Collective Entity Resolution in Relational Data

CompSci 590.03 Instructor: Ashwin Machanavajjhala



Recap: Constraints

• Transitivity:

If x and y match, y and z match, then x and z must match

Useful in deduplication

• Exclusivity:

If x matches with y, then z cannot match with y

- Useful in record linkage (matches across two datasets)
- Each dataset does not have any duplicates.

Relational Constraints:

If x and y match, then z and w should match

- If movies are the same, then directors must be the same
- (We will see in next class)



Recap: Constraint Types

	Hard Constraint	Soft Constraint
Positive Evidence	Transitivity: x=y & y=z => x=z	Relational: If x, y match then z, w are more likely to match <i>If two venues match, then their</i> <i>papers are more likely to match</i>
Negative Evidence	Exclusivity: x and y must refer to distinct entities Relational: If x,y don't match then z,w cannot match	Soft Exclusivity: x and y are very likely different elements
	<i>If two venues don't match, then their papers don't match</i>	



Match Dependencies

When matching decisions depend on other matching decisions (in other words, matching decisions are not made independently for each pair), we refer to the approach as *collective*



This Class

- Collective Entity Resolution for Relational Data
 - Problem Statement
 - Motivating Example
 - Similarity functions for Linked Data
 - Relational Clustering



Abstract Problem Statement



Deduplication-Problem Statement



Relationships are crucial



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InfoVis Co-Author Network Fragment



before

Relational Constraints



Very similar names.

Added evidence from shared co-authors

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Relational Constraints



Relational Constraints



Collective Entity Resolution



One resolution provides evidence for another => joint resolution



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Relational Features

- There are a variety of ways of improving ER performance when data is richer than a single table/entity type
- One of the simplest is to use additional information, to *enrich* model with *relational features* that will provide richer context for matching



Examples of relational features

- Value of edge or neighboring attribute (1-1)
- Aggregates (1-many)
 - Mode (sum, min, max) of related attribute
- Set similarity measures to compare nodes based on set of related nodes, e.g., compare neighborhoods
 - Overlap
 - Jaccard coefficient
 - Average similarity between set members



Preferential Attachment Score

[Liben-Nowell & Kleinberg, JASIST07]

• Based on studies, e.g. [Newman, PRL01], showing that people with a larger number of existing relations are more likely to initiate new ones.

 $s(a,b) = |N_a| \cdot |N_b|$ Set of a's neighbors



Common Neighbors

• Two nodes are likely to be connected in a graph if they share a large number of common neighbors.

 $s(a,b) = N(a) \cap N(b)$

Can be any kind of shared attributes or relationships to shared entities



Adamic/Adar Measure

[Adamic & Adar, SN03]

• Two nodes are more similar if they share more items that are overall less frequent



20

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Katz Score

• Two objects are similar if they are connected by shorter paths

$$s(a,b) = \sum_{l=1}^{\infty} \beta^{l} \cdot |\operatorname{paths}^{\langle l \rangle}(a,b)|$$

Set of paths between
a and b of length exactly ℓ

Decay factor between 0 and 1

• Since expensive to compute, often use approximate Katz, assuming some max path length of k

21

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Personalized Page Rank

- Stationary distribution of a random walk:
 - With probability (1-c), follow a random outgoing edge
 - With probability c, jump to the target node 'a'



SimRank

[Jeh & Widom, KDD02]

23

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- "Two objects are similar if they are related to similar objects"
- Defined as the unique solution to:



- Computed by iterating to convergence
- Initialization to s(a, b) = 1 if a=b and 0 otherwise

- sim(a,b) measures how soon two (reverse) random walks starting from a and b meet at the same node.
- Works best for bipartite graphs (having two types of entities)



Expected Distance

$$d(u,v) = \sum_{t:u \rightsquigarrow v} P[t]l(t)$$

$$- d(u,v) = 0$$
, if $u = v$

- t: tour (path with cycles) starting at u and ending at v
- t = [w1, w2, ..., wk]

$$P[t] = \prod_{i=1}^{k-1} \frac{1}{|O(w_i)|}$$



Expected Meeting Distance

- expected number of steps taken for 2 random walks starting from a and b to meet.
- Expected meeting distance in G is equivalent to expected distance in G².
 - Consider a graph $G^2 = (V \times V, E^2)$
 - There is an edge between (a,b) and (c,d) in E², if there are edges (a,c) and (b,d) in E

$$m(a,b) = \sum_{t:(a,b)\rightsquigarrow(x,x)} P[t]l(t)$$



Expected Meeting Distance







 $m(u,v) = \infty$

$$m(u,v) = \infty$$

m(u,w) = ∞
m(v,w) = 1

m(u,v) = 3



Expected-f Meeting Distance

• Map distance I(t) to $f(I(t), where f(z) = c^z, 0 < c < 1$

$$s'(a,b) = \sum_{t:(a,b)\rightsquigarrow(x,x)} P[t]c^{l(t)}$$

- Large distances become small similarities
- Small distances become large similarities



 s(a,b) is equivalent to s'(a,b) where in and out edges are reversed.

$$s'(a,b) = \sum_{t:(a,b) \nleftrightarrow (x,x)} P[t]c^{l(t)}$$

= $\sum_{c \in O(a)} \sum_{d \in O(b)} \sum_{t':(c,d) \twoheadrightarrow (x,x)} \frac{P[t']c^{l(t')+1}}{|O(a)||O(b)|}$
= $\frac{c}{|O(a)||O(b)|} \sum_{c \in O(a)} \sum_{d \in O(b)} s'(c,d)$
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Blocking:

• Identify similar pairs of records.

Bootstrapping:

 Create some high confidence clusters of duplicate amongst blocked pairs.

Iteration:

- Merge two closest clusters if similarity > threshold
- Update the similarities between neighboring clusters based on the fact that the cluster has been merged.



Relational Clustering using an Example





P1: "JOSTLE: Partitioning of Unstructured Meshes for Massively Parallel Machines", C. Walshaw, M.
Cross, M. G. Everett, S. Johnson

P2: "Partitioning Mapping of Unstructured Meshes to Parallel Machine Topologies", C. Walshaw, M.
Cross, M. G. Everett, S. Johnson, K. McManus

P3: "Dynamic Mesh Partitioning: A Unied Optimisation and Load-Balancing Algorithm", C. Walshaw, M. Cross, M. G. Everett

P4: "Code Generation for Machines with Multiregister Operations", Alfred V. Aho, Stephen C. Johnson, Jefferey D. Ullman

P5: "Deterministic Parsing of Ambiguous Grammars", A. Aho, S. Johnson, J. Ullman

P6: "Compilers: Principles, Techniques, and Tools", A. Aho, R. Sethi, J. Ullman

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- 1. Find similar references using 'blocking'
- 2. Bootstrap clusters using attributes and relations
- 3. Compute similarities for cluster pairs and insert into priority queue
- 4. Repeat until priority queue is empty
- 5. Find 'closest' cluster pair
- 6. Stop if similarity below threshold
- 7. Merge to create new cluster
- 8. Update similarity for 'related' clusters

O(n k log n) algorithm w/ efficient implementation

38

- Never split clusters, only merge them
 - Allows efficient implementation
 - Errors early on in the process can lead to bad clustering/resolution

- Collective Resolution
 - Two objects that are not very similar can become similar if their neighbors are clustered together.



Summary

- Many similarity metrics for relational data
 - Common Neighbors
 - Adamic/Adar
 - Katz
 - Personalized Page Rank
 - Simrank
- Need collective techniques for entity resolution on linked data
 - Relational Clustering
- Next Class
 - Collective Resolution using Markov Logic
 - Scaling Collective Entity Resolution

