Data preprocessing

- Data cleaning: fill missing values, smoothing of the data (binning method), identify outliers, resolve inconsistencies/contraddictions. To find outliers use clustering.
- Data integration: Put together multiple databases.
- Data transformation: normalize data, create new attributes.
- Data reduction: rapresent the data in a reduce format that produces similar analytical results. For example: remove unimportant attributes or cluster data and take a class rapresentative or remove randomly some data records. Simple random removal may have poor performance in the presence of skewed data. Stratified sampling: cluster data, find the percentage of the data in each class and sample using keeping in mind the percentage.
- Binning method, entropy based method. Binning method means partitioning data into bins and smooth each bin by taking the mean/mode/median/boundary...

Association rule mining Rule strength measure

$$\operatorname{support}(X \to Y) = \frac{\operatorname{count}(X \cup Y)}{n}$$
$$\operatorname{confidence}(X \to Y) = \frac{\operatorname{count}(X \cup Y)}{\operatorname{count}(X)}$$

1 Apriori frequent itemset algorithm

init-pass obtains the items in lexicographic order.

```
Algorithm Apriori(T)

C_1 \leftarrow \text{init-pass}(T);

F_1 \leftarrow \{f \mid f \in C_1, f.\text{count}/n \ge minsup\}; // n: no. of transactions in T for (k = 2; F_{k-1} \ne \emptyset; k++) do

C_k \leftarrow \text{candidate-gen}(F_{k-1}); for each transaction t \in T do

for each candidate c \in C_k do

if c is contained in t then

c.count++;

end

end

F_k \leftarrow \{c \in C_k \mid c.count/n \ge minsup\}

end

return F \leftarrow \bigcup_k F_k;
```

```
Function candidate-gen(F_{k-1})
    C_k \leftarrow \emptyset;
    forall f_1, f_2 \in F_{k-1}
           with f_1 = \{i_1, \ldots, i_{k-2}, i_{k-1}\}
            and f_2 = \{i_1, \ldots, i_{k-2}, i'_{k-1}\}
            and i_{k-1} < i'_{k-1} do
                                                           // join f_1 and f_2
        c \leftarrow \{i_1, ..., i_{k-1}, i'_{k-1}\};
        C_k \leftarrow C_k \cup \{c\};
        for each (k-1)-subset s of c do
            if (s \notin F_{k-1}) then
                delete c from C_k;
                                                           // prune
        end
    end
    return C_k;
```

$\mathbf{2}$ MS Apriori

sort in ascending

```
\begin{aligned} & \text{dgorithm MSapiori(I, MS, \varphi)} & \text{$////\varphi$ is for support difference constraint} \\ & M \leftarrow \text{sort}(l, MS); \\ & L \leftarrow \text{init}\text{pass}(M, T); \\ & F_{k} \leftarrow \{\{b \mid l \in L, Loount h \geq \text{MIS}(l)\}; \\ & \text{for } (k = 2; F_{k}, \pm \varnothing; k++) \text{ do} \\ & \text{if } k = 2 \text{ then} \\ & C_{k} \leftarrow \text{level}2\text{-candidate-gen}(L, \varphi) \\ & \text{else } C_{k} \leftarrow \text{MScandidate-gen}(F_{k+1}, \varphi); \\ & \text{end}; \end{aligned}
                                                  nd;

r each transaction t ∈ T do
for each candidate c ∈ C<sub>c</sub> do

if c is contained in t then
c.count++;

if c − {c[1]} is contained in t then
c.tailCount++
                                F_k \leftarrow \{c \in C_k \mid c.count/n \ge MIS(c[1])\}
         end
return F \leftarrow \bigcup_k F_k;
```

Figure 1: MsApriori

data set T and the sorted items M, to produce the seeds L for generating candidate itemsets of length 2, i.e., C_2 : init-pass() has two steps:

- 2. It then follows the sorted order to find the first item *i* in *M* that meets MIS(*j*). *i* is inserted into *L*. For each subsequent item *j* in *M* after *i*, if *j*-countin *j* e MIS(*j*), then *j* is also inserted into *L*. The subsequent item *j* in *M* after *i*, if scountin *j* e MIS(*j*), then *j* is also inserted into *L*, where *j*-count is the support count of *j*, and *n* is the total number of transactions in *T*.

Figure 2: Init Pass

Function level2-candidate-gen(L, φ) // initialize the set of candidates $C_2 \leftarrow \emptyset$; // initial for each item l in L in the same order do (at least the first of the same state of the first of th Fig. 2.7. The level2-candidate-gen function Function MScandidate-gen (F_{k-1}, φ) // initialize the set of candidates // find all pairs of frequent itemsets // that differ only in the last item Fig. 2.8. The MScandidate-gen function

Figure 3: Candidate Gen MS

GSP mining algorithm

Very similar to the Apriori algorithm

```
Algorithm GSP(S)
     C_1 \leftarrow \text{init-pass}(S);
                                                            // the first pass over S
     F_1 \leftarrow \{\langle \{f\} \rangle | f \in C_1, f.\text{count}/n \ge minsup\}; // n \text{ is the number of sequences in } S
     for (k = 2; F_{k-1} \neq \emptyset; k++) do
                                                            // subsequent passes over S
4
        C_k \leftarrow \text{candidate-gen-SPM}(F_{k-1});
5
        for each data sequence s \in S do
                                                            // scan the data once
             for each candidate c \in C_k do
7
                 if c is contained in s then
8
                                                            // increment the support count
                    c.count++;
9
            end
10
        F_k \leftarrow \{c \in C_k \mid c.count/n \ge minsup\}
11
12 end
13 return \bigcup_k F_k;
```

Fig. 12. The GSP Algorithm for generating sequential patterns

When he says is the same as the subsequence... Means that you have to remove the parenthesis and see if the elements are in the same order.

Function candidate-gen-SPM(F_{k-1})

- 1. Join step. Candidate sequences are generated by joining F_{k-1} with F_{k-1}. A sequence s₁ joins with s₂ if the subsequence obtained by dropping the first item of s₁ is the same as the subsequence obtained by dropping the last item of s₂. The candidate sequence generated by joining s₁ with s₂ is the sequence s₁ extended with the last item in s₂. There are two cases:
 - the added item forms a separate element if it was a separate element in s₂, and is appended at the end of s₁ in the merged sequence, and
 - the added item is part of the last element of s₁ in the merged sequence otherwise.

When joining F_1 with F_1 , we need to add the item in s_2 both as part of an itemset and as a separate element. That is, joining $\langle \{x\} \rangle$ with $\langle \{y\} \rangle$ gives us both $\langle \{x,y\} \rangle$ and $\langle \{x\} \{y\} \rangle$. Note that x and y in $\{x,y\}$ are ordered.

 Prune step. A candidate sequence is pruned if any one of its (k-1)subsequence is infrequent (without minimum support).

Fig. 13. The candidate-gen-SPM() function

4 Index for Classifiers

Metric	Description / Formula
Accuracy	Correctly Classified Examples Total Examples
Precision	$\frac{TP}{TP+FP}$: Of predicted positives, how many are correct.
Recall/sensitivity (TPR)	$\frac{TP}{TP+FN}$: Of actual positives, how many are identified.
F_1 Score	$2 \times \frac{p \cdot r}{p+r}$: Combines precision and recall.
Specificity (TNR)	$\frac{TN}{TN+FP}$: True negative rate.
False Positive Rate (FPR)	1-TNR
ROC Curve	Plot: FPR (x-axis) vs. TPR (y-axis).

Table 1: Summary of Classification Metrics

5 Decision trees

```
. Algorithm decisionTree(D, A, T)
1
      if D contains only training examples of the same class c_i \in C then
2
          make T a leaf node labeled with class c_j;
3
      elseif A = \emptyset then
4
          make T a leaf node labeled with c_f, which is the most frequent class in D
5
      else // D contains examples belonging to a mixture of classes. We select a single
            // attribute to partition D into subsets so that each subset is purer
7
           p_0 = impurityEval-1(D);
8
           for each attribute A_i \in \{A_1, A_2, ..., A_k\} do
9
               p_i = \text{impurityEval-2}(A_i, D)
10
           Select A_g \in \{A_1, A_2, ..., A_k\} that gives the biggest impurity reduction,
11
               computed using p_{\theta} - p_i;
12
           if p_{\theta} - p_{\varrho} \le threshold then
                                            //A_g does not significantly reduce impurity p_\theta
13
              make T a leaf node labeled with c_j, the most frequent class in D.
14
                                             // A_g is able to reduce impurity p_\theta
15
               Make T a decision node on A_g;
               Let the possible values of A_g be v_1, v_2, ..., v_m. Partition D into m
16
                   disjoint subsets D_1, D_2, ..., D_m based on the m values of A_g.
               for each D_i in \{D_1, D_2, ..., D_m\} do
17
                   if D_j \neq \emptyset then
18
19
                      create a branch (edge) node T_i for v_i as a child node of T;
20
                      decisionTree(D_i, A-\{A_g\}, T_i)//A_g is removed
21
22
               end
23
           end
24
      end
```

Entropy D is the dataset. C are the different classes in the dataset.

$$\mathrm{entropy}(D) = -\sum_{j=1}^{|C|} \Pr(C_j) \log_2 \Pr(C_j)$$

Is a positive value. The sum of the probabilities of the classes is 1. The more the data is purer, the more the entropy value is close to zero. Worst entropy is 1 (all the classes are equally distributed).

$$\operatorname{entropy}_{A_i}(D) = \sum_{j=1}^v \frac{|D_j|}{|D|} \times \operatorname{entropy}(D_j)$$

This value is the entropy if we choose to partition the data over attribute A. There are v possible values for the attribute A.

$$gain(D, A_i) = entropy(D) - entropy_{A_i}(D)$$

The higher the gain the better

6 Naive based classification

Product rule

$$Pr(a1, a2) = Pr(a1)Pr(a2|a1)$$

Product rule for conditional probability

$$Pr(a1, a2|c) = Pr(a2|c)Pr(a2|a1, c)$$

General case

$$\Pr(a_1 \cdots a_n \mid c) = \Pr(a_1 \mid c) \Pr(a_2 \mid c, a_1) \cdots \Pr(a_n \mid c, a_1 \cdots a_{n-1})$$

Conditional independence assumption

$$Pr(a_i \mid c, a_1 \cdots a_{i-1}) = Pr(a_i \mid c)$$

Goal:

$$\Pr(c \mid (a_1 \cdots a_n)) = \frac{\Pr(c)\Pr((a_1 \cdots a_n) \mid c)}{\Pr(a_1 \cdots a_n)}$$

Using the law of total probability

$$\Pr(a_1 \cdots a_n) = \sum_{r=1}^{|C|} \Pr(c) \Pr((a_1 \cdots a_n) \mid c)$$

Using the product rule and the conditional independence assumption

$$\Pr(c \mid (a_1 \cdots a_n)) = \frac{\Pr(c) \prod_{i=1}^n \Pr(a_i \mid c)}{\sum_{r=1}^{|C|} \Pr(c) \prod_{i=1}^n \Pr(a_i \mid c)}$$

Adjusting the probability to account for attribute values that don't occur with that class.

$$\Pr(A_i = a_i \mid C = c_j) = \frac{n_{ij} + \lambda}{n_j + \lambda m_i}$$
Instead if there are missing values we just ignore them and use what we have

- □ n_i: # examples with C=c_i in training data
- $\neg n_{ii}$: # examples with both $A_i = a_i$ and $C = c_i$
- m_i : # possible values of attribute A_i .
- \square Normally, we use λ = 1

Naive based text classification 7

$$\Pr(c|d) = \frac{\Pr(c)\Pr(d|c)}{\Pr(d)}$$

$$= \frac{\Pr(c) \prod_{k=1}^{\text{all word in d}} \Pr(w_k|c)}{\sum_{r=1}^{|C|} \Pr(c_r) \prod_{k=1}^{\text{all word in d}} \Pr(w_k|c)}$$

$$\Pr(c) = \frac{\sum_{i=1}^{|D|} \Pr(c|d_i)}{|D|}$$

$$\Pr(w|c) = \frac{\sum_{s=1}^{|D|} N_i \Pr(c|d_i)}{\sum_{s=1}^{|V|} \sum_{i=1}^{|D|} N_{si} \Pr(c|d_i)}$$