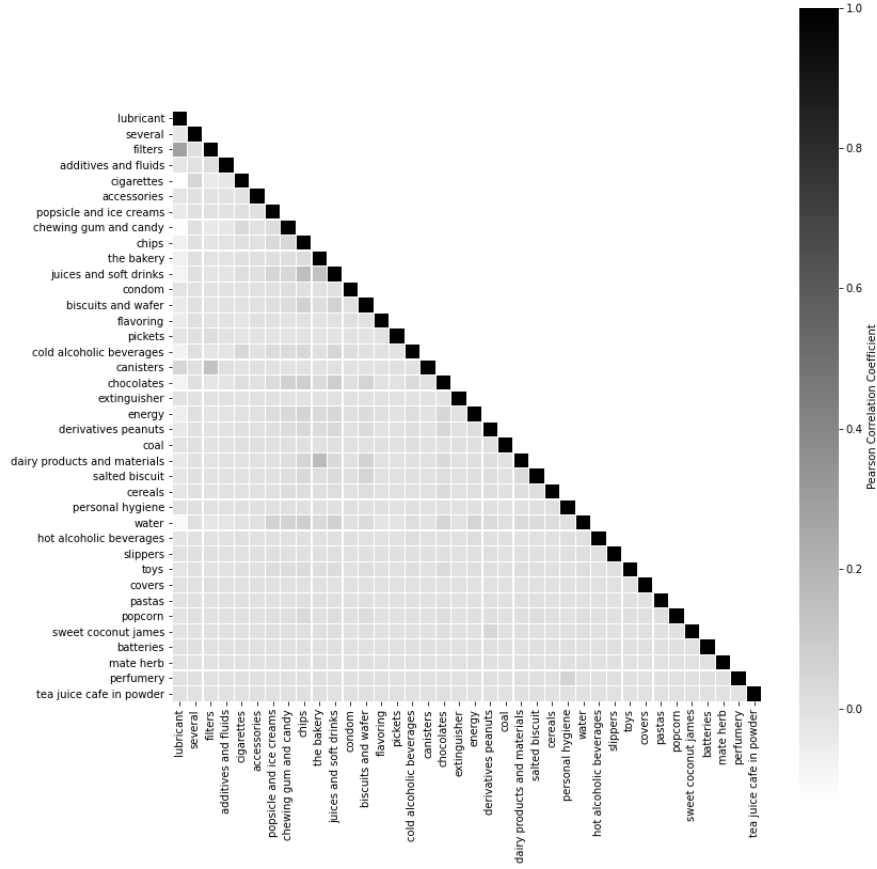


Figure 3.2: Correlation Matrix from Binary Purchase Vectors

A minimum spanning tree is a graph composed of the *shortest* path connecting all vertices, and therefore the values in the correlation matrix would need to be transformed via a function such that the higher the correlation value, the lower the output value. This would allow the MST to capture the strongest associations between the product categories. As used by **M. A. Valle et al.** in (**Valle et al. 2018**), the distance function (Equation 3.1) was used to make this transformation - where ϕ_{ij} refers to the correlation value for two given product categories i and j in Figure 3.2). However, whereas **M. A.**

Valle et al. left their ϕ_{ij} unaltered, we have converted it to an absolute value (i.e. $|\phi_{ij}|$). The reason behind this is to capture the strongest relationships between product categories, positive and negative alike. By leaving the value unaltered, the MST would not capture strong negative associations as they would have the highest weights after the distance function was applied. Figure 3.2 illustrates the effect of this function, where the stronger the correlation, the lower the distance function's output. The figure also demonstrates that because the Pearson's Correlation Coefficient is bound to the interval $[-1, 1]$, the distance function's output is therefore bound to the interval $[0, \sqrt{2}]$.

$$d_{ij} = \sqrt{2(1 - |\phi_{ij}|)} \quad (3.1)$$

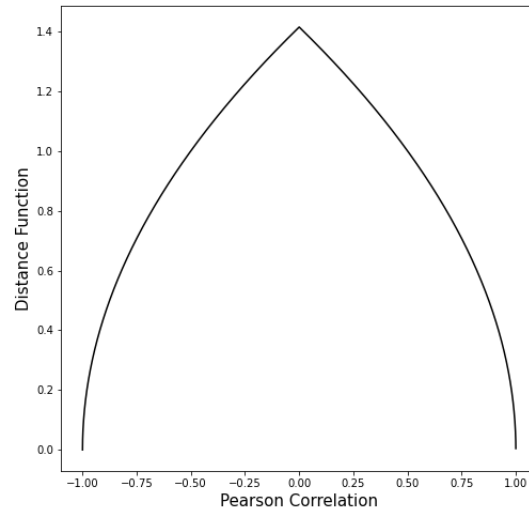


Figure 3.3: The effect of the distance function $\sqrt{2(1 - |x|)}$

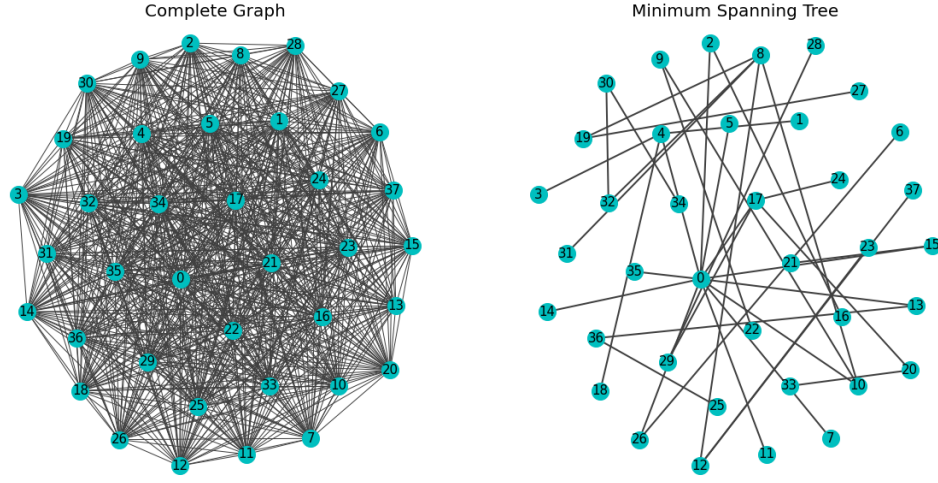


Figure 3.4: Product Category Graph and MST

With the correlation matrix transformed using Equation 3.1, a graph $G = (V, E)$ was generated such that the vertices V represent the product categories, and the edges E the associations between them. Employing Kruskal's algorithm, a minimum spanning tree was then extracted from this graph. Both the complete graph and the MST are illustrated in Figure 3.4. The value of each node is an integer, which corresponds with the index of the product category in the binary purchase vector dataset. The length of each edge is directly proportionate to its value, such that the greater the value, the greater the length of the edge.

3.3 Markov Clustering

The MST was then clustered using the Markov Clustering algorithm. To identify the most modular clustering configuration (modularity being the minimality of connectedness between dense clusters), an array of inflation scores between 1.5 and 2.5 (inclusive) were tested at increments of 0.1, leading us to determine that an inflation score of 1.6 resulted in the most optimal modularity. The Markov Clustering was performed using this inflation score, and the clustering results are illustrated in Figure 3.5 (note that while the disposition of the nodes is different from that illustrated in Figure 3.4, the values of all nodes and edges are the same).

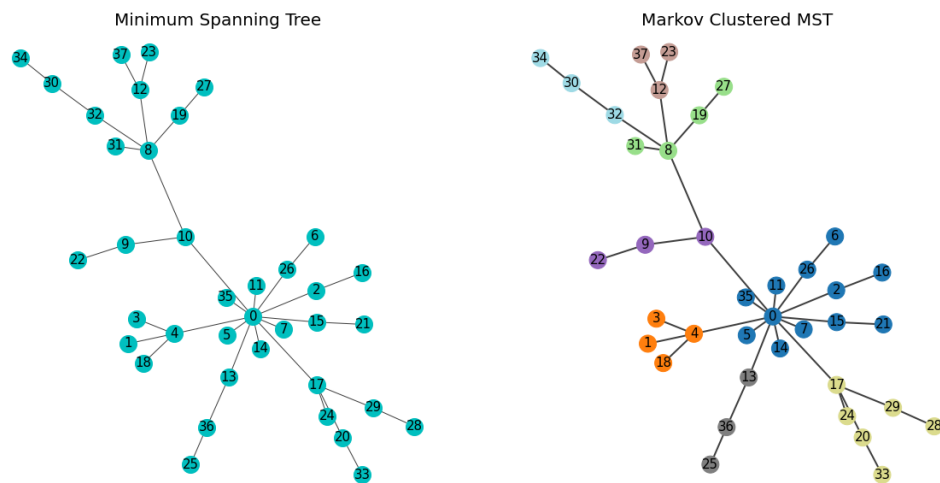


Figure 3.5: MST before and after Markov Clustering

The Markov Clustering algorithm segmented the nodes into 8 distinct clusters. Figure 3.6 illustrates the names of the product categories in each cluster, color-coded in accordance with the graphs in Figure 3.5.



Figure 3.6: Product categories by cluster

3.4 Rule Generation

3.4.1 Itemset Generation

For every cluster identified, all possible itemsets are generated for the n product categories in a given cluster such that the length of the itemsets range from 1 to $n - 1$. Listing 3.1 is the excerpt of code that generates all possible itemsets for a cluster.

```

1 # Source: arm_cython.pyx
2 from itertools import combinations
3 ...
4 cpdef __generate_itemsets_by_cluster():
5     global itemsets_by_cluster
6     cdef set items
7     for index, cluster in enumerate(clusters):
8         items = set()
9         for set_size in range(1, len(cluster)):
10             items.update(combinations(cluster, set_size))
11         itemsets_by_cluster[index] = items

```

Listing 3.1: Cluster Itemset Generation

Once the minimum spanning tree has been successfully clustered, association rules can be generated from the clusters in two distinct ways.

3.4.2 Bi-Cluster Rule Generation

Bi-cluster rules are those where the antecedent and consequent originate from distinct and separate clusters. All possible bi-cluster permutations nP_2 are calculated for n clusters. Possible association rules are then generated for these combinations of clusters.

```

1 # Source: arm_cython.pyx
2 cpdef list generate_biclust_rules(min_support=0.005, min_confidence=0.6):
3     cdef list rules = []
4     cdef list cluster_combinations = list(combinations(list(range(len(clusters))), 2))
5
6     for comb in cluster_combinations:
7         a_items = itemsets_by_cluster[comb[0]]
8         b_items = itemsets_by_cluster[comb[1]]
9         current_rules = list(product(a_items, b_items))

```

```

10     rules.extend(current_rules)
11
12     rules = __add_reverse(rules)
13     return __filter_rules(rules, min_support=min_support, min_confidence=min_confidence)

```

Listing 3.2: Cluster Itemset Generation

Listing 3.2 is the excerpt of code used to generate all the bi-cluster rules. On line 4, we have used the `combinations` function instead of `permutations`.

If we were to use permutations, we would have to generate each itemset i such that:

$$\{i_1 \in A\} \rightarrow \{i_2 \in B\}$$

for clusters A and B , and then later:

$$\{i_1 \in B\} \rightarrow \{i_2 \in B\}$$

which would essentially be the same combinations in a different configuration. Instead, by using `combinations` on line 4 of Listing 3.2, we only compute the former, and then re-add the rule to the ruleset with the antecedent and consequent swapped (line 12). This method essentially reduces our computational time by half.

3.4.3 Intra-Cluster Rule Generation

Intra-cluster rules are those where the both antecedent and consequent originate from the same cluster. Similar to the bi-cluster rules, the intra-cluster rules used the itemsets generated via the `__generate_itemsets_by_cluster()` function in Listing 3.1, however with the constraint:

$$\{i_1\} \rightarrow \{i_2\}; i \in A$$

for every itemset i in cluster A .

3.4.4 Rule Pruning

As described in Section ??, the *anti-monotone property of support* implies that the support for a subset of a given set should be equal to or greater than the support of the set:

$$\text{support}(\{x, y\}) \geq \text{support}(\{x, y, z\})$$

To take advantage of this, we first sort our rules in ascending order by the number of elements in the antecedent. When iterating through the potential rules, if either the antecedent or consequent is found to have a support score below the pre-defined minimum tolerance, the itemsets are added to a set `below_threshold`. Any future rules iterated are then cross-checked against `below_threshold`, and if it is found that any set in `below_threshold` is a subset of the rule, the rule is discarded as we know that its support is below our minimum tolerance. This allows us to avoid computing the support for several rules - the time complexity of which $O(n)$ at worst, thereby significantly reducing computation time. For both the *bi-cluster* and *intra-cluster* rule generation, the rules are pruned accordingly, and those that satisfy the thresholds specified are returned.

Chapter 4

Evaluation and Analysis

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