Course Information

Course: CS 570 Data Mining **Instructor:** Dr. Wei Jin **University:** Emory University

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1 Preprocessing

Data integration Redundancy

- 1. Chi-Square (χ^2) Test (Categori**cal**): $\chi^2 = \sum_{i}^{n} \frac{(O_i - E_i)^2}{E_i}$ H_0 : Variable distributions are independent. Higher χ^2 suggests correlation (not causality). df = (#row - 1)(#col - 1)
- 2. Variance & Covariance (Numeri**cal**): $\hat{\sigma}^2 = \frac{1}{n} \sum (x_i - \hat{\mu})^2$, $\hat{\sigma}_{12} = \frac{1}{n} \sum (x_{i1} - \hat{\mu}_1)(x_{i2} - \hat{\mu}_2)$ Expectation form: $\sigma_{12} = E[X_1 X_2] E[X_1]E[X_2]$
- 3. Covariance matrix $\sigma_{12} = E[(X_1 X_2)]$ $\mu_1)(X_2 - \mu_2)^T$
- 4. **Pearson Correlation**: Normalized ar relation only. $\hat{\rho}_{12} = \frac{\hat{\sigma}_{12}}{\hat{\sigma}_1 \hat{\sigma}_2} =$ $\frac{\sum (x_{i1} - \hat{\mu}_1)(x_{i2} - \hat{\mu}_2)}{\sqrt{\sum (x_{i1} - \hat{\mu}_1)^2 \sum (x_{i2} - \hat{\mu}_2)^2}}$

1.1 Data Reduction

- Parametric: Regression
- Non-parametric: Histogram, Clustering, Sampling

1.2 Data Transformation

Min-Max: $v' = \frac{v - min_A}{max_A - min_A} (new_max - max)$ new_min) + new min **Z-Score:** $v' = \frac{v - \mu_A}{\sigma_A}$

Decimal Scaling: $v' = \frac{v}{10^j}$, where *j* is the smallest integer s.t. max(|v'|) < 1

2 Dimensionality Reduction **Curse of Dimensionality**

Data becomes sparse, distance/density lose meaning, subspace combinations grow exponentially: $1 - (1 - \epsilon)^d$

PCA (Principal Component Analysis)

Unsupervised, linear. Maximizes variance preservation: $L = g^T S g - \lambda (g^T g - 1)$

Sample covariance matrix: S = $\frac{1}{N-1}\sum_{i=1}^{N}(x_i-\overline{x})(x_i-\overline{x})^T$

Find first p eigenvectors $\{g_i\}_{i=1}^p$ from: $|S - \lambda I| = 0$, $Sg = \lambda g$

Construct transformation matrix G = $[g_1, g_2, ..., g_p]$ and transform: $x \in \mathbb{R}^d \to$ $G^T x \in \mathbb{R}^p$

Attribute Subset Selection

Choose best attribute by significance tests under independence assumption. Selection or Elimination.

3 Pattern Mining

3.1 Concept

Support (Count): Freq. of itemset *X* in transactions. Relative Support: Fraction of transactions containing X. An itemset X is *frequent* if X's support is no less than a minimum support threshold.

Frequent Itemset: *X* is frequent if $Supp(X) \ge min support.$ Association Ru-Covariance, range: [-1,1]. Line- le: $X \to Y$, where X, Y are itemsets.

Supp
$$(X \to Y) = \frac{|T(X \cup Y)|}{|T|}$$

Conf $(X \to Y) = \frac{|T(X \cup Y)|}{|T(X)|}$

3.2 Apriori

BFS, iterative DB scans.

Downward Closure: Any subset of a frequent itemset must be frequent. So we can prune supersets of itemset found infrequent. $\forall X, Y : (X \subseteq Y) \Rightarrow supp(X) \ge$

Candidate generation: a. $F_{k-1} \times F_1$ b.

Rule generation: low confidence rule $conf(ABC \rightarrow D) \ge conf(AB \rightarrow CD) \ge$ $conf(A \rightarrow BCD)$

3.3 FPGrowth

DFS, 2 scans. Grow long patterns from short ones using **local** frequent items. 1. FPTree Construction: Find frequent 1itemsets, sort into f-list, insert transactions into FP-Tree maintaining freq. counts. 2. **FPTree Mining**: Construct conditional FP-Tree for every suffix until only a single path remains.

3.4 Evaluation

$$Lift(X \to Y) = \frac{p(X \cup Y)}{p(X)p(Y)} \text{ Lift}(X \to Y) \text{ vs.}$$

$$Lift(X \to \neg Y)$$

Closed Pattern: *X* is frequent and has no super-pattern $Y \supset X$ with the same support. **Max-Pattern**: *X* is frequent and has no frequent super-pattern $\dot{Y} \supset X$.

3.5 Sequential Pattern Mining

Sequence Data: sequence \rightarrow element $(transaction) \rightarrow event (item). k-sequence:$ A sequence with k events/items. **Support of subsequence** *w*: Fraction of data sequences that contain w. Generalized Sequential Pattern (GSP) Algorithm ~ Apriori: Merging k-1 patterns to form k-sequences: - Merge two frequent (k-1)sequences w_1, w_2 if the first event of w_1 matches the last event of w_2 after deletion. - The resulting *k*-sequence extends w_1 : - If the last two events in w_2 share an element, append its last event to w_1 's last element. - Else, append it as a new element.

4 Similarity Search

4.1 Proximity measure

Minkowski Distance: $\left(\sum_{k=1}^{n} |p_k - q_k|^r\right)^{\overline{r}} - r = 1 \quad (Manhat$ tan, L_1 norm), r = 2 (Euclidean), $r \to \infty$ $(L_{\text{max}} \text{ norm}).$

Mahalanobis Distance: Penalizes covariance matrix.

SMC vs. Jaccard:
$$SMC = \frac{M_{11}+M_{00}}{M_{11}+M_{00}+M_{10}+M_{01}}$$
, $J = \frac{M_{11}}{M_{11}+M_{10}+M_{01}}$ - One-hot encoding \rightarrow Jaccard. - Cosine Similarity. - Correlation: standar-

- Cosine Similarity. - Correlation: standar $dize \rightarrow dot product (fails for nonlinear).$

4.2 Similarity Search

Collaborative Filtering:

User-based NN: $pred(u,i) = \overline{r}_u +$ $\sum_{n \in neighbors(u)} sim(u,n) \cdot (r_{ni} - \overline{r}_n)$ $\sum_{n \in neighbors(u)} sim(u,n)$

Item-based NN:

 $\sum_{j \in ratedItems(u)} sim(i,j) \cdot r_{uj}$ $\sum_{i \in ratedItems(u)} sim(i,j)$

k-D Trees: Partition data recursively: 1. Choose dim. w/ max variance. 2. Split at median. 3. Partition into two halves. 4. Repeat for next highest variance.

4.3 Hashing for Dim Reduction Locality Sensitive Hashing (LSH) Similar

items ⇒ High probability of same bucket MinHash: Increases the probability of collision when documents are similar. Generate a random permutation of the columns. For data d_i , the MinHash value is the **first permuted column** that has a "1". \Rightarrow Reduces the number of pairwise comparisons.

5 Classification

5.1 Decision Tree

Induction Algorithm (overfitting → preor post-pruning), Prediction: Majority

Entropy (Expected information needed to classify a tuple in D): Info(D) = $-\sum_{i=1}^{m} p_i \log_2(p_i)$ Information needed after splitting D using attribute A: $Info_A(D) = \sum_{j=1}^{v} \frac{|D_j|}{|D|} \times Info(D_j)$ Infor-

mation Gain (reduction in entropy): $Gain(A) = Info(D) - Info_A(D)$ **Stopping Conditions**: All nodes belong

to same class, no attributes left, or no samples left.

5.2 Bayesian classification

Generative classifier P(Y,X), incremental (incorporate prior knowledge) Naïve Assumption: Attributes are conditionally independent

Categorical: $p(X|C_i) = \prod_k p(x_k|C_i)$ Continuous-valued $p(x_k = v_k|C_i) =$

$$N(x_k|\mu_{C_i}, \sigma_{C_i}) = \frac{1}{\sqrt{2\pi\sigma_{C_i}^2}} e^{\frac{-(x-\mu_{C_i})^2}{2\sigma^2}}$$

5.3 Linear Classifier

Discriminative Classifier P(Y|X)

Logistic Regression Turns linear predictions into probabilities.

Sigmoid function: $S(x) = \frac{1}{1+e^{-x}} =$ $\frac{e^x}{e^x+1}$ Likelihood Function: l(w) =

$$\sum_{i=1}^{N} y_i \log p(Y = 1|x_i; w) + (1 - y_i) \log(1 - p(Y = 1|x_i; w))$$
• **Optimization**: No closed-form solution \rightarrow Use gradient descent $a_{n+1} = a_n - \gamma \nabla F(a_n)$.

kNN: Non-parametric, lazy learner.

5.4 Bayesian belief networks

Chain Rule of Probability: Prob. factorization based on Directed Acyclic Graph (DAG). $P(x_1, x_2, ..., x_N) =$ $\prod_i P(x_i|Parent(x_i))$

Maximize margin $\frac{2}{||w||} \rightarrow \text{Minimize}$ $E(\mathbf{w}) = \frac{\|\mathbf{w}\|^2}{2}$ Subject to:

$$v_i(\mathbf{w}^\top \mathbf{x}_i + b) \ge 1, \quad i = 1, 2, \dots, N$$

Lagrange Formulation

$$L(\mathbf{w}) = \frac{\|\mathbf{w}\|^2}{2} - \sum_{i} \lambda_i [y_i(\mathbf{w}^{\top} \mathbf{x}_i + b) - 1]$$

Derivatives:

$$\frac{\partial L}{\partial \mathbf{w}} = 0 \Rightarrow \mathbf{w} = \sum_{i} \lambda_{i} y_{i} \mathbf{x}_{i}, \quad \frac{\partial L}{\partial b} = 0 \Rightarrow \sum_{i} \lambda_{i} y_{i} = 0$$
Constraints:

$$\lambda_i \ge 0$$
, $\lambda_i[y_i(\mathbf{w}^\top \mathbf{x}_i + b) - 1] = 0$

(Complementary slackness) **Quadratic Programming:**

$$\max_{\lambda} L(\lambda) = \sum_{i} \lambda_{i} - \frac{1}{2} \sum_{i,j} \lambda_{i} \lambda_{j} y_{i} y_{j}(\mathbf{x}_{i} | \mathbf{x}_{j}), \quad \lambda \geq 0$$

Support Vectors: Points where $\lambda_i \neq 0$. **Kernel Trick**

Gain linear separation by mapping to a higher dimension: **Applying Kernel:**

$$L_d = \sum \alpha_i - \frac{1}{2} \sum \alpha_i \alpha_j y_i y_j K(\mathbf{x}_i, \mathbf{x}_j)$$

1 Clustering

Unsupervised learning.

1.1 Partitioning Approach

K-means: Each cluster is represented by the mean of its points.

Algorithm: Select *K* initial centroids. **Repeat:** a. Assign each point to the closest centroid. b. Update centroids as cluster means. **Until** centroids don't change.

Complexity: O(nKId) (n: points, K: clusters, I: iterations, d: dimensions) **P1. initialization of centroids** k-means++: For each point x_i , compute its distance to the nearest centroid: $d(x_i,C)^2 = \min_{x_c \in C} d(x_i,x_c)^2$ Choose the next centroid x_k with probability: $P(x_k) = \frac{d(x_k,C)^2}{\sum_i d(x_i,C)^2}$

P2. Choosing K: Cross-validation. P3. Non-globular clusters \rightarrow Use many

clusters. P4. Sensitive to outliers \rightarrow Use K-

P4. Sensitive to outliers → Use K-Medoids (medoid instead of mean).
P5. Empty clusters → Select the point contributing most to SSE as a new centroid.

*Bisecting K-Means (Hierarchical) Split the cluster using basic K-means. → Add resulting clusters with the highest SSE.

1.2 Hierarchical Approach

Irreversible, $O(N^2)$ space, $O(N^3)$ time. **Divisive** (**Top-Down**): Better global structure, early stopping avoids overclustering. (*Can use K-means for efficiency*) **Agglomerative** (**Bottom-Up**): Clear merging patterns, useful for structured data. **Inter-Cluster Similarity**:

MIN (Single-Linkage): Handles nonelliptical shapes, sensitive to noise/outliers.

MAX (Complete-Linkage): Less noisesensitive, splits large clusters, favors globular clusters.

Group Average: Reduces noise impact, favors globular clusters.

Centroid-Based Distance: Noise-robust, simple, not suitable for categorical data.

1.3 Density-Based Approach

Clusters based on density reachability, detecting outliers and non-globular shapes. **DBSCAN:** Label points as **core**, **border**, or **noise**. Remove noise, connect core

points within ϵ , assign border points to nearest core cluster. **Parameters:** Eps (radius), **MinPts** (min neighbors). **Properties:** Noise-robust, handles arbitrary shapes, sensitive to Eps/MinPts. Sort points by k-th nearest neighbor distance, use elbow method for optimal ϵ .

1.4 Clustering Evaluation

Extrinsic (Supervised) BCubed Precision & Recall measure correctness of object relations: Precision(o_i) = $\frac{\text{#objects in } C(o_i) \text{ with } L(o_i)-1}{\text{#objects in } C(o_i) \text{ with } L(o_i)-1}$, Recall(o_i) = $\frac{\text{#objects in } C(o_i) \text{ with } L(o_i)-1}{\text{Total # objects with } L(o_i)-1}$. BCubed Pre-

cision and Recall: $\frac{1}{n}\sum_{i=1}^{n}\operatorname{Precision}(o_i)$, $\frac{1}{n}\sum_{i=1}^{n}\operatorname{Recall}(o_i)$.

Intrinsic (Unsupervised) Com-

pactness (Intra-cluster Distance): $a(o) = \frac{\sum_{o' \in C_i, o \neq o'} dist(o, o')}{|C_i| - 1}$, measures how

 $a(o) = \frac{}{|C_i|-1}$, measures now close o is to other points in the same cluster. Lower is better.

Separateness (Inter-cluster Distance):

$$b(o) = \min_{C_j: j \neq i} \frac{\sum_{o' \in C_j} dist(o,o')}{|C_j|}, \text{ measu-}$$

res how far o is from the closest other cluster. Higher is better.

Silhouette Score: $s(o) = \frac{b(o) - a(o)}{\max(a(o), b(o))} \in [-1, 1]$. $s(o) \approx 1$: Well-clustered, s(o) < 0: Incorrect clustering.

2 Outlier Detection

Types: Global, contextual (attribute-based), collective.

2.1 Statistical Methods

Parametric: Probability density function, Grubb's test.

Univariate: Fit normal distribution, outliers deviate by $\geq 3\sigma$. Grubb's Test: z =

$$\frac{|x-\mu|}{\sigma}, \quad z \ge \frac{N-1}{\sqrt{N}} \times \frac{t_{\alpha/(2N),N-2}}{\sqrt{N-2+t_{\alpha/(2N),N-2}^2}}$$

Multivariate: Mahalanobis Distance accounts for covariance: $MDist(o, \delta) = \sqrt{(o - \delta)^{\top} \Lambda (o - \delta)}$

2.2 Non-Parametric Methods

Histogram and Kernel Density Estimation (KDE) Gaussian kernel smoo-

thing:
$$K(u) = \frac{1}{h\sqrt{2\pi}}e^{-\frac{1}{2}(\frac{u}{h})^2}$$
, Density estimation for dataset $\{x_1, x_2, ..., x_N\}$: $\hat{f}(o_i) =$

$$\sum_{j=1}^{N} K\left(\frac{n-kj}{h}\right)$$

2.3 Proximity-based Methods

Distance-based: An object is an outlier if $\leq k$ objects exist within its r-neighborhood. $k = \pi \cdot |D|$. **Density-Based:** k-distance $\mathrm{dist}_k(o)$: Distance from o to its k-th nearest neighbor. $N_k(o)$: Set of objects within k-distance. **Local Outlier Factor (LOF)** Density:

$$\begin{split} d(o) &= \frac{|N_k(o)|}{\sum_{o' \in N_k(o)} \max(\operatorname{dist}(o,o'),\operatorname{dist}_k(o'))} \text{ LOF:} \\ LOF(o) &= \frac{1}{|N_k(o)|} \sum_{o' \in N_k(o)} \frac{d(o')}{d(o)} \text{ LOF} \approx 1: \end{split}$$

Not an outlier. LOF
$$\gg$$
 1: Local outlier. **2.4 Clustering-based Methods**

DBSCAN: Detects objects not belonging to any cluster. **Distance-based**

Outliers: Score $\propto \frac{\operatorname{dist}(o,c_n)}{\operatorname{avg dist}(c_n)}$. CBLOF

(Cluster-Based Local Outlier Factor): Sort clusters by size. CBLOF: $|C| \times \text{similarity}(p, C_{\text{closest}})$. Example: If p in small cluster C_3 has low similarity to its closest large cluster C_2 , then p is an outlier.