Clustering Item Data Sets with Association-Taxonomy Similarity

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Abstract

We explore in this paper the efficient clustering of item data. Different from those of the traditional data, the features of item data are known to be of high dimensionality and sparsity. In view of the features of item data, we devise in this paper a novel measurement, called the association-taxonomy similarity, and utilize this measurement to perform the clustering. With this association-taxonomy similarity measurement, we develop an efficient clustering algorithm, called algorithm AT (standing for Association-Taxonomy), for item data. Two validation indexes based on association and taxonomy properties are also devised to assess the quality of clustering for item data. As validated by both real and synthetic datasets, it is shown by our experimental results that algorithm AT devised in this paper significantly outperforms the prior works in the clustering quality, indicating the usefulness of association-taxonomy similarity in item data clustering.

1 Introduction

Data clustering is an important technique for exploratory data analysis. Data clustering is an application dependent issue and certain applications may call for their own specific requirements. Different from those of the traditional data, the features of market-basket data are known to be of high dimensionality and sparsity. There are several clustering technologies which addressed the issue of clustering market-basket data [2][3][4][5][6].

Explicitly, the support of item \( i \) is defined as the percentage of transactions which contain \( i \). Note that in mining association rules, a large item is basically an item with frequent occurrence in transactions. Thus, item \( i \) is called a large item if the support of item \( i \) is larger than the pre-given minimum support threshold. In market-basket data, the taxonomy of items defines the generalization relationships for the concepts in different abstraction levels.

In view of the features of item data, we devise in this paper a novel measurement, called the association-taxonomy similarity, and utilize this measurement to perform the clustering for shelf-space organization. With this association-taxonomy similarity measurement, we develop an efficient clustering algorithm, called algorithm AT (standing for Association-Taxonomy), for item data. Two validation indexes, association index (abbreviated as AI) and taxonomy index (abbreviated as TI), are also devised in this paper for clustering item data. As validated by real data, it is shown by our experimental results, with the association and taxonomy information, algorithm AT devised in this paper significantly outperforms the prior works [2][3] in the clustering quality.

2 Preliminaries

In market-basket data, a database of transactions is denoted by \( D = \{ t_1, t_2, ..., t_v \} \), where each transaction \( t_h \) is a set of items \( \{ i_1, i_2, ..., i_w \} \). In mining association rules [1], the minimum support \( Sup \) is given to identify the large itemsets. In addition, the support of an itemset in database \( D \) is defined as the number of transactions which contain this itemset in database \( D \). An itemset is called a large itemset if its support is larger than or equal to the minimum support \( Sup \). In this paper, an association itemset is defined as a large itemset that contains at least two items and is not contained by any other large itemset. The set of association itemsets is denoted by \( L_A = \{ I_1, I_2, ..., I_m \} \). Items in the transactions can be generalized to multiple concept levels of the taxonomy and represented as a taxonomy tree. In the taxonomy tree, the leaf nodes are called the item nodes and the internal nodes are called the category nodes.

In view of the features of item data, the items are categorized into three kinds of items which are association items (represented as \( I_A \)), single large items (represented as \( I_S \)),
and rare items (represented as $I_R$). An association item is an item which appears in at least one association itemset. A single large item is a large item but not an association item. In essence, a single large item can be viewed as a large 1-item which is not contained by any large 2-itemset. A rare item is not a large item (i.e., not frequently purchased). Explicitly, the rare item is an item whose support is smaller than the minimum support.

In this paper, a clustering $U = \{C_1, C_2, ..., C_k\}$ is a partition of items into $k$ clusters, where $C_j$ is a cluster consisting of a set of items. Note that purchasing relationships (i.e., association) and taxonomy relationships are important for the shelf-space organization. In this paper, the objective of clustering item data is to cluster the items with high association relationships and high taxonomy relationships together.

In view of the features of item data, we propose association index and taxonomy index, which are defined below, to assess the qualities of the clustering results.

**Definition 1: (Association Index)** The association index of the clustering $U$ is defined as:

$$AI(U) = \frac{\sum_{i_x, i_y \in C_p} A(i_x, i_y)}{|U|},$$

where $A(i_x, i_y)$ is the association value of item $i_x$ and item $i_y$. Explicitly, $A(i_x, i_y) = 1$, if $i_x$ and $i_y$ are in the same association itemset based on the minimum support $Sup$, and $A(i_x, i_y) = 0$, otherwise.

**Definition 2: (Taxonomy Index)** The taxonomy index of the clustering $U$ is defined as:

$$TI(U) = \frac{\sum_{C_p \in U} \frac{\sum_{i_x, i_y \in C_p} T(i_x, i_y)}{|C_p|(|C_p|-1)}}{|U|},$$

where $T(i_x, i_y)$ is the taxonomy value of item $i_x$ and item $i_y$. Explicitly, $T(i_x, i_y) = 1$, if $i_x$ and $i_y$ are in the same category under the cluster level $Lev^C$, and $T(i_x, i_y) = 0$, otherwise. In this paper, the cluster level $Lev^C$ is defined as the level where the number of categories is equal to the number of clusters $k$.

### 3 Design of Algorithm AT (Association Taxonomy)

In this paper, we devise algorithm AT for clustering item data. The similarity measurement of AT will be described in Section 3.1. Section 3.2 describes the procedure of AT.

#### 3.1 Similarity Measurement

The similarity measurement employed by algorithm AT is called association-taxonomy similarity which consists of the association similarity and the taxonomy similarity. As described before, the set of association itemsets is denoted by $L_A = \{I_1, I_2, ..., I_m\}$. For each association itemset, the association relationships of items can be represented as a complete graph $I_p = \{V_p, E_p\}$, consisting of a set of vertices $V_p$ and a set of edges $E_p$. In each complete graph, each vertex represents an item in the association itemset and each edge represents the association between two items. In mining association rules, an association rule $i_x \rightarrow i_y$ holds in transaction database $D$ with confidence $Con(i_x \rightarrow i_y)$ if $Con(i_x \rightarrow i_y)$ of transactions in $D$ that contain $i_x$ also contain $i_y$. In this paper, we use co-confidence as the measurement of the association between two items.

**Definition 3: (Co-Confidence between Association Items)** The co-confidence between $i_x$ and $i_y$ is defined as:

$$e(i_x, i_y) = \frac{1}{2}(Con(i_x \rightarrow i_y) + Con(i_y \rightarrow i_x))$$

$$= \frac{1}{2}\left(\frac{Sup(i_x \rightarrow i_y)}{Sup(i_x)} + \frac{Sup(i_y \rightarrow i_x)}{Sup(i_y)}\right)$$

where $Sup(i_x)$ is the support of item $i_x$. The co-confidence $e(i_x, i_y)$ represents the association between item $i_x$ and item $i_y$.

Each association itemset is viewed as a cluster of items (i.e., $I_p = I_p$). For notational simplicity, the union cluster of $C_p$ and $C_q$ is denoted as $C_{p,q}$. The set of overlapped items in $C_{p,q}$ is denoted as $C_{p,q}^{op}$ and the set of non-overlapped items in $C_{p,q}$ is denoted as $C_{p,q}^{nn}$. In addition, $E_{C_{p,q}}$ denotes the set of edges in $C_{p,q}$, $E_{C_{p,q}}^{op}$ denotes the set of edges connecting the overlapped items in $C_{p,q}$, $E_{C_{p,q}}^{nn}$ denotes the set of edges connecting the non-overlapped items in $C_{p,q}$.

**Definition 4: (Association Similarity between overlapped items)** The association similarity between overlapped items of $C_p$ and $C_q$ is defined as:

$$AS_{op}(C_p, C_q) = \sum_{i_x \in C_{p,q}^{op}, i_y \in C_{p,q}^{op}} e(i_x, i_y)$$

**Definition 5: (Association Similarity between overlapped items and non-overlapped items)** The association similarity between overlapped items and non-overlapped items of $C_p$ and $C_q$ is defined as:

$$AS_{on}(C_p, C_q) = \sum_{i_x \in C_{p,q}^{op}, i_y \in C_{p,q}^{nn}} e(i_x, i_y)$$

$$= \frac{\sum_{i_x \in C_{p,q}^{op}, i_y \in C_{p,q}^{nn}} e(i_x, i_y)}{|E_{C_{p,q}}^{op}| + |E_{C_{p,q}}^{nn}|}$$
For the similarity measurements in Definition 4 and Definition 5, $|E^m_{p,q}|$ is a normalization factor for considering the effect of the edges of non-overlapped items in decreasing the similarity between two clusters. Explicitly, the existence of non-overlapped items represents the dissimilarity between two clusters. Thus, an edge between the non-overlapped items increases the association dissimilarity between two clusters.

**Definition 6: (Association Similarity) The association similarity between $C_p$ and $C_q$ is defined as:**

$$AS(C_p, C_q) = \alpha_{oo} \cdot AS_{oo}(C_p, C_q) + \alpha_{on} \cdot AS_{on}(C_p, C_q),$$

where $\alpha_{oo}$ is the weight of the association similarity between overlapped items and $\alpha_{on}$ is the weight of the association similarity between overlapped items and non-overlapped items.

**Definition 7: (Taxonomy similarity of an overlapped Item) The taxonomy similarity of overlapped item $i_x$ to union cluster $C_{p,q}$ is defined as:**

$$T_o(i_x, C_{p,q}) = \frac{\sum_{k=1}^{N^{lev}} |C_{p,q}(i_x, k)|}{k},$$

where $N^{lev}$ is the number of levels in the taxonomy tree and $C_{p,q}(i_x, k)$ is the set of items which is in the same category with item $i_x$ in level $k$ in $C_{p,q}$.

**Definition 8: (Taxonomy Similarity of overlapped items) The taxonomy similarity of overlapped items of $C_p$ and $C_q$ is defined as:**

$$TS_o(C_p, C_q) = \sum_{i_x \in C_{p,q}} T_o(i_x, C_{p,q})$$

**Definition 9: (Taxonomy similarity of a non-overlapped Item) Let $i_y$ be an item in $C_p$ and $i_y$ is not overlapped with any item in $C_q$. The taxonomy similarity of non-overlapped item $i_y$ in cluster $C_p$ to cluster $C_q$ is defined as:**

$$T_n(i_y, C_q) = \frac{\sum_{k=1}^{N^{lev}} |C_q(i_y, k)|}{k},$$

where $C_q(i_y, k)$ is the set of items which is in the same category with item $i_y$ in level $k$ in $C_q$.

**Definition 10: (Taxonomy Similarity of non-overlapped items) The taxonomy similarity of non-overlapped items of $C_p$ and $C_q$ is defined as:**

$$TS_n(C_p, C_q) = \frac{\sum_{i_y \in C_p} T_n(i_y, C_q) + \sum_{i_x \in C_{p,q}} T_n(i_x, C_{p,q})}{|C_p| \cdot |C_q| + |C^p_{p,q}| \cdot |C_p|},$$

**Definition 11: (Taxonomy Similarity) The taxonomy similarity between $C_p$ and $C_q$ is defined as:**

$$TS(C_p, C_q) = \beta_o \cdot TS_o(C_p, C_q) + \beta_n \cdot TS_n(C_p, C_q) \cdot \frac{1}{N^{lev}},$$

where $\beta_o$ is the weight of the taxonomy similarity of overlapped items and $\beta_n$ is the weight of the taxonomy similarity of non-overlapped items. If each item in $C_p$ and each item in $C_q$ only have the root node as the same category, $C_p$ is totally dissimilar to $C_q$ according to the taxonomy tree and $TS(C_p, C_q)$ should be zero. Hence, because there are no overlapped item between $C_p$ and $C_q$, the constant $\frac{1}{N^{lev}}$ is subtracted in the non-overlapped part for normalization purpose.

**Definition 12: (Association-Taxonomy Similarity) The association-taxonomy similarity between $C_p$ and $C_q$ is denoted as $SIM(C_p, C_q)$ defined as:**

$$SIM(C_p, C_q) = \omega_A \cdot AS(C_p, C_q) + \omega_T \cdot TS(C_p, C_q),$$

where $\omega_A$ is the weight of the association similarity and $\omega_T$ is the weight of the taxonomy similarity. The determination of values of $\omega_A$ and $\omega_T$ is in fact application-dependent.

### 3.2 Procedure of Algorithm AT

Algorithm AT is designed to consist of three phases: the segmentation phase, the association-taxonomy phase, and the pure-taxonomy phase. Note that the association items consist of the elements in association itemsets. The overall procedure of algorithm AT is outlined as follows.

**Procedure of Algorithm AT (Association-Taxonomy)**

1. **(1) The Segmentation Phase:**
   - **Step 1.** Identify the set of association itemsets, the set of single large items, and the set of rare items.

2. **(2) The Association-Taxonomy Phase:**
   - **Step 2.** For each pair in the set of association itemsets, calculate the corresponding association-taxonomy similarity.
   - **Step 3.** Merge the pair which has the largest association-taxonomy similarity as a new cluster.
   - **Step 4.** Repeat Step 2 and Step 3 until the dendrogram is constructed.

3. **(3) The Pure-Taxonomy Phase:**
   - **Step 5.** Identify $k$ clusters in the dendrogram.
   - **Step 6.** For each single large item, allocate it to the cluster with the largest taxonomy similarity.
   - **Step 7.** For each rare item, allocate it to the cluster with the largest taxonomy similarity.
   - **Step 8.** Repeat Step 6 and Step 7 until no item is moved between clusters.
The advantageous features of algorithm AT are twofold. The first one is on employing the association-taxonomy similarity to effectively improve the quality of clustering association items. The second one is to allocate the single large items and rare items into clusters by calculating the taxonomy similarity. As such, these items can be efficiently and effectively allocated into the clusters. Note that the numbers of single large items and rare items are usually large as compared to the number of association itemsets. If we take each single large item (or each rare item) as a cluster and put them into the procedure from Step 2 to Step 4, the execution time will be prohibitive. In addition, lack of large association similarity with other clusters, these clusters with only one single large item (or one rare item) would never be merged until most of the association itemsets are merged. These problems are avoided in algorithm AT.

4 Experimental Studies

To assess the efficiency of AT, we conducted experiments to compare AT with the k-modes algorithm [3] and the ROCK algorithm [2]. We use the real market-basket data from a large bookstore company for performance study. In this real data set, there are $|D| = 100K$ transactions, $|I| = 58909$ items, and $N^{Lev} = 3$ levels. In addition, the number of the taxonomy level in this real data set is 3. In the real data, the items with the same category are usually purchased together. Thus, the association relationships and taxonomy relationships are related to each other.

Figure 1 shows the relative quality of clustering results of AT, ROCK, and k-modes in real data set where the database size $|D|$ varies from 20K to 100K. When we vary $|D|$ from 20K to 100K in ROCK, the numbers of clusters are, respectively, 576, 524, 468, 413, and 519. With association-taxonomy similarity measurement, AT significantly outperforms other algorithms as validated by $AI(U)$ in Figure 1(a) and by $TI(U)$ in Figure 1(b). In this real data set, because the items with high taxonomy relationships are usually purchased together while the items with low taxonomy relationships are not, AT has higher taxonomy index than association index, i.e., $AI(U) > AI(U)$.

5 Conclusion

In this paper, with the association-taxonomy similarity measurement proposed, we developed algorithm AT for item data. Two validation indexes based on association and taxonomy features of items was also devised in this paper to assess the quality of clustering for item data. As validated by real data, it was shown by our experimental results that algorithm AT devised in this paper significantly outperforms the prior works in the clustering quality of item data.

![Graphs showing Association Index and Taxonomy Index](image)

Figure 1. $AI(U)$ and $TI(U)$ for algorithms when $|D|$ varies.

References