

Entity Resolution: Tutorial

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http://www.cs.umd.edu/~getoor/Tutorials/ER_ASONAM2012.pdf

What is Entity Resolution?

Problem of identifying and linking/grouping different manifestations of the same real world object.

Examples of manifestations and objects:

- Different ways of addressing (names, email addresses, FaceBook accounts) the same person in text.
- Web pages with differing descriptions of the same business.
- Different photos of the same object.
- ...

What is Entity Resolution?



Article [Talk](#)

[Read](#) [Edit](#) [View history](#)

Search



Record linkage

From Wikipedia, the free encyclopedia

(Redirected from [Entity resolution](#))

Record linkage (RL) refers to the task of finding **records** in a data set that refer to the same **entity** across different data sources (e.g., data files, books, websites, databases). Record linkage is necessary when **joining** data sets based on entities that may or may not share a common identifier (e.g., **database key**, **URI**, **National identification number**), as may

Name resolution

From Wikipedia, the free encyclopedia

Coreference

From Wikipedia, the free encyclopedia

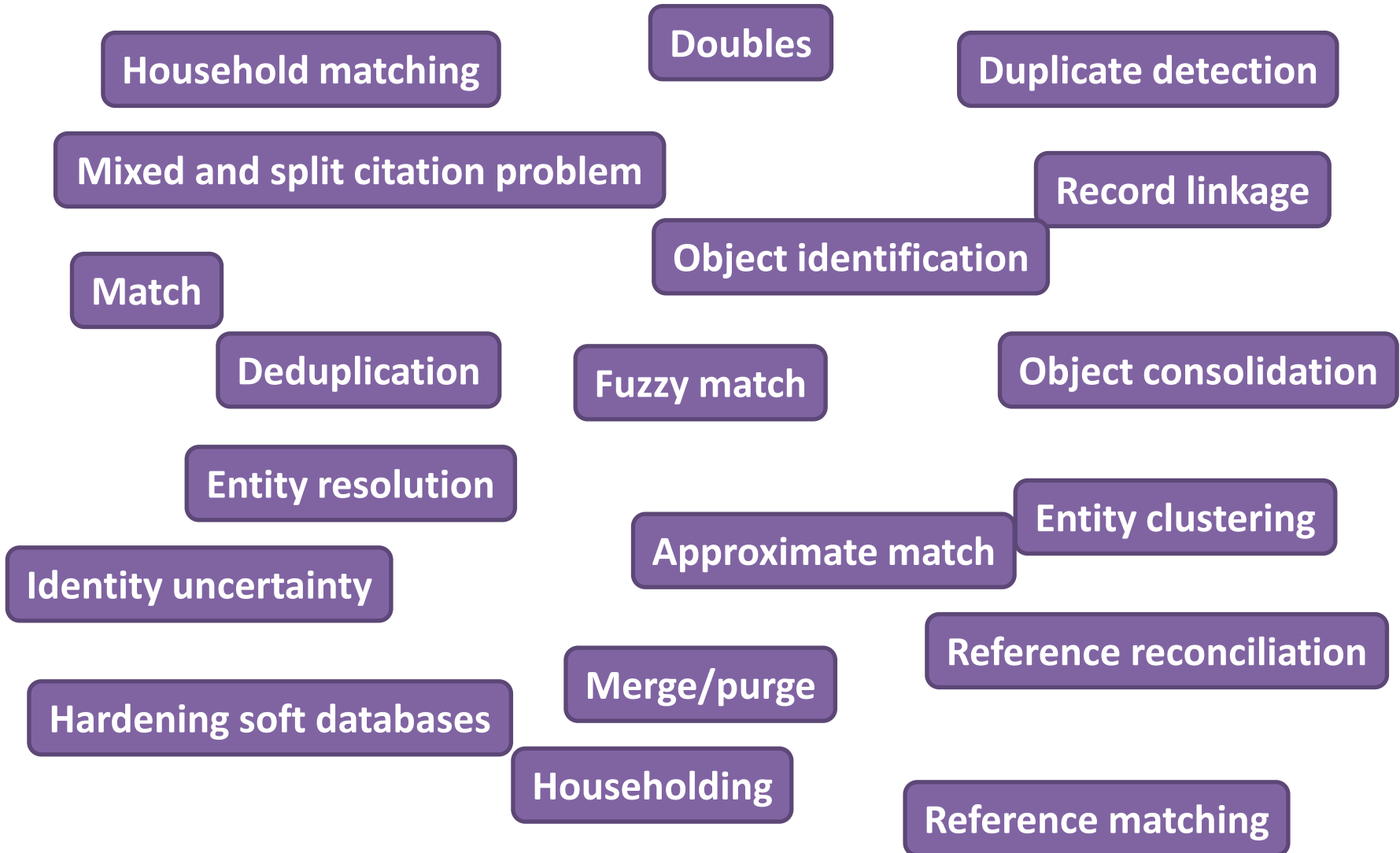
Deduplication

From Wikipedia, the free encyclopedia

Identity resolution

From Wikipedia, the free encyclopedia

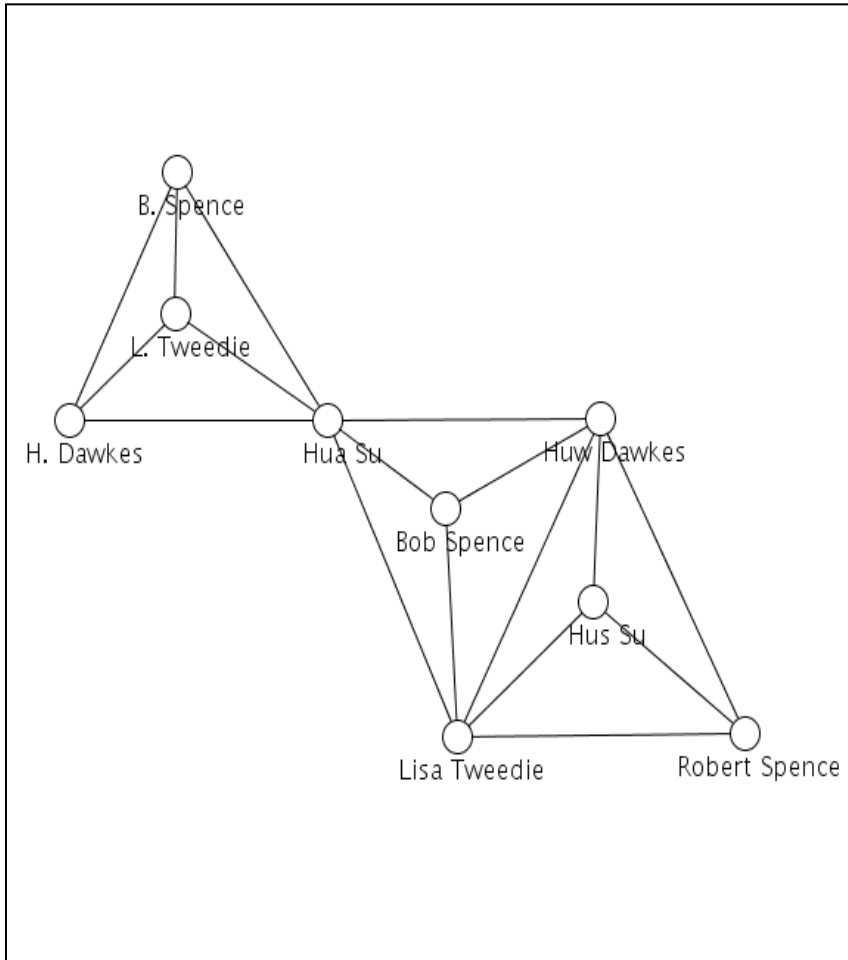
Ironically, Entity Resolution has many duplicate names



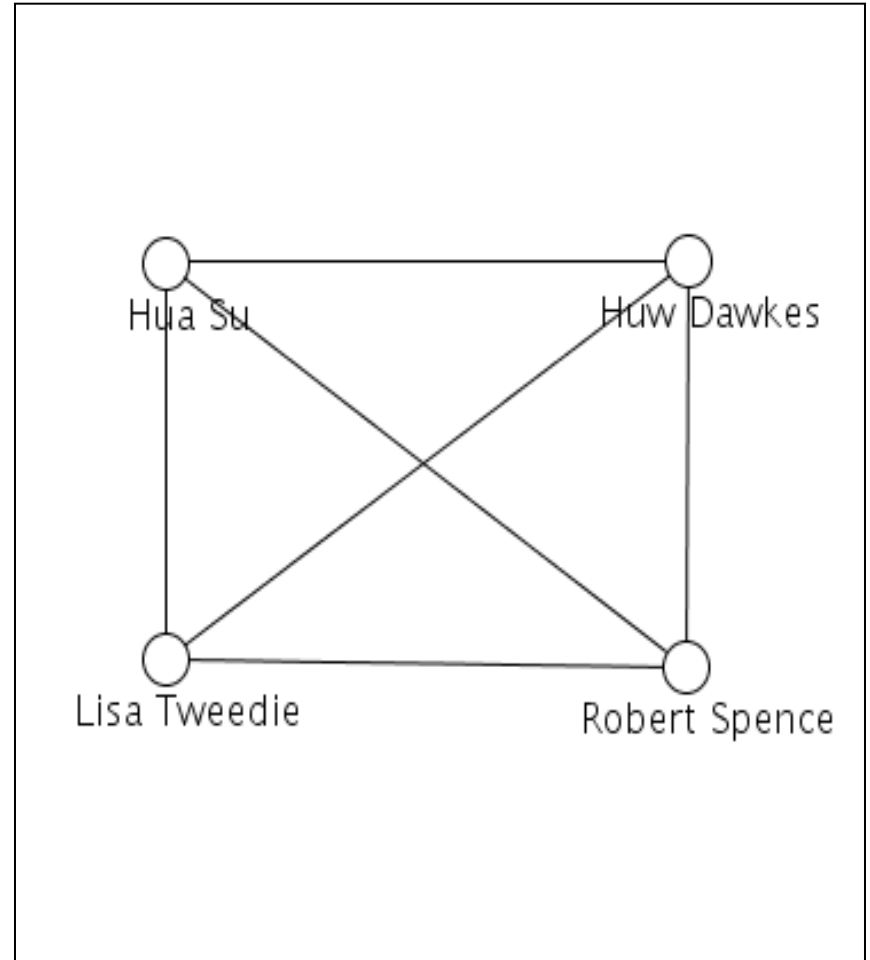
ER Motivating Examples

- *Linking Census Records*
- *Public Health*
- *Web search*
- *Comparison shopping*
- *Counter-terrorism*
- *Spam detection*
- *Machine Reading*
- ...

ER and Network Analysis



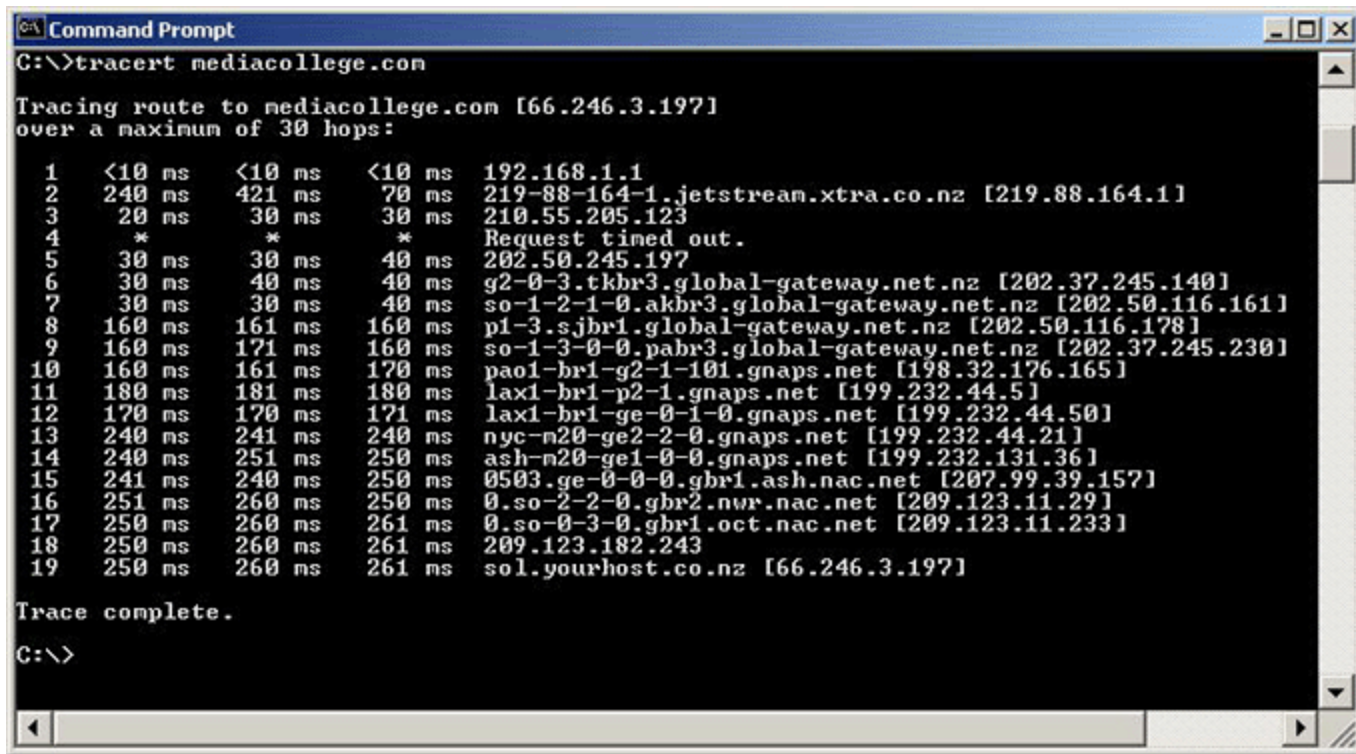
before



after

Motivation: Network Science

- Measuring the topology of the internet ... using traceroute



```
C:\>tracert mediacollege.com

Tracing route to mediacollege.com [66.246.3.197]
over a maximum of 30 hops:

  0  <10 ms  <10 ms  <10 ms  192.168.1.1
  1  240 ms  421 ms  70 ms  219-88-164-1.jetstream.xtra.co.nz [219.88.164.1]
  2  20 ms  30 ms  30 ms  210.55.205.123
  3  *      *      *      Request timed out.
  4  30 ms  30 ms  40 ms  202.50.245.197
  5  30 ms  40 ms  40 ms  g2-0-3.tkbr3.global-gateway.net.nz [202.37.245.140]
  6  30 ms  30 ms  40 ms  so-1-2-1-0.akbr3.global-gateway.net.nz [202.50.116.161]
  7  160 ms  161 ms  160 ms  p1-3.sjbr1.global-gateway.net.nz [202.50.116.178]
  8  160 ms  171 ms  160 ms  so-1-3-0-0.pabr3.global-gateway.net.nz [202.37.245.230]
  9  160 ms  161 ms  170 ms  pao1-br1-g2-1-101.gnaps.net [198.32.176.165]
 10  180 ms  181 ms  180 ms  lax1-br1-p2-1.gnaps.net [199.232.44.5]
 11  170 ms  170 ms  171 ms  lax1-br1-ge-0-1-0.gnaps.net [199.232.44.50]
 12  240 ms  241 ms  240 ms  nyc-n20-ge2-2-0.gnaps.net [199.232.44.21]
 13  240 ms  251 ms  250 ms  ash-n20-ge1-0-0.gnaps.net [199.232.131.36]
 14  241 ms  240 ms  250 ms  0503.ge-0-0-0.gbr1.ash.nac.net [207.99.39.157]
 15  251 ms  260 ms  250 ms  0.so-2-2-0.gbr2.nwr.nac.net [209.123.11.29]
 16  250 ms  260 ms  261 ms  0.so-0-3-0.gbr1.oct.nac.net [209.123.11.233]
 17  250 ms  260 ms  261 ms  209.123.182.243
 18  250 ms  260 ms  261 ms  sol.yourhost.co.nz [66.246.3.197]

Trace complete.

C:\>
```

IP Aliasing Problem [Willinger et al. 2009]

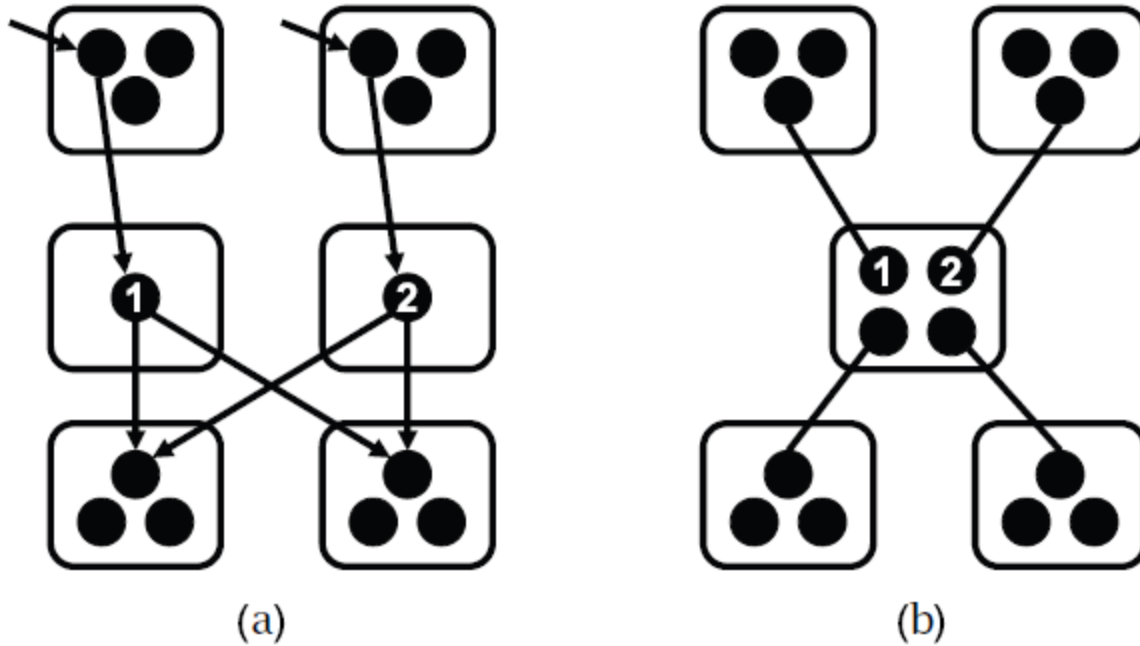


Figure 2. The IP alias resolution problem. Paraphrasing Fig. 4 of [50], traceroute does not list routers (boxes) along paths but IP addresses of input interfaces (circles), and alias resolution refers to the correct mapping of interfaces to routers to reveal the actual topology. In the case where interfaces 1 and 2 are aliases, (b) depicts the actual topology while (a) yields an “inflated” topology with more routers and links than the real one.

IP Aliasing Problem [Willinger et al. 2009]

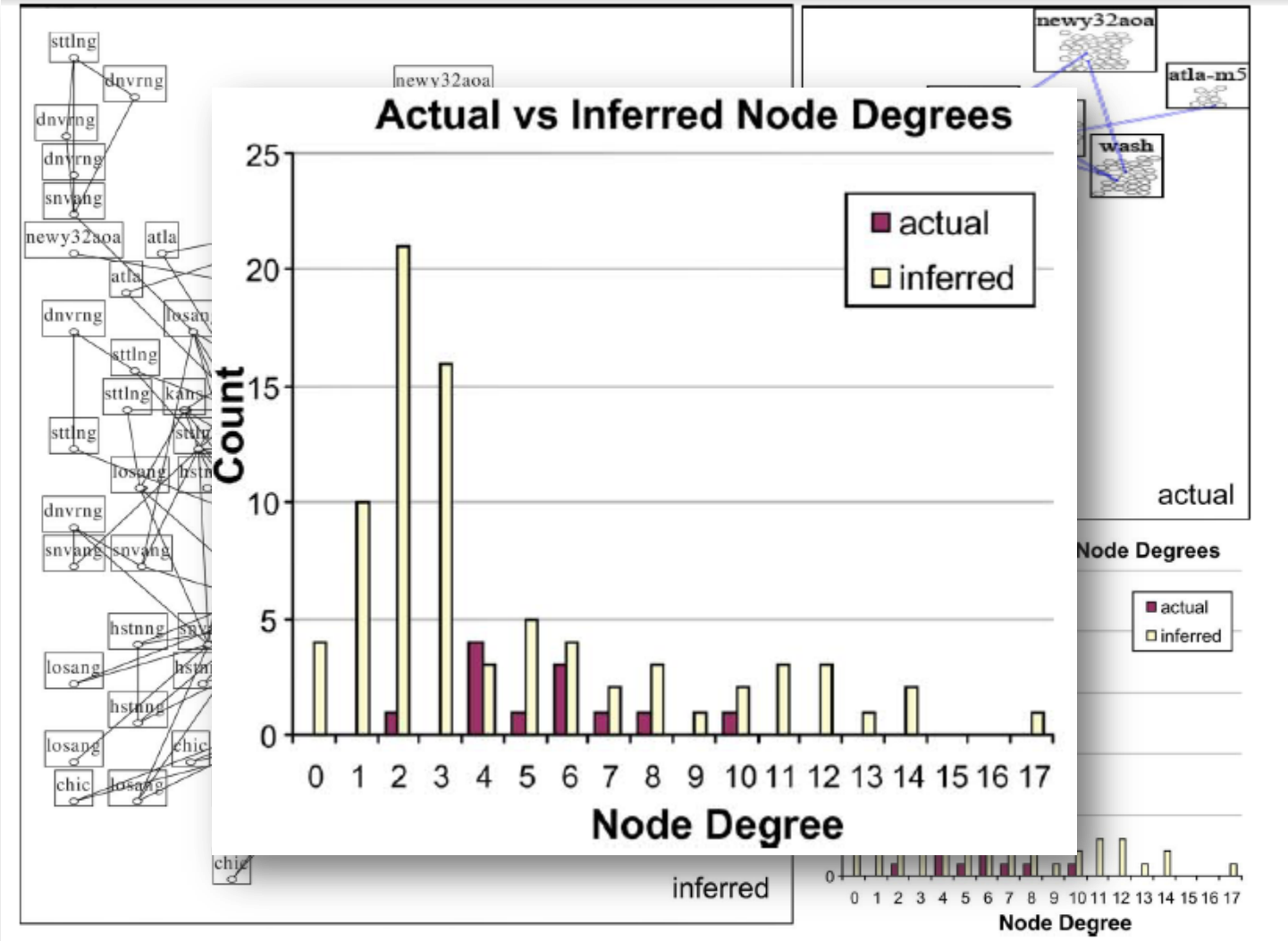
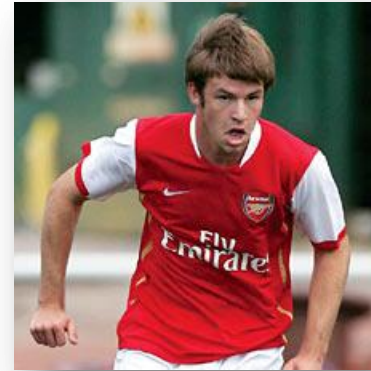


Figure 3. The IP alias resolution problem in practice. This is re-produced from [48] and shows a comparison between the Abilene/Internet2 topology inferred by Rocketfuel (left) and the actual topology (top right). Rectangles represent routers with interior ovals denoting interfaces. The histograms of the corresponding node degrees are shown in the bottom right plot. © 2008 ACM.

Traditional Challenges in ER

- Name/Attribute ambiguity

Thomas Cruise



Michael Jordan



Traditional Challenges in ER

- Name/Attribute ambiguity
- Errors due to data entry



↓	C1	C2
	Total Cholesterol_1	Total Cholesterol_2
682	214.4	214.4
683	184.4	184.4
684	183.5	183.5
685	240.7	240.7
686	215.1	215.1
687	198.6	198.6
688	2800.0	280.0
689	210.8	210.8
690	182.5	182.5
691	192.6	192.6

Traditional Challenges in ER

- Name/Attribute ambiguity
- Errors due to data entry
- Missing Values

Exhibit 2: Examples of variables that are set to unknown values

Administrative dates: set to 0101YY, 010199, 999999

Date of Birth 0101YY, 1506YY, 3006YY, 0107YY, 1507YY, 0101YEAR

Names: set to spaces, NK, UNKNOWN, or ZZZZ
BABY, MALE, FEMALE, TWIN, TRIPLET, INFANT

Other variables: set to 9, 99, 9999, -1

NK (Not Known)
NA (Not applicable)
NC (Not coded)
U (Unknown)

Traditional Challenges in ER

- Name/Attribute ambiguity
- Errors due to data entry
- Missing Values
- Changing Attributes

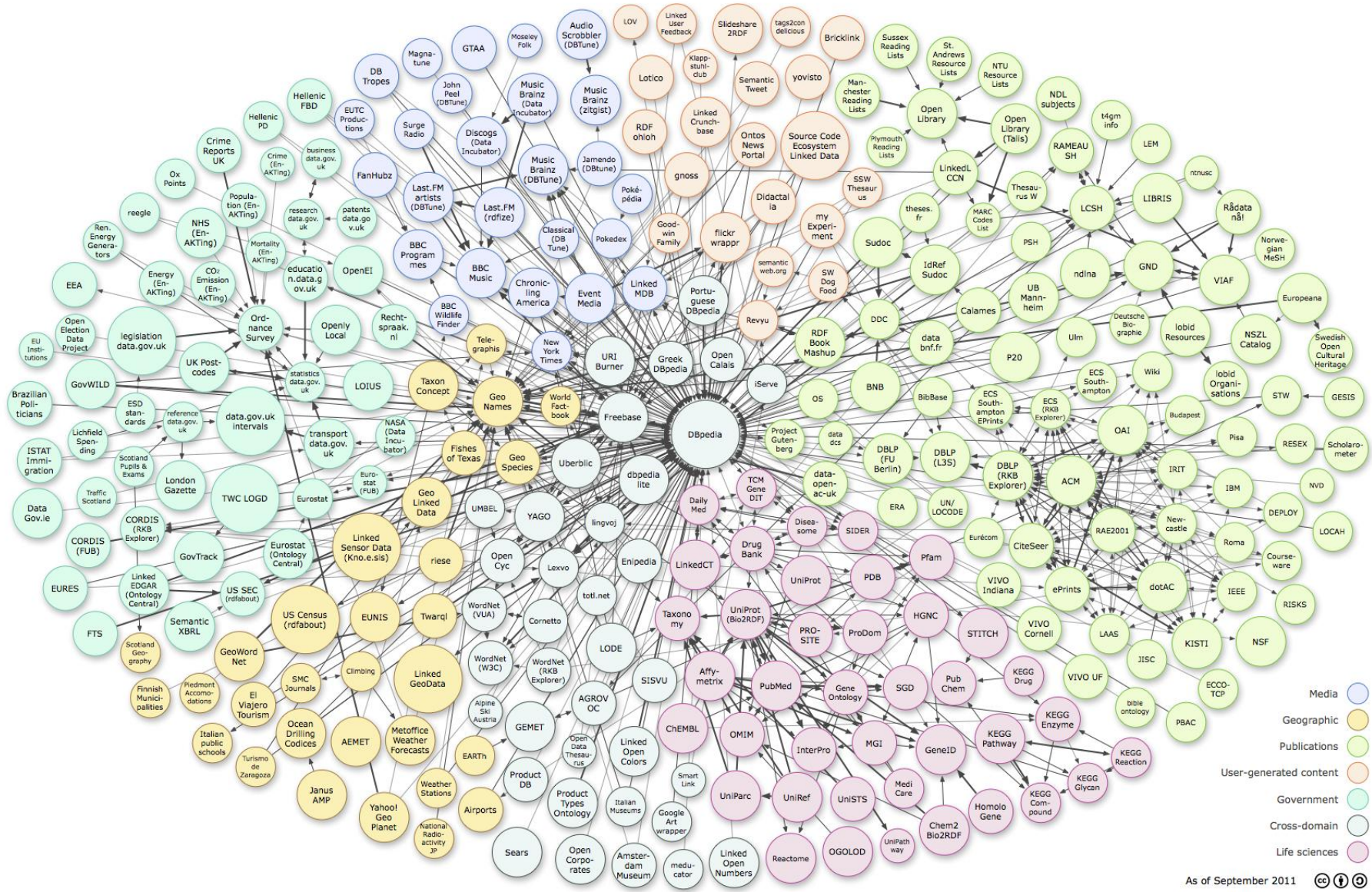


- Data formatting



- Abbreviations / Data Truncation

Big-Data ER Challenges



Big-Data ER Challenges

- Larger and more Datasets
 - Need efficient parallel techniques
- More Heterogeneity
 - Unstructured, Unclean and Incomplete data. Diverse data types.
 - No longer just matching names with names, but Amazon profiles with browsing history on Google and friends network in Facebook.

Big-Data ER Challenges

- Larger and more Datasets
 - Need efficient parallel techniques
- More Heterogeneity
 - Unstructured, Unclean and Incomplete data. Diverse data types.
- More linked
 - Need to infer relationships in addition to “equality”
- Multi-Relational
 - Deal with structure of entities (Are Walmart and Walmart Pharmacy the same?)
- Multi-domain
 - Customizable methods that span across domains
- Multiple applications (web search versus comparison shopping)
 - Serve diverse application with different accuracy requirements

Outline

1. Classical Single Entity ER
2. Relational & MultiEntity ER
3. Efficiency: Blocking/Canopies
4. Challenges & Future Directions

PART 1

CLASSICAL SINGLE ENTITY ER

Outline

1. Classical Single Entity ER
 - a) Problem Statement
 - b) Data Preparation & Matching Features
 - c) Algorithms for Single-Entity ER
 - d) Canonicalization
2. Relational & MultiEntity ER
3. Efficiency: Blocking/Canopies
4. Challenges & Future Directions

PART 1-a

ER PROBLEM STATEMENT

Abstract Problem Statement

Real World



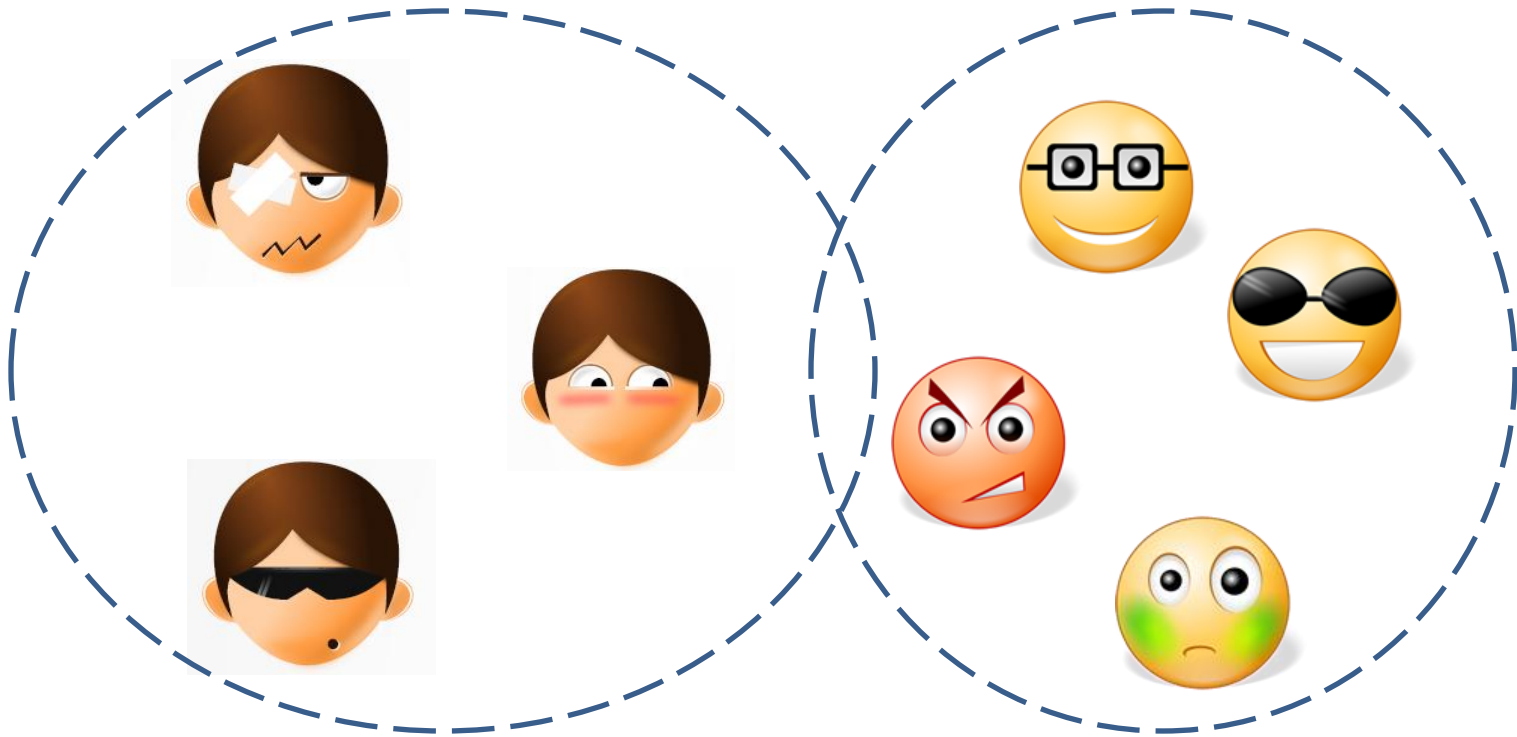
Digital World



Records /
Mentions

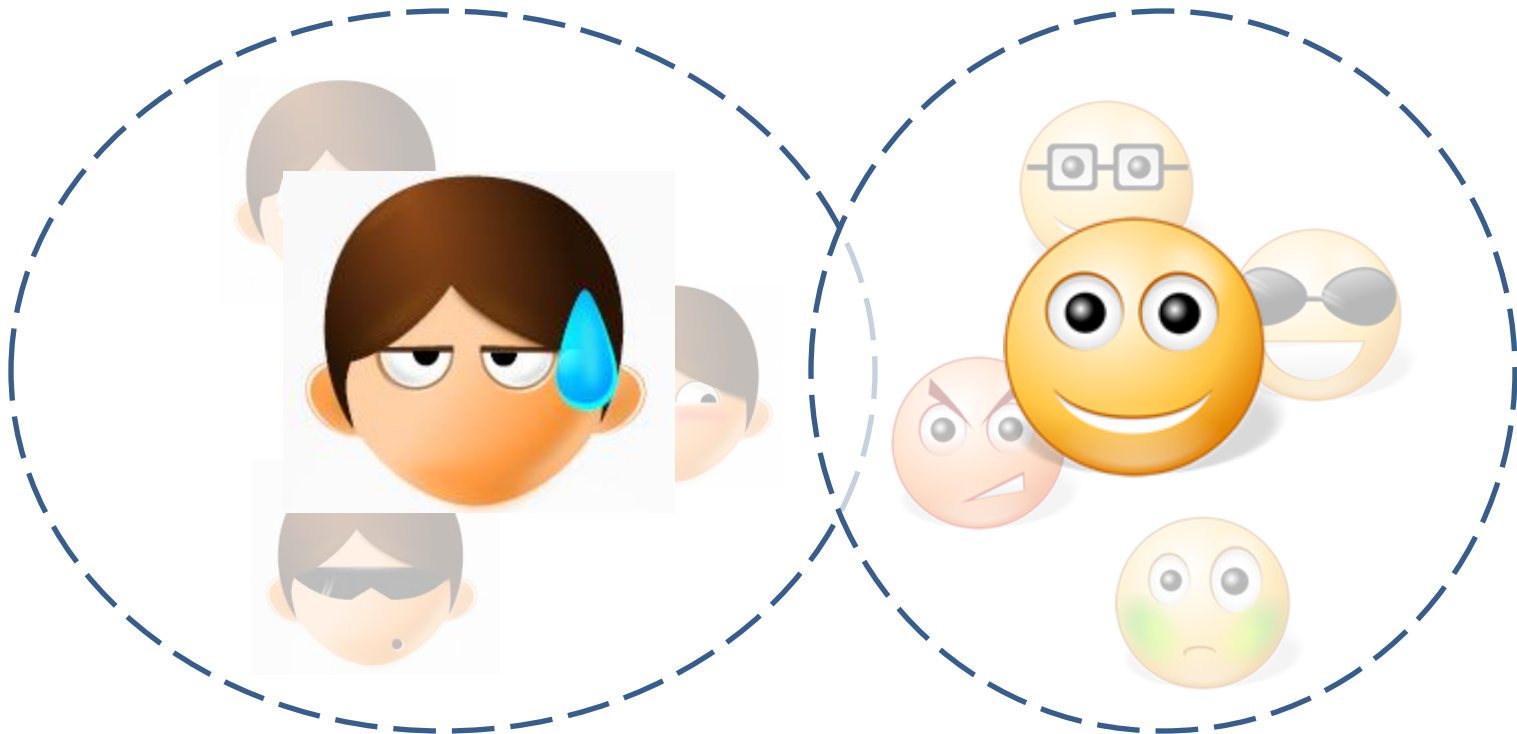
Deduplication Problem Statement

- Cluster the records/mentions that correspond to same entity



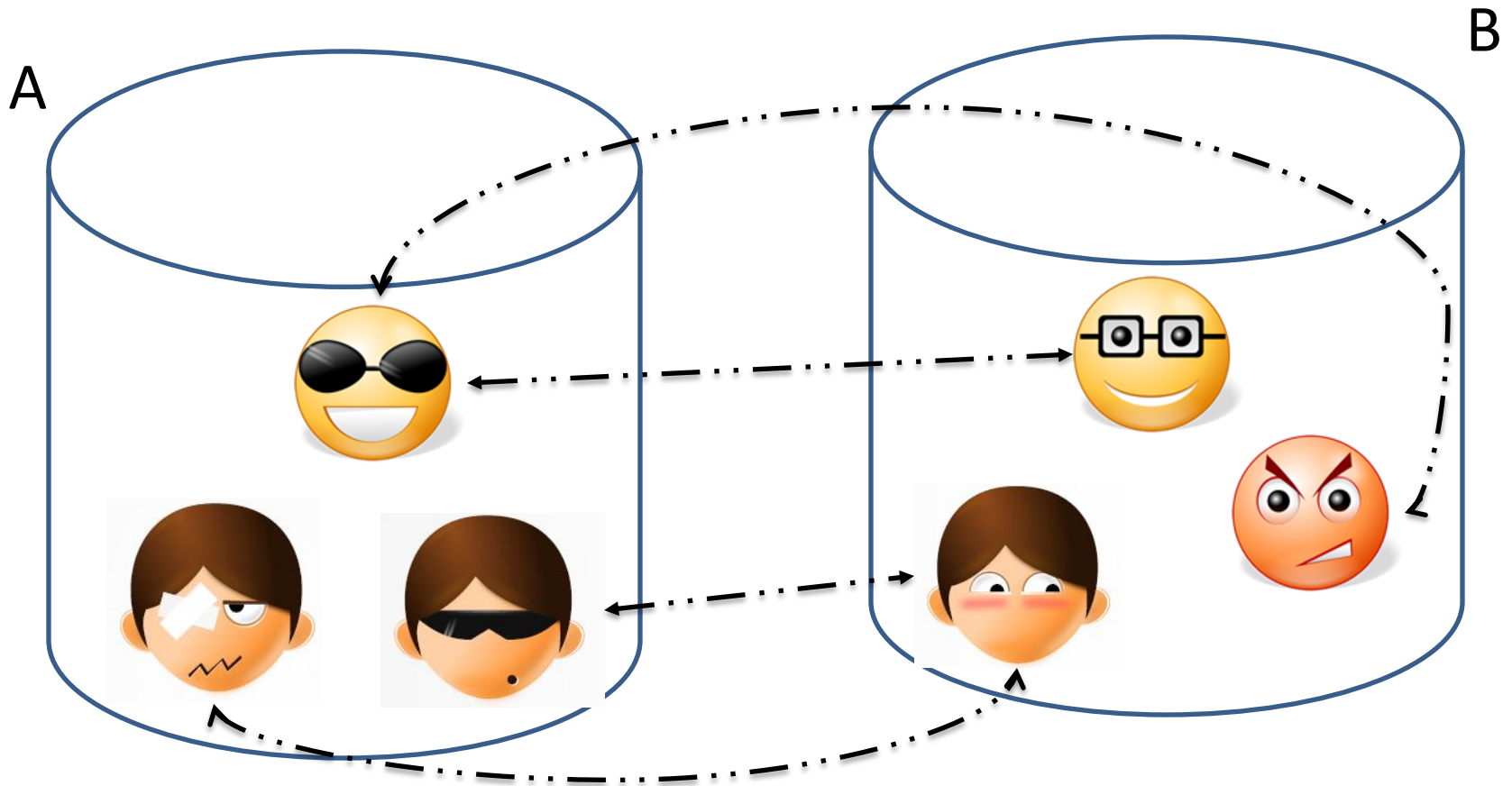
Deduplication Problem Statement

- Cluster the records/mentions that correspond to same entity
 - **Intensional Variant:** Compute cluster representative



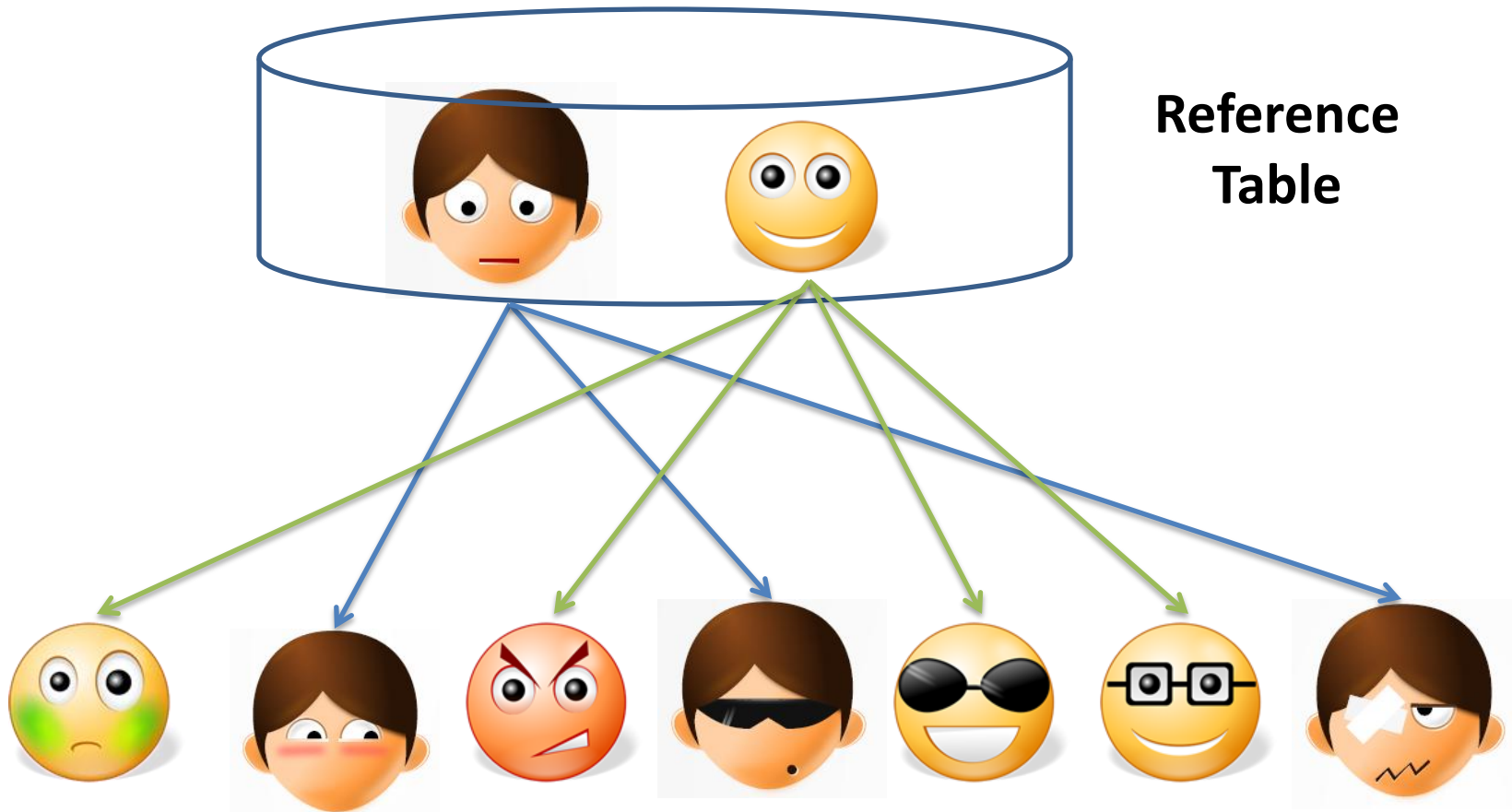
Record Linkage Problem Statement

- Link records that match across databases



Reference Matching Problem

- Match noisy records to clean records in a reference table



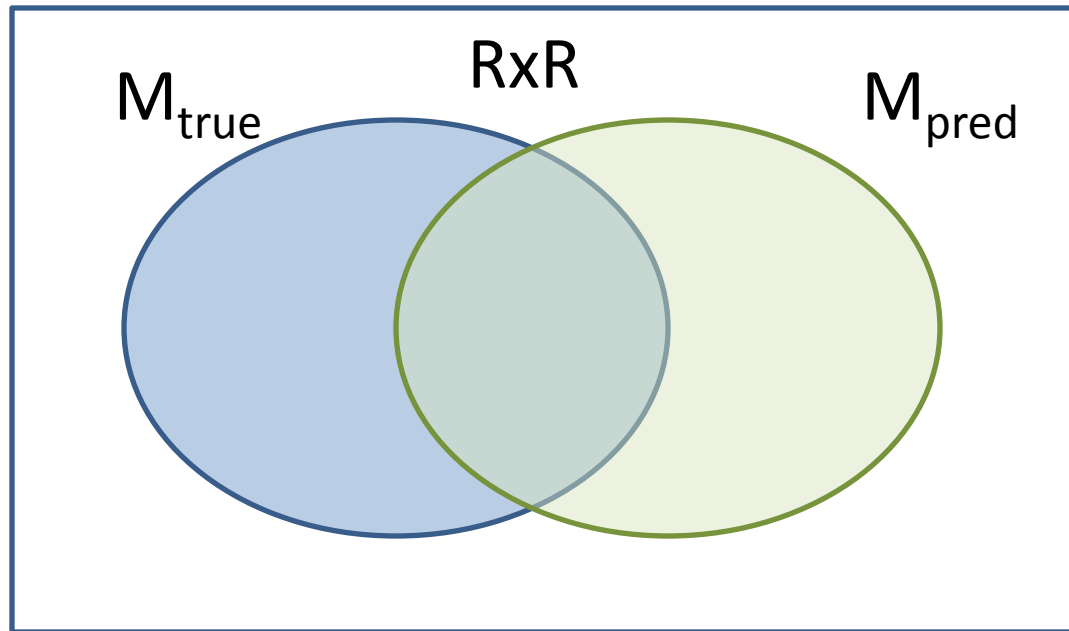
Notation & Assumptions

- R : set of records / mentions
- M : set of *matches* (record pairs that correspond to same entity)
- N : set of *non-matches* (record pairs corresponding to different entities)
- E : set of entities

- True ($M_{true}, N_{true}, E_{true}$): according to real world
vs Predicted ($M_{pred}, N_{pred}, E_{pred}$): by algorithm

Relationship between M_{true} and M_{pred}

- M_{true} (SameAs , Equivalence)
- M_{pred} (Similar representations and similar attributes)

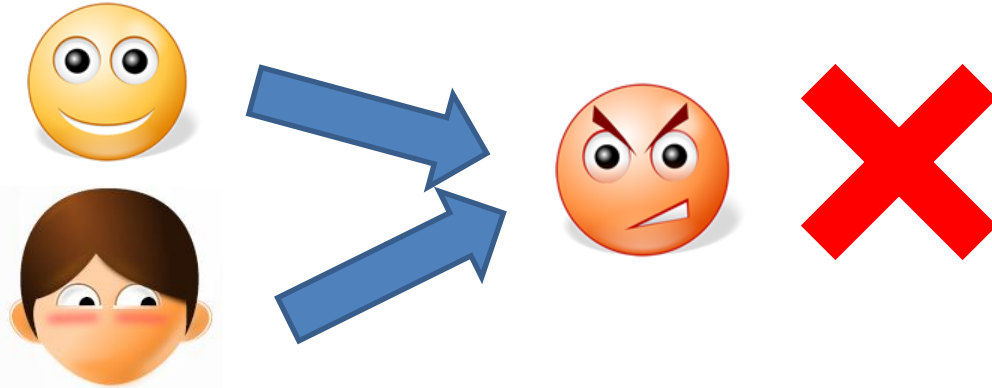


Metrics

- Pairwise metrics
 - Precision/Recall, F1
 - # of predicted matching pairs
- Cluster level metrics
 - purity, completeness, complexity
 - Precision/Recall/F1: Cluster-level, closest cluster, MUC, B^3 , Rand Index
 - Generalized merge distance [Menestrina et al, PVLDB10]

Typical Assumptions Made

- *Each record/mention is associated with a single real world entity.*



- *In record linkage, no duplicates in the same source*
- *If two records/mentions are identical, then they are true matches*

$$\left(\text{👤}, \text{👤} \right) \in M_{\text{true}}$$

ER versus Classification

Finding matches vs non-matches is a classification problem

- Imbalanced: typically $O(R)$ matches, $O(R^2)$ non-matches
- Instances are pairs of records. Pairs are not IID

 $\in M_{\text{true}}$

AND



 $\in M_{\text{true}}$

 $\in M_{\text{true}}$

ER vs Clustering

Computing entities from records is a clustering problem

- In typical clustering algorithms (k-means, LDA, etc.) *number of clusters is a constant or sub linear in R .*
- In ER: *number of clusters is linear in R , and average cluster size is a constant. Significant fraction of clusters are singletons.*

PART 1-b

**DATA PREPARATION &
MATCH FEATURES**

Normalization

- Schema normalization
 - Schema Matching – e.g., contact number and phone number
 - Compound attributes – full address vs str,city,state,zip
 - Nested attributes
 - List of features in one dataset (air conditioning) vs each feature a boolean attribute
 - Set valued attributes
 - Set of phones vs phone
 - Record segmentation
- Data normalization
 - Often case sensitive/all lower/all upper; remove whitespace
 - detecting and correcting values that contain known typographical errors or variations,
 - expanding abbreviations and replacing them with standard forms; replacing nicknames with their proper name forms
 - Usually done based on dictionaries (e.g., commercial dictionaries, postal addresses, etc.)

Initial data prep big part of the work; smart normalization can go long way!

Matching Functions

- For two references x and y , compute a “comparison” vector, typically similarity of each component attribute.
- Distance metric:
 - Idempotent
 - Non-negative
 - Symmetric
 - Triangle inequality
- Not all commonly used ER distance functions are metrics
 - non-linear elastic matching (NEM)
- From distance, can convert to similarity:
 - $S = 1 / d$, or if d is normalized, $s = 1-d$

Summary of Matching Functions

Handle
Typographical errors

- Equality on a boolean predicate
- Edit distance
 - Levenstein, Smith-Waterman, Affine

- Set similarity
 - Jaccard, Dice
- Vector Based
 - Cosine similarity, TFIDF

Good for Text like
reviews/ tweets

- Useful packages:
 - SecondString, <http://secondstring.sourceforge.net/>
 - Simmetrics: <http://sourceforge.net/projects/simmetrics/>
 - LingPipe, <http://alias-i.com/lingpipe/index.html>

Good for Names

- Alignment-based or Two-tiered
 - Jaro-Winkler, Soft-TFIDF, Monge-Elkan
- Phonetic Similarity
 - Soundex
- Translation-based
- Numeric distance between values
- Domain-specific

Useful for
abbreviations,
alternate names.

PART 1-c

ALGORITHMS FOR SINGLE-ENTITY ER

Matching Algorithms

- Pairwise Matching
 - Given a vector of comparison scores,
Independently compute a (probability) score indicative of whether a pair of records/mentions match.
- Record Linkage
 - Each record from one database matches at most one record from other database.
 - Weighted k-partite matching
- Deduplication
 - Transitivity constraints must be satisfied.
 - Correlation Clustering

PAIRWISE MATCHING

Pairwise Match Score

Problem: Given a vector of component-wise similarities for a pair of records (x,y), compute P(x and y match).

Solutions:

1. Weighted sum or average of component-wise similarity scores.
Threshold determines match or non-match.
 - $0.5 * \text{Last-name-match-score} + 0.2 * \text{address-match-score} + 0.3 * \text{login-match-score}$.
 - Hard to pick weights.
 - Match on last name match *more predictive* than login name.
 - Match on “Smith” *less predictive* than match on “Getoor”.
 - Hard to tune a threshold.

Pairwise Match Score

Problem: Given a vector of component-wise similarities for a pair of records (x,y) , compute $P(x \text{ and } y \text{ match})$.

Solutions:

1. Weighted sum or average of component-wise similarity scores.
Threshold determines match or non-match.
2. Formulate rules about what constitutes a match.
 - (Last-name-score > 0.7 AND address-match-score > 0.8)
OR (login-match-score > 0.9 AND address-match-score > 0.9)
 - Manually formulating the right set of rules is hard.

ML Pairwise Approaches

- Supervised machine learning algorithms
 - Decision trees
 - [Cochinwala et al, IS01]
 - Support vector machines
 - [Bilenko & Mooney, KDD03]; [Christen, KDD08]
 - Ensembles of classifiers
 - [Chen et al., SIGMOD09]
 - Conditional Random Fields (CRF)
 - [Wellner & McCallum, NIPS04]
- Issues:
 - **Training set generation**
 - Imbalanced classes – many more negatives than positives (even after eliminating obvious non-matches ... using *Blocking*)
 - Misclassification cost

Creating a Training Set is a key issue

- Constructing a training set is hard – since most pairs of records are “easy non-matches”.
 - 100 records from 100 cities.
 - Only 10^6 pairs out of total 10^8 (1%) come from the same city
- Some pairs are hard to judge even by humans
 - Inherently ambiguous
 - E.g., Paris Hilton (person or business)
 - Missing attributes
 - Starbucks, Toronto vs Starbucks, Queen Street ,Toronto

Avoiding Training Set Generation

- Unsupervised / Semi-supervised Techniques
 - Fellegi-Sunter Model
 - [Newcombe et al Science '59, Fellegi & Sunter JASA 69, Winkler '06, Herzog et al '07]
 - Generative Models
 - [Ravikumar & Cohen, UAI04]

Fellegi & Sunter Model

- $r = (x, y)$ is record pair, γ is comparison vector, M matches, U non-matches
- In the original work, γ is binary, 0/1, match/not match

- Decision rule
$$R = \frac{P(\gamma | r \in M)}{P(\gamma | r \in U)}$$

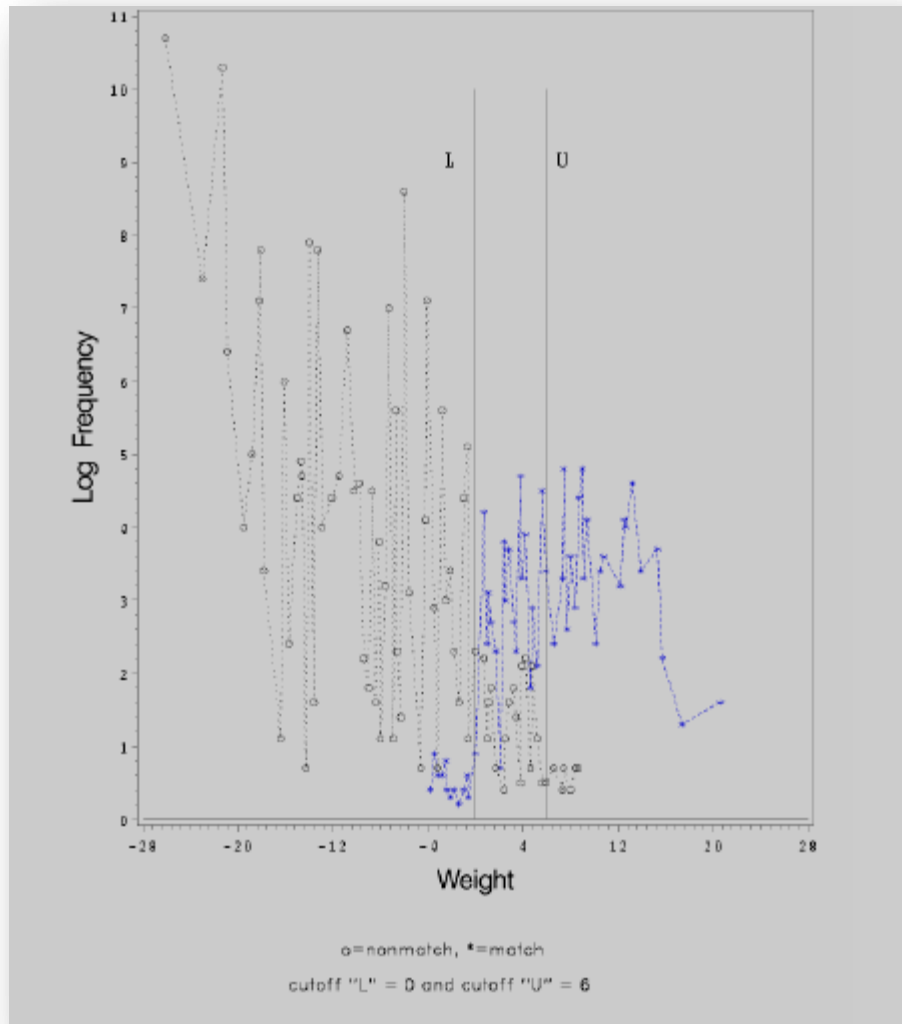
$$R \geq t_u \Rightarrow r \rightarrow \text{Match}$$

$$t_l < R < t_u \Rightarrow r \rightarrow \text{Potential Match}$$

$$R \leq t_u \Rightarrow r \rightarrow \text{Non - Match}$$

- Thresholds t_u and t_l determined by apriori bounds on false matches and false non-matches

Fellegi & Sunter Model



[Winkler 2006]

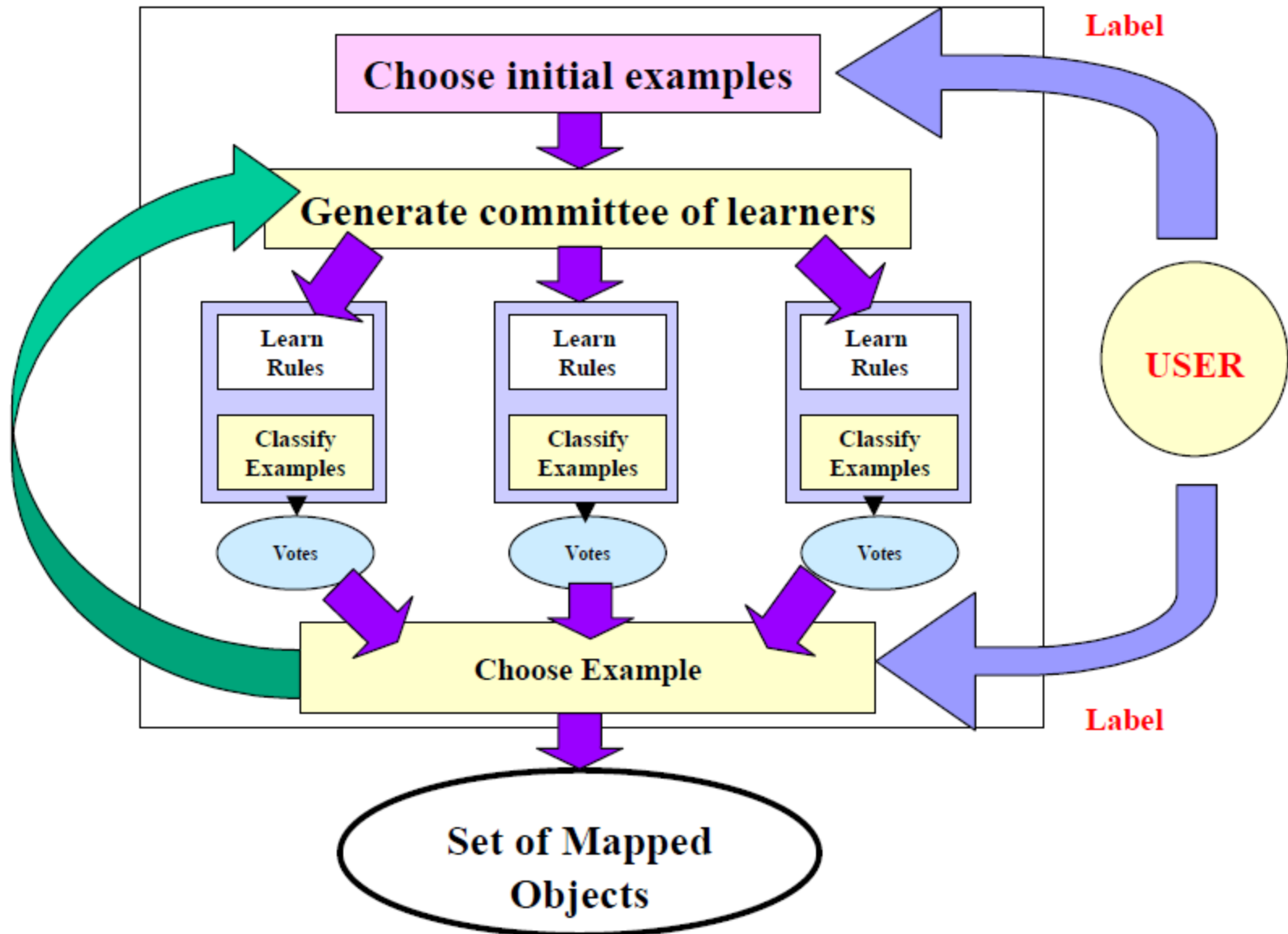
Computing Probabilities

- Typically make an independence assumption
- Agreement weight w_i is calculated for each attribute i based on m and u probabilities:
 - $m_i = P(x_i = y_i \mid r \in M)$
 - $u_i = P(x_i = y_i \mid r \in U)$
- Probabilities can be estimated using EM
 - See [Winkler 2006] for a survey of techniques used in the US Census.

Avoiding Training Set Generation

- Unsupervised / Semi-supervised Techniques
 - Fellegi-Sunter Model
 - [Newcombe Science '59, Fellegi & Sunter JASS 69, Winkler '99, Herzog et al '07]
 - Generative Models
 - [Ravikumar & Cohen, UAI04]
- Active Learning
 - Committee of Classifiers
 - [Sarawagi et al KDD '00, Tajeda et al IS '01]
 - Provably optimizing precision/recall
 - [Arasu et al SIGMOD '10, Bellare et al KDD '12]

Committee of Classifiers [Tejada et al, IS '01]



Active Learning with Provable Guarantees

- Most active learning techniques minimize 0-1 loss
[Beygelzimer et al NIPS 2010].

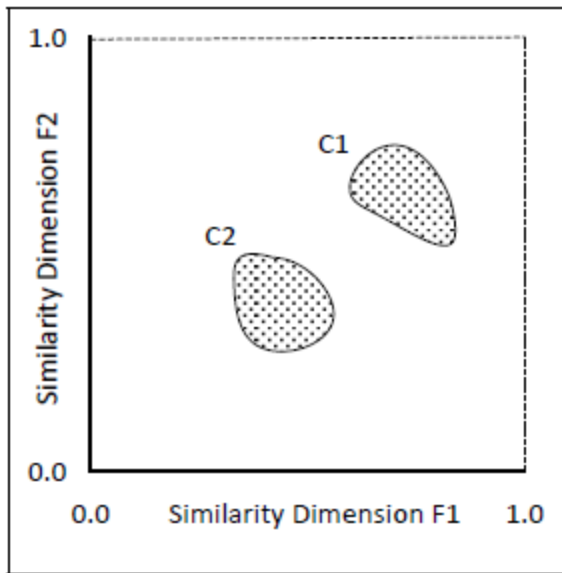
$$\text{minimize } \frac{f_n(h) + f_p(h)}{n}$$

- However, ER is very imbalanced:
 - Number of non-matches \gg number of matches.
 - Classifying all pairs as “non-matches” has low 0-1 loss.
- Hence, need active learning techniques that minimize precision/recall.

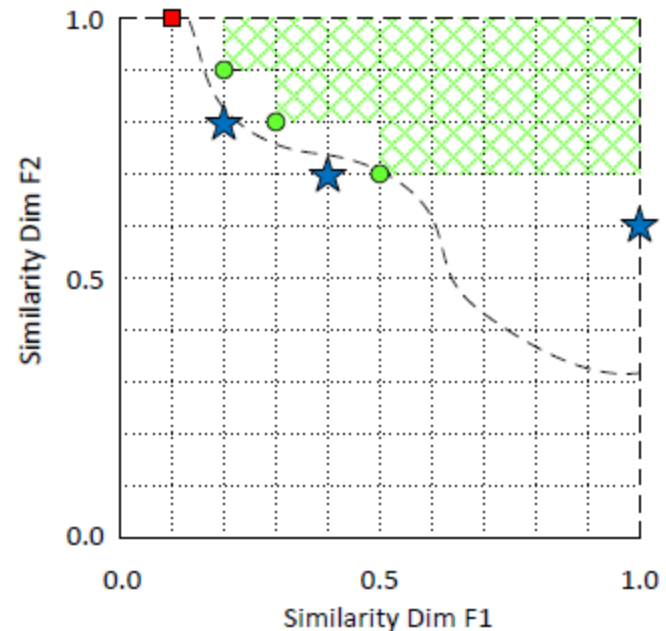
$$\begin{array}{ll} \text{maximize} & \text{recall}(h) \\ \text{subject to} & \text{precision}(h) \geq \tau \end{array}$$

Active Learning with Provable Guarantees

- Monotonicity of Precision [Arasu et al SIGMOD '10]



There is a larger fraction of matches in C1 than in C2.



Algorithm searches for the optimal classifier using binary search on each dimension

Active Learning with Provable Guarantees

[Bellare et al KDD '12]

$O(\log^2 n)$ calls to a blackbox 0-1 loss active learning algorithm.

**Exponentially smaller label complexity than [Arasu et al SIGMOD '10]
(in the worst case).**

1. Precision Constrained \rightarrow Weighted 0-1 Loss Problem
(using a Lagrange Multiplier λ).
2. Given a fixed value for λ , weighted 0-1 Loss can be optimized by a balckbox active learning classifier.
3. Right value of λ is computed by searching over all optimal classifiers.
 - Classifiers are embedded in a 2-d plane (precision/recall)
 - Search is along the convex hull of the embedded classifiers

Open challenge

- Handling errors in human judgements:
 - In an experiment on Amazon Mechanical Turk:
 - Each pairwise judgment given to 5 different people
 - Majority of workers agreed on truth on only 90% of pairwise judgements.

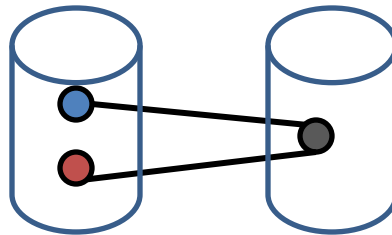
Using pairwise ER

- ER applications need more than independent classification of pairs of records as match/non-match.
- Record Linkage
- Deduplication

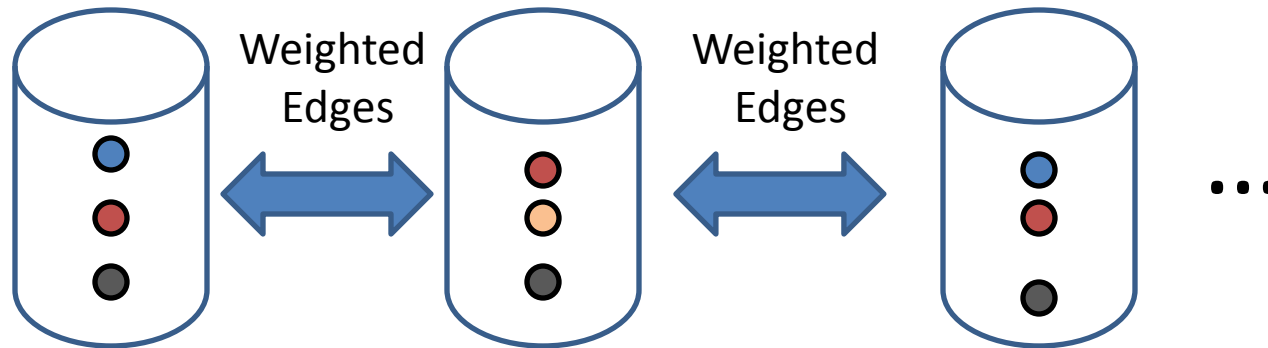
RECORD LINKAGE

1-1 assumption

- Matching between (almost) deduplicated databases.
- Each record in one database matches at most one record in another database.
- Pairwise ER may match a record in one database with more than one record in second database

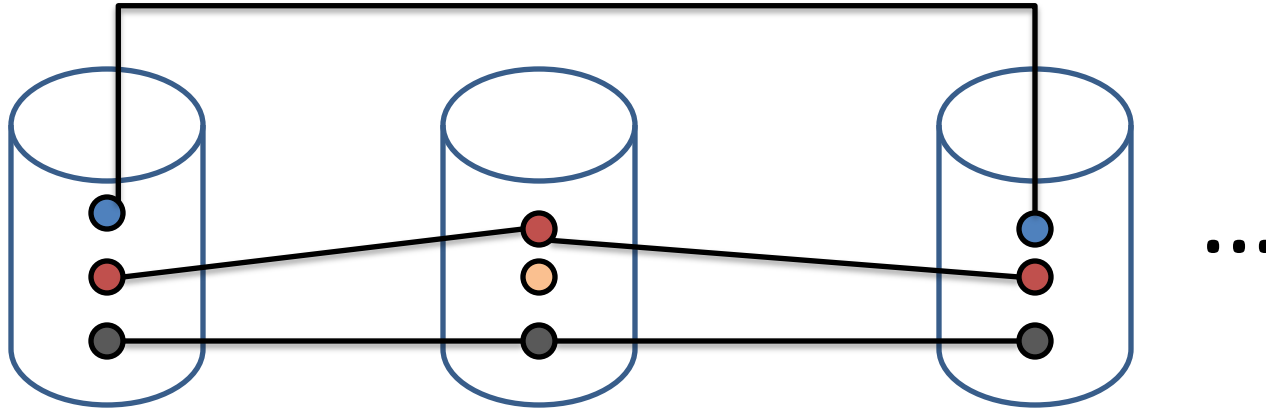


Weighted K-Partite Matching



- Edges between pairs of records from different databases
- Edge weights
 - Pairwise match score
 - Log odds of matching

Weighted K-Partite Matching



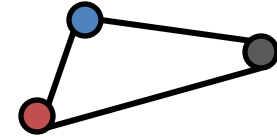
- Find a matching (each record matches at most one other record from other database) that maximize the sum of weights.
- General problem is NP-hard (3D matching)
- Successive bipartite matching is typically used. [Gupta & Sarawagi, VLDB '09]

DEDUPLICATION

Deduplication => Transitivity

- Often pairwise ER algorithm output “inconsistent” results

– $(x, y) \in M_{\text{pred}}, (y, z) \in M_{\text{pred}}, \text{ but } (x, z) \notin M_{\text{pred}}$

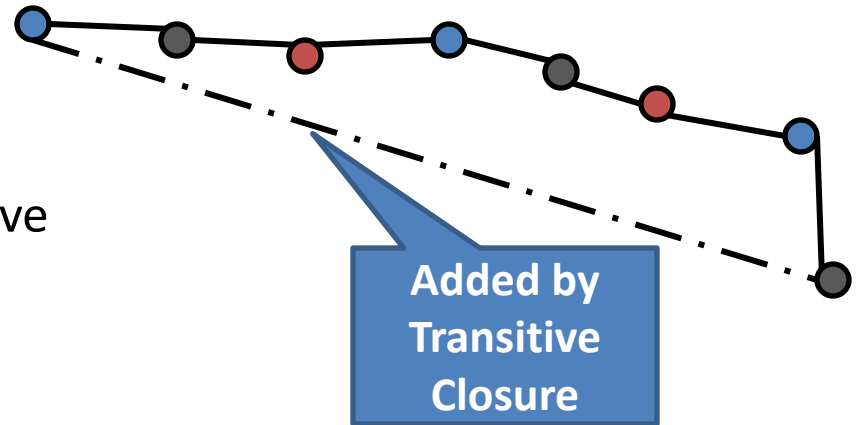


- Idea: Correct this by adding additional matches using transitive closure

- In certain cases, this is a bad idea.

– Graphs resulting from pairwise ER have diameter > 20

[Rastogi et al Corr'12]



- Need clustering solutions that deal with this problem directly by reasoning about records jointly.

Clustering-based ER

- Resolution decisions are not made independently for each pair of records
- Based on variety of clustering algorithms, but
 - Number of clusters unknown a priori
 - Many, many small (possibly singleton) clusters
- Often take a pair-wise similarity graph as input
- May require the construction of a *cluster representative* or *canonical entity*

Clustering Methods for ER

- Hierarchical Clustering
 - [Bilenko et al, ICDM 05]
- Nearest Neighbor based methods
 - [Chaudhuri et al, ICDE 05]
- **Correlation Clustering**
 - [Soon et al CL'01, Bansal et al ML'04, Ng et al ACL'02, Ailon et al JACM'08, Elsner et al ACL'08, Elsner et al ILP-NLP'09]

Integer Linear Programming view of ER

- $r_{xy} \in \{0,1\}$, $r_{xy} = 1$ if records x and y are in the same cluster.
- $w_{xy}^+ \in [0,1]$, cost of clustering x and y together
- $w_{xy}^- \in [0,1]$, cost of placing x and y in different clusters

$$\text{minimize } \sum r_{xy} w_{xy}^+ + (1 - r_{xy}) w_{xy}^-$$

$$\text{s. t. } \forall x, y, z \in R,$$

$$r_{xy} + r_{xz} + r_{yz} \neq 2$$

Transitive
closure

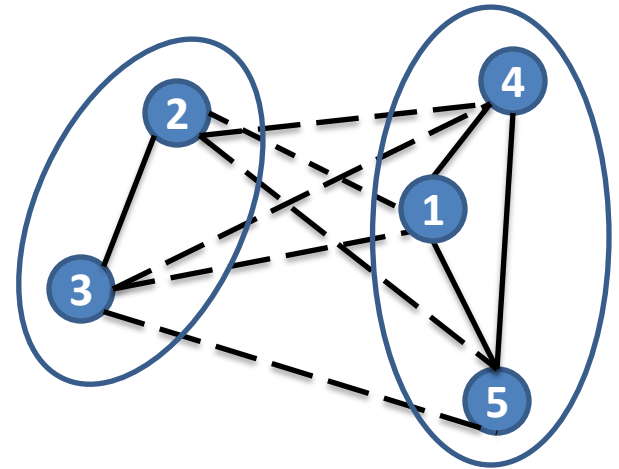
Correlation Clustering

$$\begin{aligned} & \text{minimize } \sum r_{xy} w_{xy}^+ + (1 - r_{xy}) w_{xy}^- \\ & \text{s.t. } \forall x, y, z \in R, \\ & \quad r_{xy} + r_{xz} + r_{yz} \neq 2 \end{aligned}$$

- Cluster mentions such that total cost is minimized

Solid edges contribute w_{xy}^+ to the objective

Dashed edges contribute w_{xy}^- to the objective



- Cost based on pairwise similarities

$$\{p_{xy} \mid \forall (x, y) \in R \times R\}$$

- Additive: $w_{xy}^+ = p_{xy}$ and $w_{xy}^- = (1 - p_{xy})$
- Logarithmic: $w_{xy}^+ = \log(p_{xy})$ and $w_{xy}^- = \log(1 - p_{xy})$

Correlation Clustering

- Solving the ILP is NP-hard [Ailon et al 2008 JACM]
- A number of heuristics [Elsner et al 2009 ILP-NLP]
 - Greedy BEST/FIRST/VOTE algorithms
 - Greedy PIVOT algorithm (5-approximation)
 - Local Search

Greedy Algorithms

Step 1: Permute the nodes according a random π

Step 2: Assign record x to the cluster that maximizes *Quality*

Start a new cluster if *Quality* < 0

Quality:

- BEST: Cluster containing the closest match $\max_{y \in C} w_{xy}^+$
 - [Ng et al 2002 ACL]
- FIRST: Cluster contains the most recent vertex y with $w_{xy}^+ > 0$
 - [Soon et al 2001 CL]
- VOTE: Assign to cluster that minimizes objective function.
 - [Elsner et al 08 ACL]

Practical Note:

- Run the algorithm for many random permutations , and pick the clustering with best objective value (better than average run)

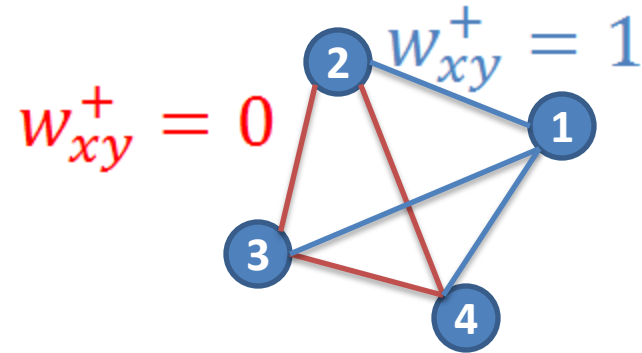
Greedy with approximation guarantees

PIVOT Algorithm

[Ailon et al 2008 JACM]

- Pick a random (*pivot*) record p .
- New cluster = $\{x \mid w_{px}^+ > 0\}$

- $\pi = \{1,2,3,4\}$ $C = \{\{1,2,3,4\}\}$
- $\pi = \{2,4,1,3\}$ $C = \{\{1,2\}, \{4\}, \{3\}\}$
- $\pi = \{3,2,4,1\}$ $C = \{\{1,3\}, \{2\}, \{4\}\}$



When weights are 0/1,

$E(\text{cost}(\text{greedy})) < 3 \text{ OPT}$

For $w_{xy}^+ + w_{xy}^- = 1$,

$E(\text{cost}(\text{greedy})) < 5 \text{ OPT}$

[Elsner et al, ILP-NLP '09] : Comparison of various correlation clustering algorithms

Summary of Single-Entity ER Algorithms

- Many algorithms for independent classification of pairs of records as match/non-match
 - ML based classification & Fellegi-Sunter
 - **Pro: Advanced state of the art**
 - **Con: Building a training set is an open problem**
 - Active learning is becoming popular
- ER applications need more than pairwise classification
 - **Record linkage**: each record matched to at most one record from other database.
 - **Weighted K-Partite Matching**
 - **Deduplication**: transitivity requires clustering based algorithms.
 - **Correlation Clustering**

PART 1-d

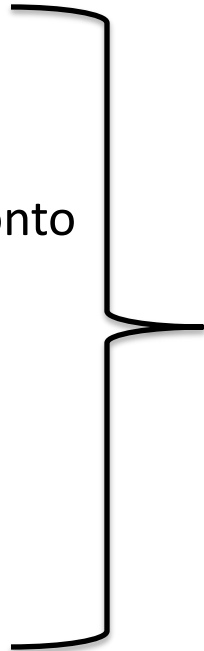
CANONICALIZATION

Canonicalization

- Merge information from duplicate mentions to construct a cluster representative with *maximal* information

- Starbucks,
123 Queen St, Toronto
Ph: *null*

- Starbucks,
null, Toronto
Ph: 333-4444



**Starbucks,
123 Queen St, Toronto
Ph: 333-4444**

Canonicalization

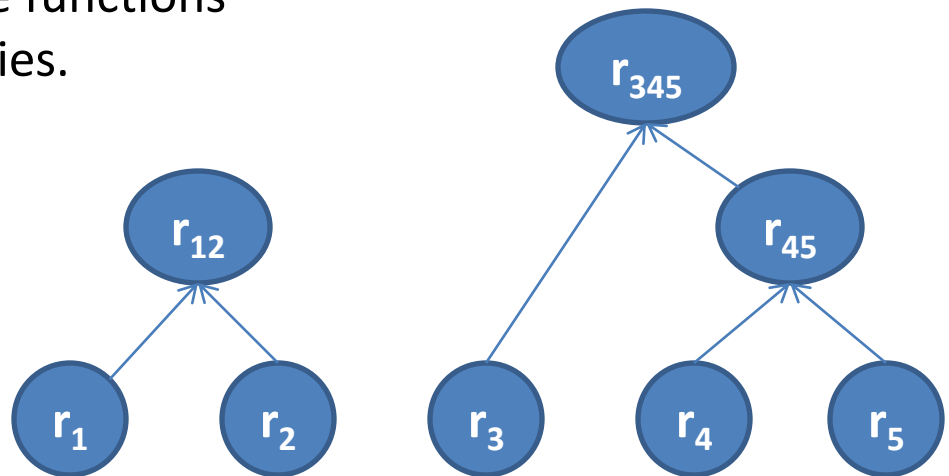
- Critically important in Web portals where users must be shown a consolidated view of the duplicate cluster.
 - Each mention only contains a subset of the attributes.
 - Mentions contain variations (of names, addresses).
 - Some of the mentions can have wrong values.

Canonicalization Algorithms

- Rule based:
 - For names: typically longest names are used.
 - For set values attributes: UNION is used.
- For strings, [Culotta et al KDD07] learn an edit distance for finding the most representative “centroid”.
- Can use “majority rule” to fix errors
(if 4 out of 5 say a business is closed, then business is closed).
 - **This may not always work due to copying [Dong et al VLDB09], or when underlying data changes [Pal et al WWW11]**

Canonicalization for Efficiency

- Stanford Entity Resolution Framework [Benjelloun VLDBJ09]
 - Consider a blackbox match and merge function
 - Match is a pairwise boolean operator
 - Merge: construct canonical version of a matching pair
- Can minimize time to compute matches by interleaving matching and merging
 - esp., when match and merge functions satisfy **monotonicity** properties.



Summary of Canonicalization

- Critically important in Web portals where users must be shown a consolidated view of the duplicate cluster.
- Canonicalization can also help speed up ER in certain cases.

PART 2

RELATIONAL & MULTIENTITY ER

Outline

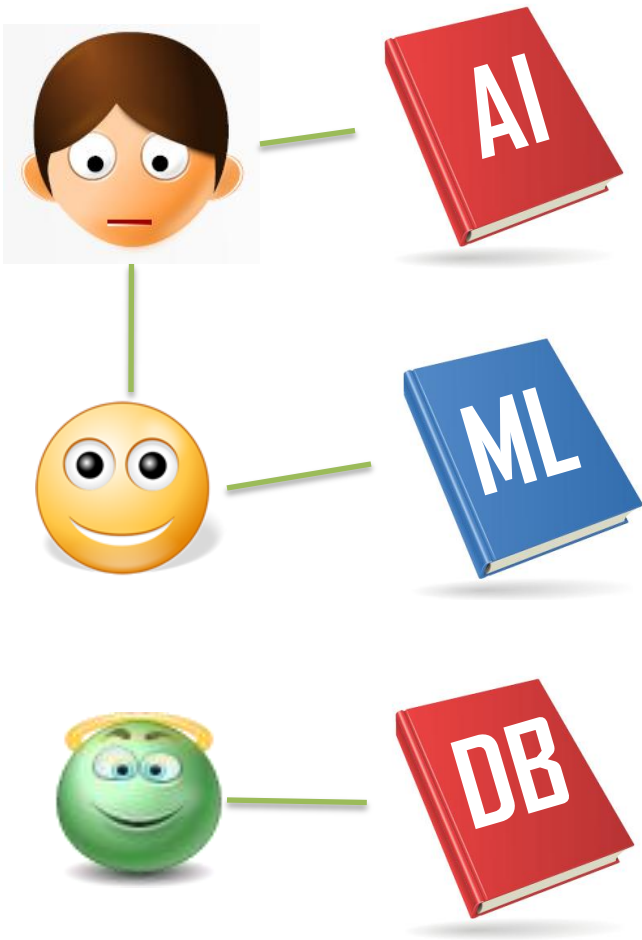
1. Classical Single Entity ER
2. Relational & MultiEntity ER
 - a) Problem Statement
 - b) Relational Features & Constraints
 - c) Non-Probabilistic Approaches: Similarity Propagation
 - d) Non-Probabilistic Approaches: Constraint Optimization
 - e) Probabilistic Approaches: Generative Models
 - f) Probabilistic Approaches: Undirected Models
3. Efficiency: Blocking/Canopy Generation
4. Challenges & Future Directions

PART 2-a

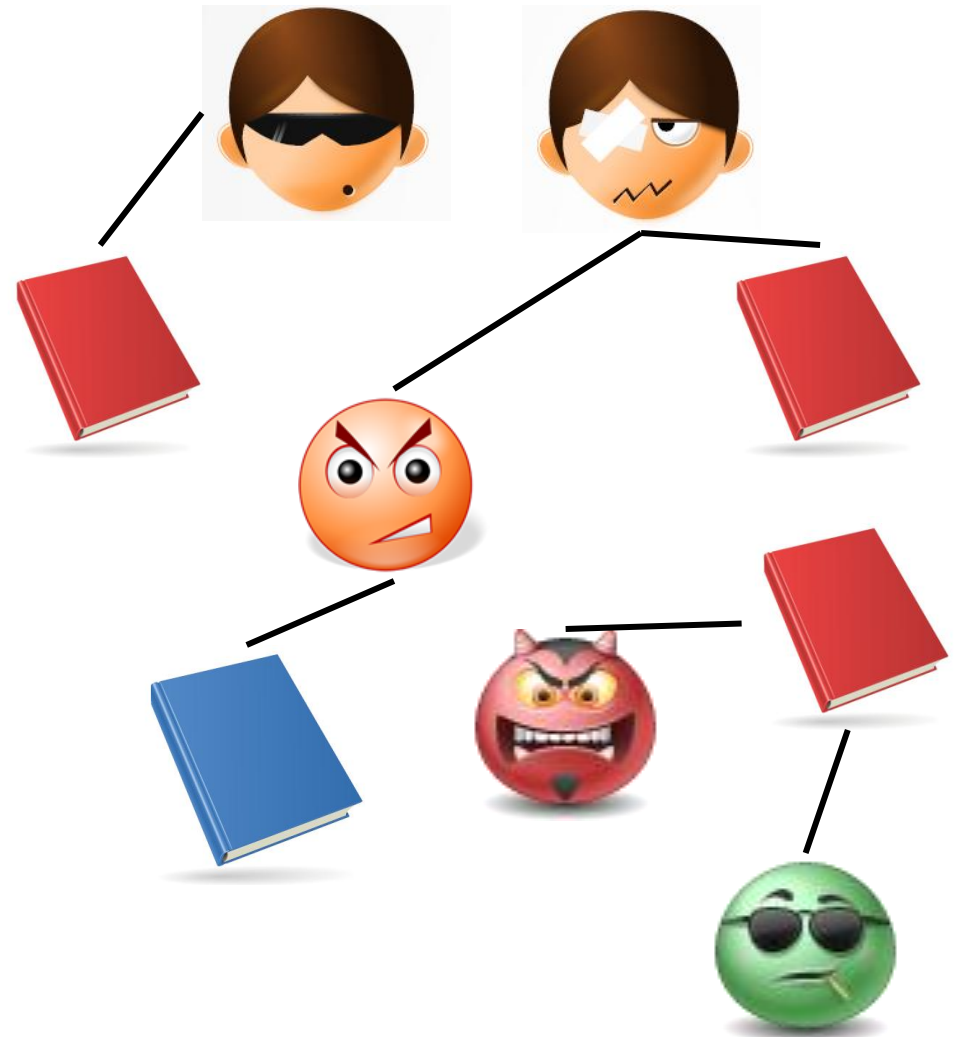
PROBLEM DEFINITION

Abstract Problem Statement

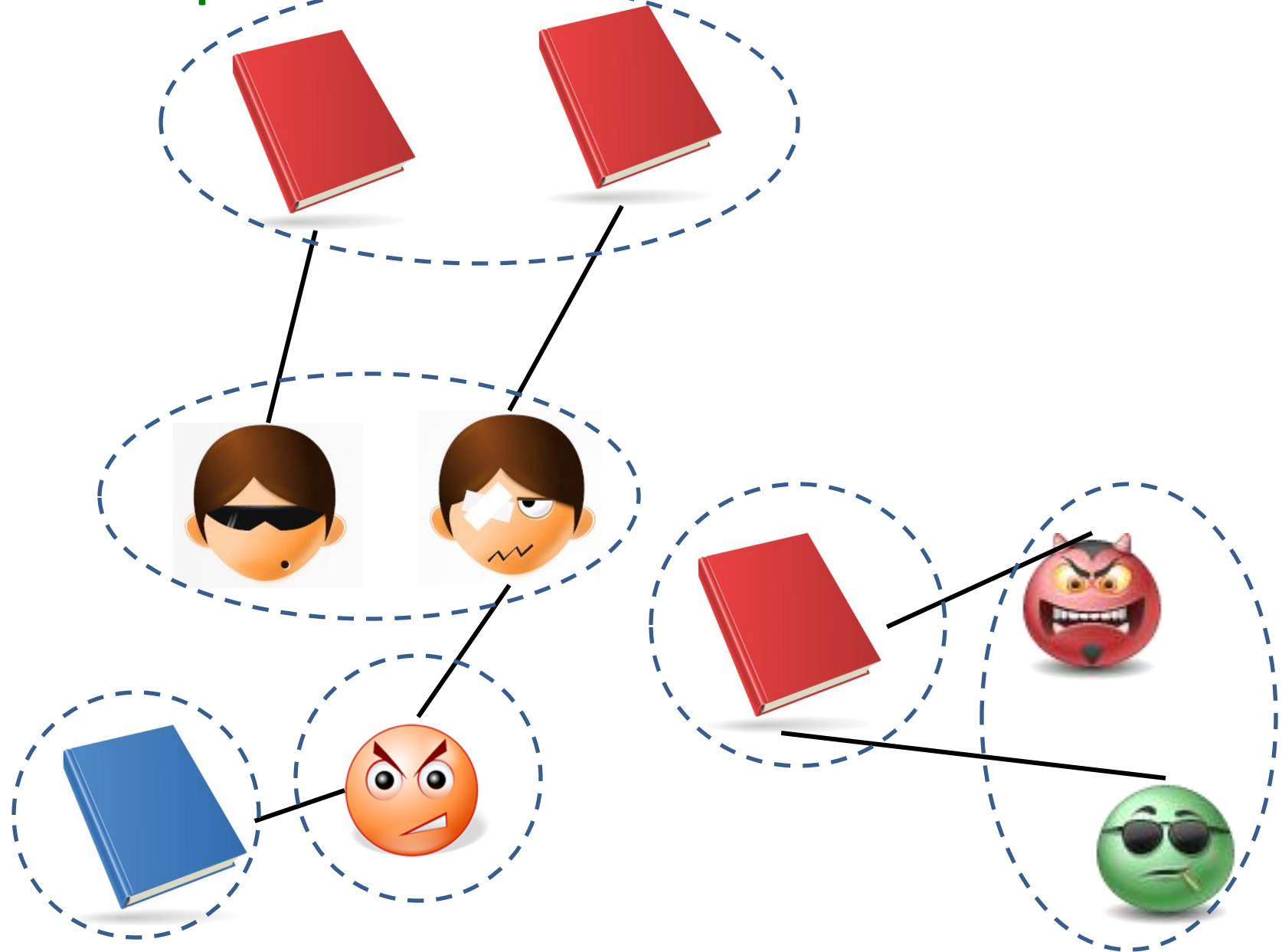
Real World



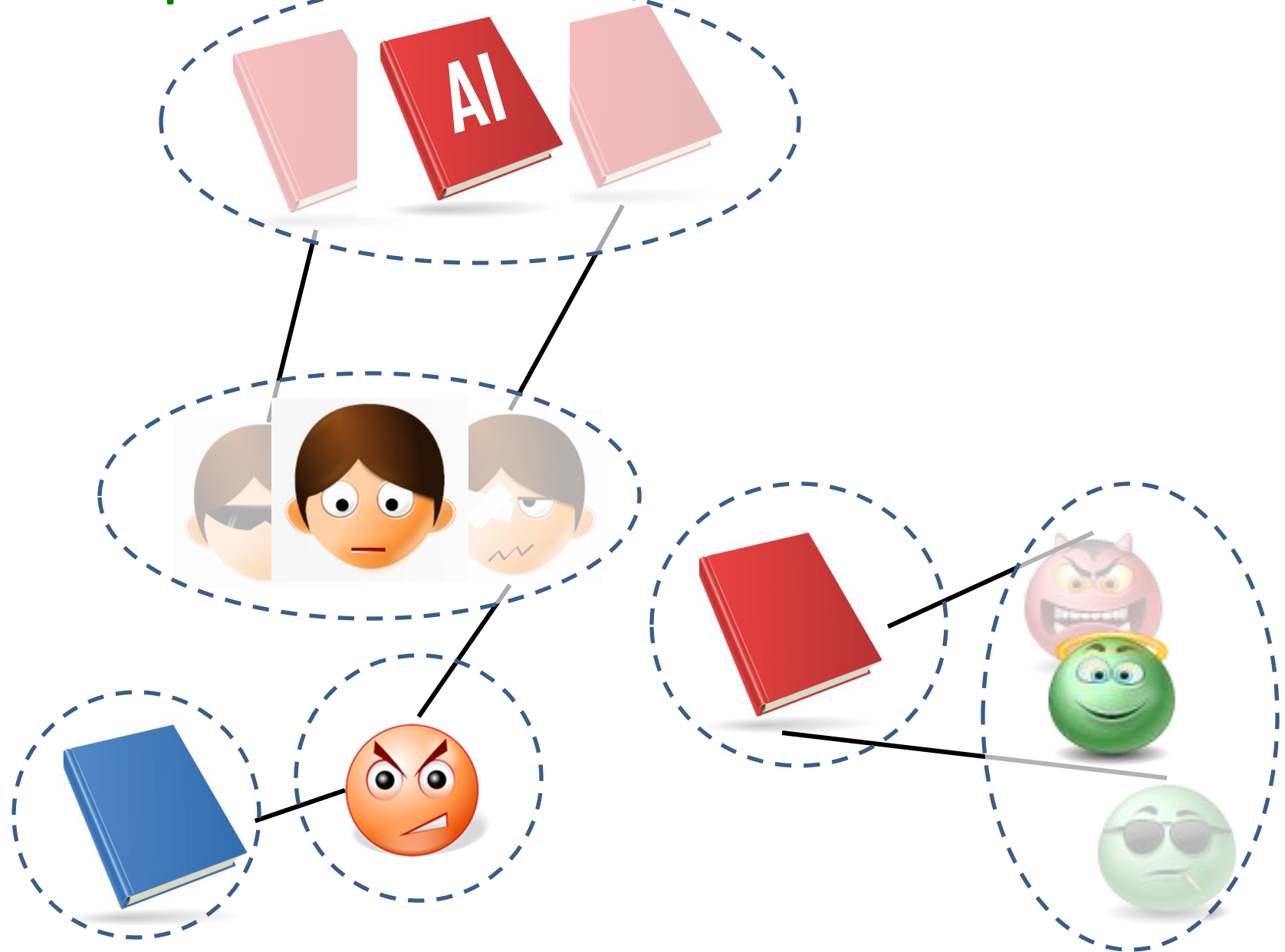
Digital World



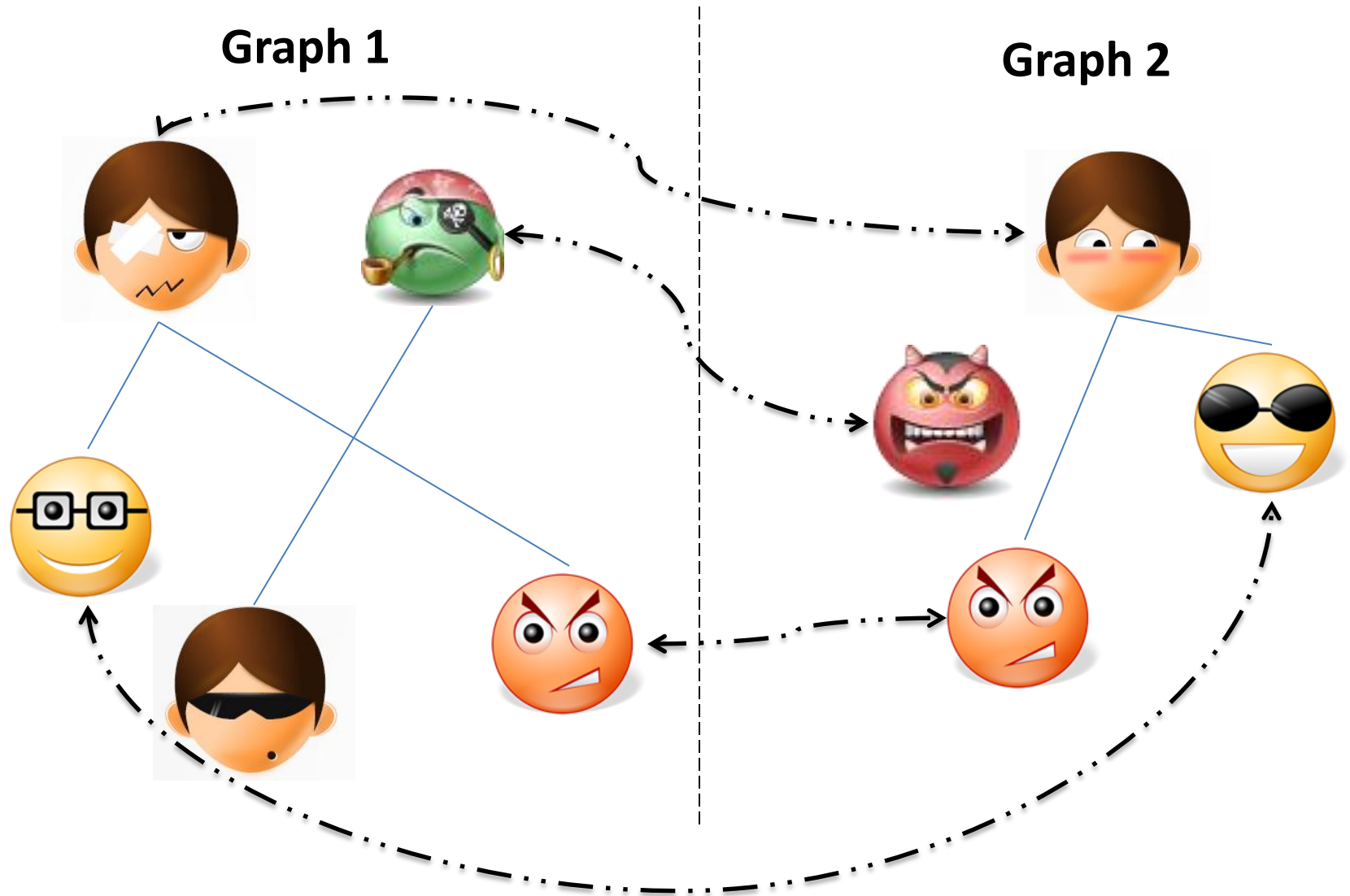
Deduplication Problem Statement



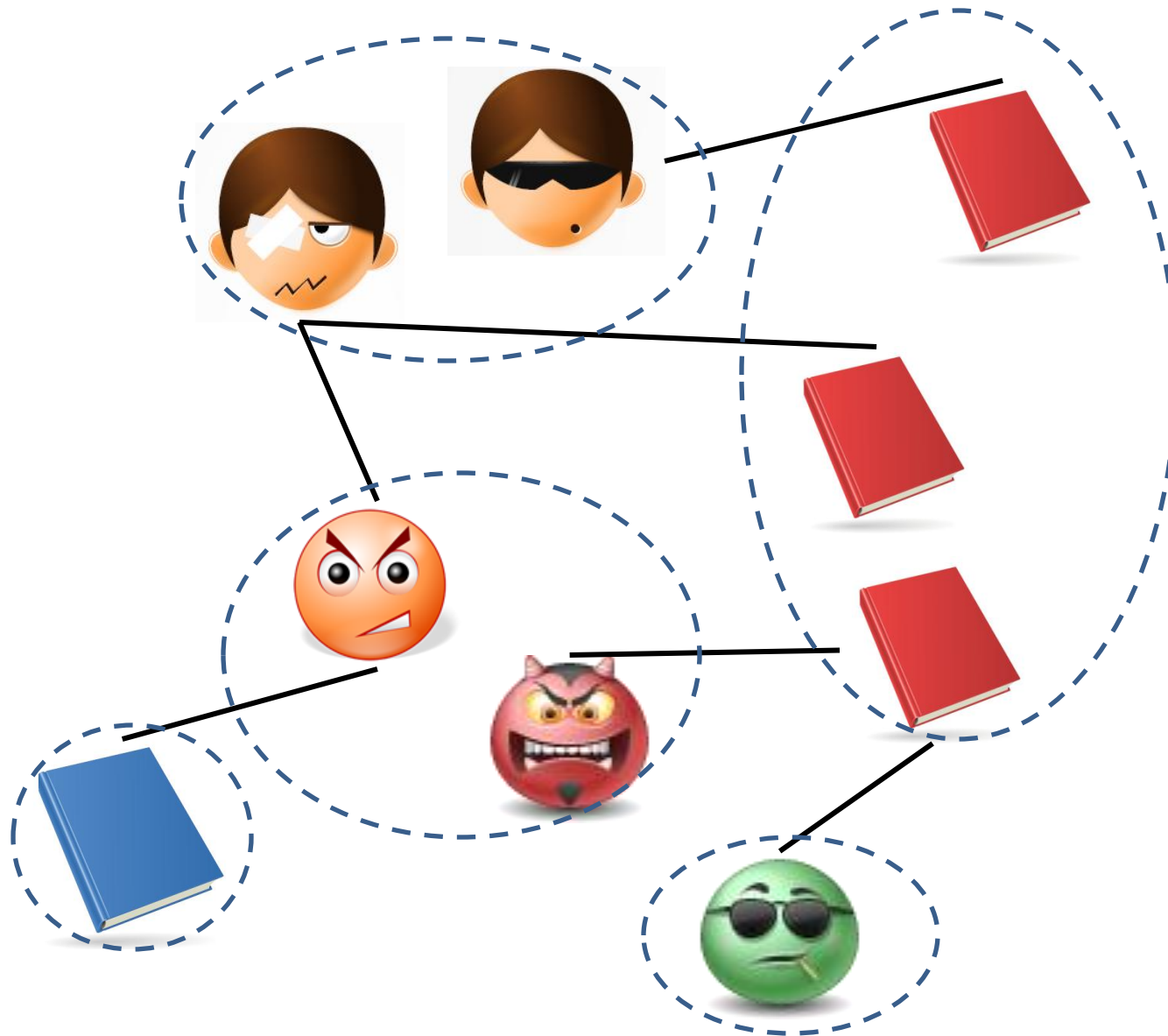
Deduplication with Canonicalization



Graph Alignment (& motif search)



Relationships are crucial



Notation & Assumptions

- R : set of records / mentions (typed)
- H : set of relations / hyperedges (typed)
- M : set of *matches* (record pairs that correspond to same entity)
- N : set of *non-matches* (record pairs corresponding to different entities)
- E : set of entities
- L : set of links

- True ($M_{true}, N_{true}, E_{true}, L_{true}$): according to real world
vs Predicted ($M_{pred}, N_{pred}, E_{pred}, L_{pred}$): by algorithm

Metrics

- Most algorithms use pairwise and cluster-based measures on each entity type
- Little work that evaluations correct prediction of links

**MOTIVATING EXAMPLE:
BIBLIOGRAPHIC DOMAIN**

Bibliography Domain

- Entities:
 - Papers
 - Authors
 - Organizations/Author Affiliations
 - Venues
 - Conference Locations
- Relations:
 - Author-Of
 - Associated-With
 - AppearsIn
 - Cites
- Co-occurrence relationships
 - Co-authors
 - Papers in same conference
 - Papers by same author
 - etc.

PART 2-b

**RELATIONAL FEATURES &
CONSTRAINTS**

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
 - This will often lead to increased precision
 - Relational information can help to distinguish references, add avoid false positives
 - It may also lead to increased recall
 - The best threshold will be different, and it may be, with the additional information, one can get increased recall as well.

Set-based Relational Features

- Relational features are often set-based
 - Set of coauthors for a paper
 - Set of cities in a country
 - Set of products manufactured by manufacturer
- Can use set similarity functions mentioned earlier
 - Common Neighbors: Intersection size
 - Jaccard's Coefficient: Normalize by union size
 - Adar Coefficient: Weighted set similarity
- Can reason about similarity in sets of values
 - Average or Max
 - Other aggregates

Constraints

- In single entity case, we already saw two important forms of constraints:
 - Transitivity: If M1 and M2 match, M2 and M3 match, then M1 and M3 match
 - Exclusivity: If M1 matches with M2, then M3 cannot match with M2
- Transitivity is key to deduplication
- Exclusivity is key to record linkage

Relational Constraints

- In multi-relational domains, matching decisions often propagate
 - Constraints may be hard constraints
 - If M1, M2 match then M3, M4 must match
 - If two papers match, their venues match
 - If two cities match, then their countries match
 - If M1, M2 don't match then M3, M4 cannot match
 - If two venues don't match, then their papers don't match
 - Or soft constraints
 - If M1, M2 match then M3, M4 more likely to match
 - If two venues match, then their papers are more likely to match
 - If M1, M2 don't match then M3, M4 less likely to match
 - If institutions don't match, then authors less likely to match

Terminology

- **Positive evidence:** If M1, M2 match then M3, M4 match
- **Negative evidence:** If M1, M2 match then M3, M4 don't match
- When matching decisions depend on other matching decisions (in other words, matching decisions are not made independently), we refer to the approach as ***collective***

Match Propagation

- **Global:** In two papers match, then their venues match
 - This constraint can be applied to all instances of venue mentions
 - All occurrences of 'SIGMOD' can be matched to 'International Conference on Management of Data'
- **Local:** If two papers match, then their authors match
 - This constraint can only be applied locally
 - Don't want to match all occurrences of 'J. Smith' with 'Jeff Smith', only in the context of the current paper

Additional Relational Constraints

- Constraints can also encode a variety of additional forms of integrity constraints
 - Uniqueness Constraints
 - Mention M1 and M2 must refer to distinct entities
 - Coauthors are distinct
 - Count Constraints
 - Entity A can link to at most N Bs
 - Authors have at most 5 papers at any conference
- Again, these can be either hard or soft constraints

Ex. Semantic Integrity Constraints

Type	Example
Aggregate	C1 = No researcher has published more than five AAAI papers in a year
Subsumption	C2 = If a citation X from DBLP matches a citation Y in a homepage, then each author mentioned in Y matches some author mentioned in X
Neighborhood	C3 = If authors X and Y share similar names and some co-authors, they are likely to match
Incompatible	C4 = No researcher exists who has published in both HCI and numerical analysis
Layout	C5 = If two mentions in the same document share similar names, they are likely to match
Key/Uniqueness	C6 = Mentions in the PC listing of a conference is to different researchers
Ordering	C7 = If two citations match, then their authors will be matched in order
Individual	C8 = The researcher with the name “Mayssam Saria” has fewer than five mentions in DBLP (new graduate student)

COLLECTIVE APPROACHES

Collective Approaches

- Decisions for cluster-membership depends on other clusters
 - Non-probabilistic approaches
 - Similarity Propagation
 - Constraint Optimization
 - Probabilistic Models
 - Generative Models
 - Undirected Models

PART 2-c

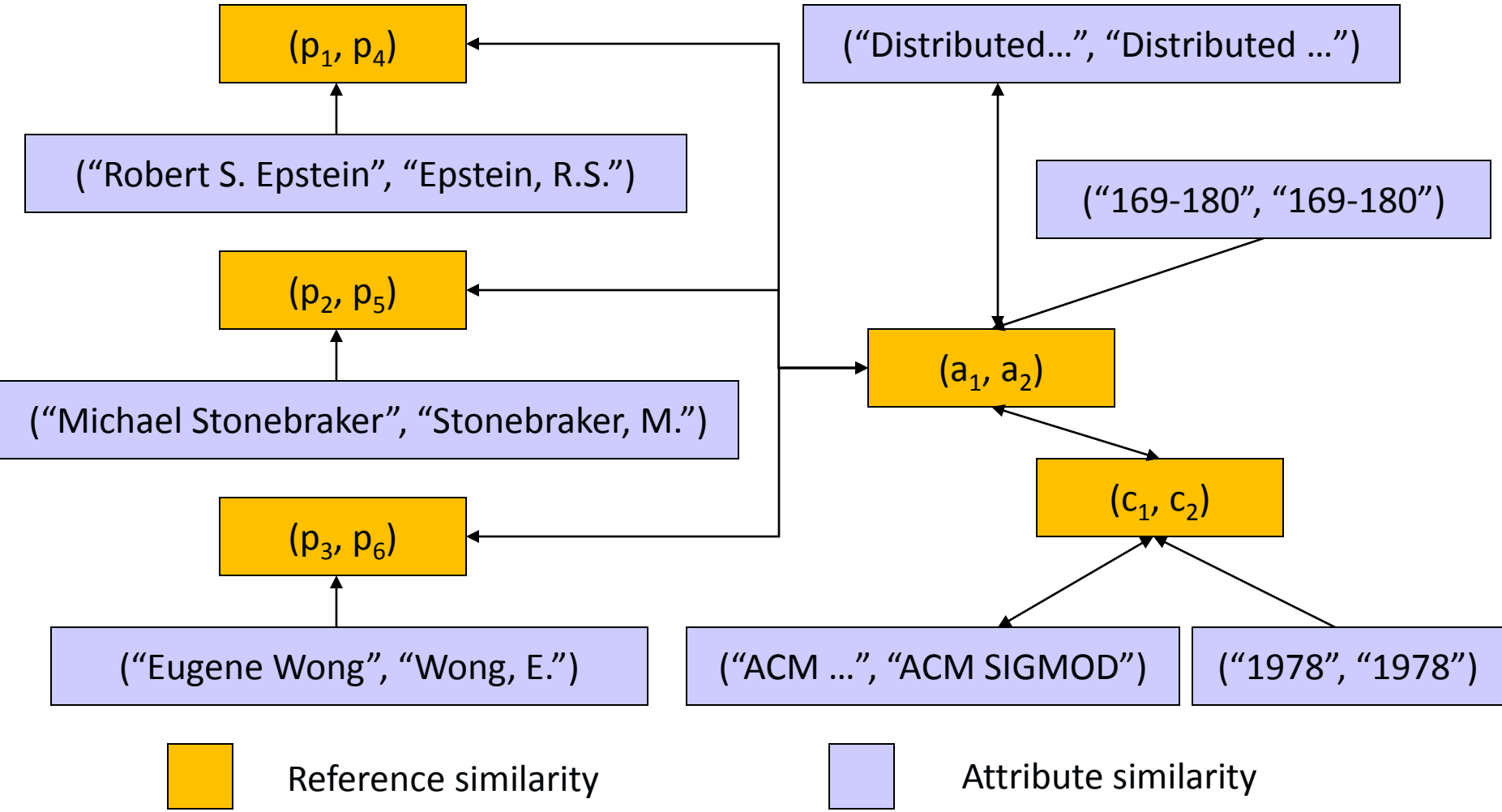
**NON-PROBABILISTIC APPROACHES:
SIMILARITY PROPAGATION**

Similarity Propagation Approaches

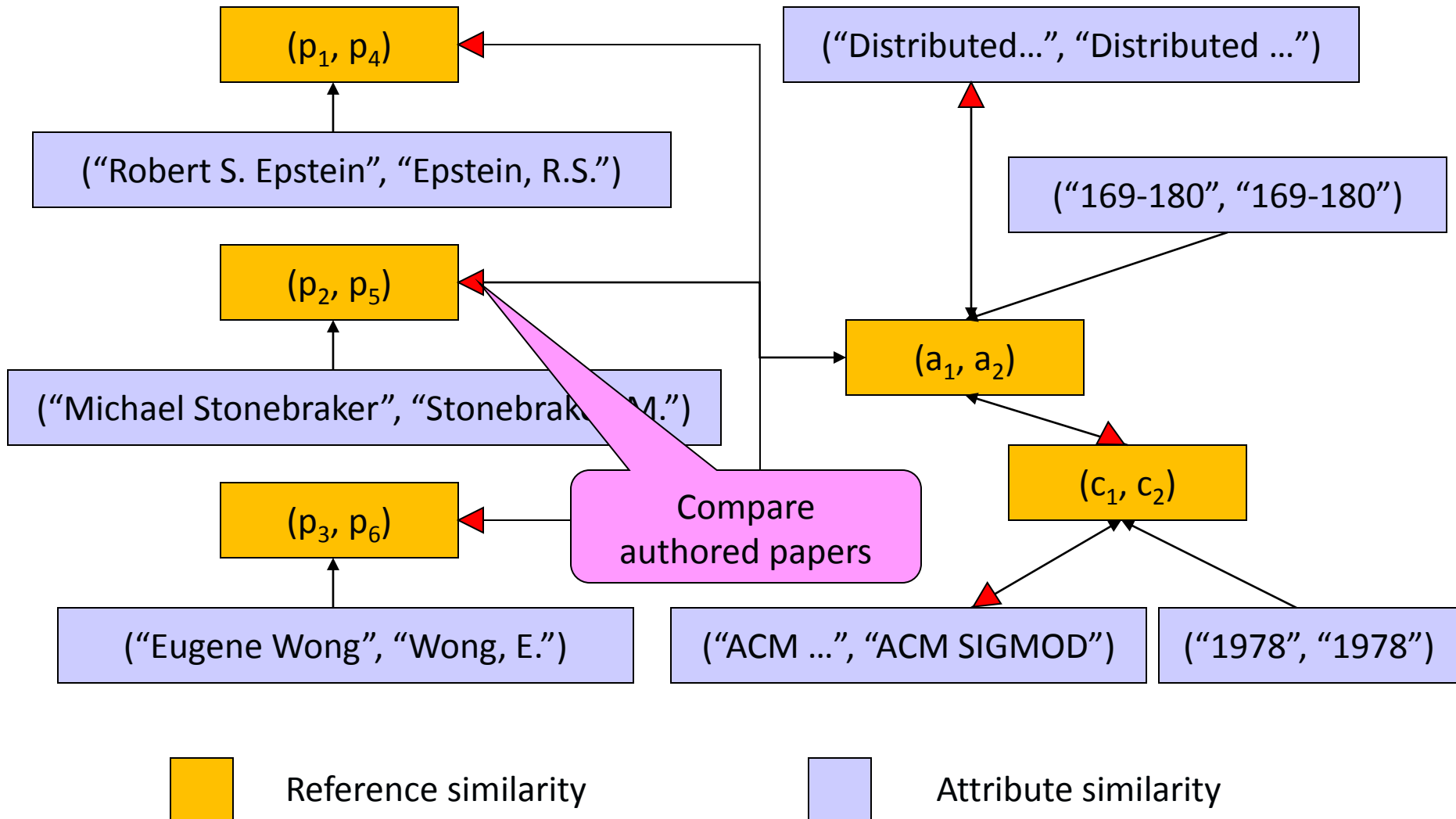
- Similarity propagation algorithms define a graph which encodes the entity mentions and matching decisions, and compute matching decisions by propagating similarity values.
 - Details of what type of graph is constructed, and how the similarity is computed varies
 - Algorithms are usually defined procedurally
 - While probabilities may be encoded in various ways in the algorithms, there is no global probabilistic model defined
- Approaches often more scalable than global probabilistic models
- Examples
 - Dependency Graphs [Dong et al, SIGMOD05]
 - Collective Relational Clustering [Bhattacharya & Getoor, TKDD07]

Dependency Graph

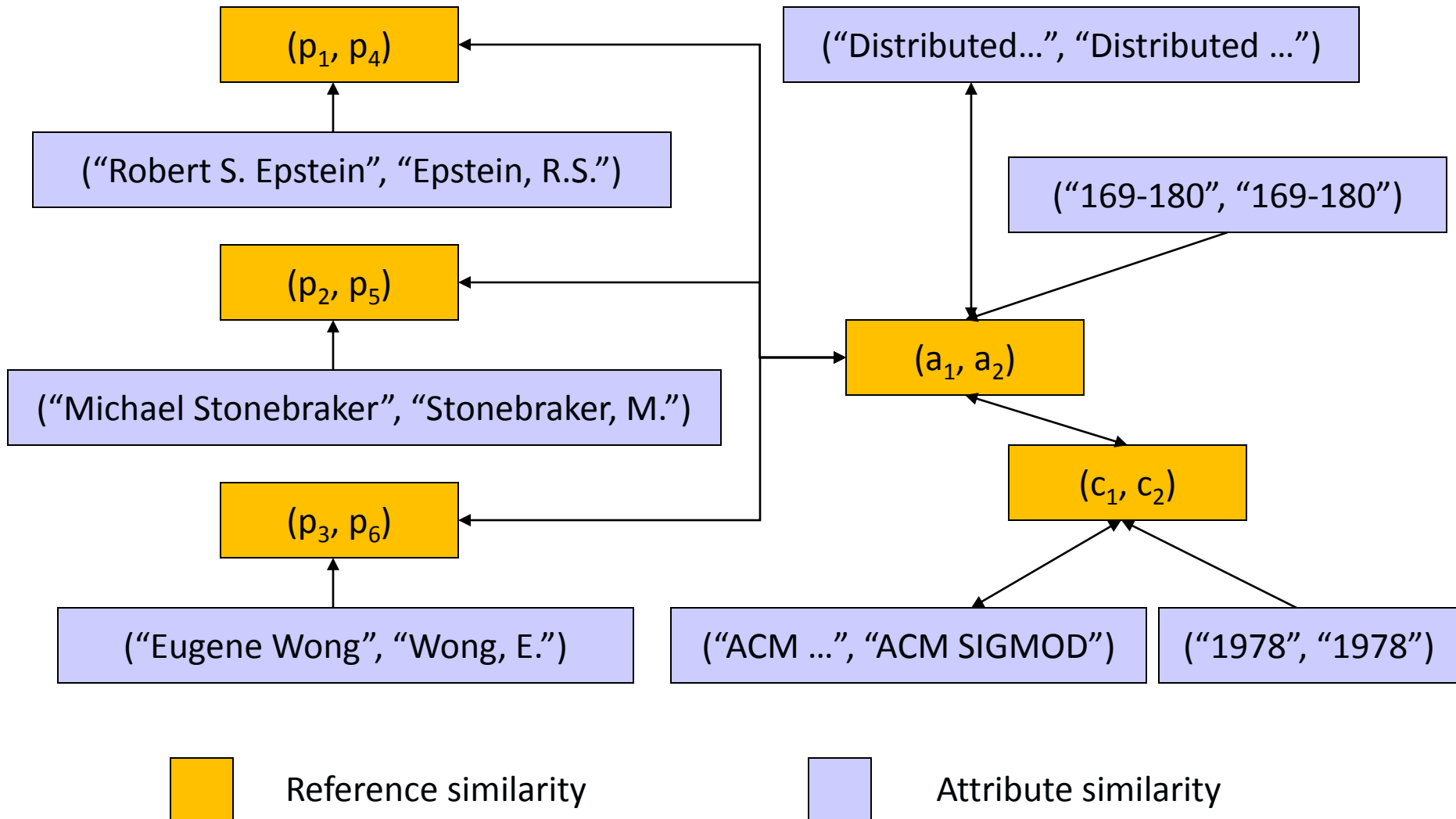
[Dong et al., SIGMOD05]



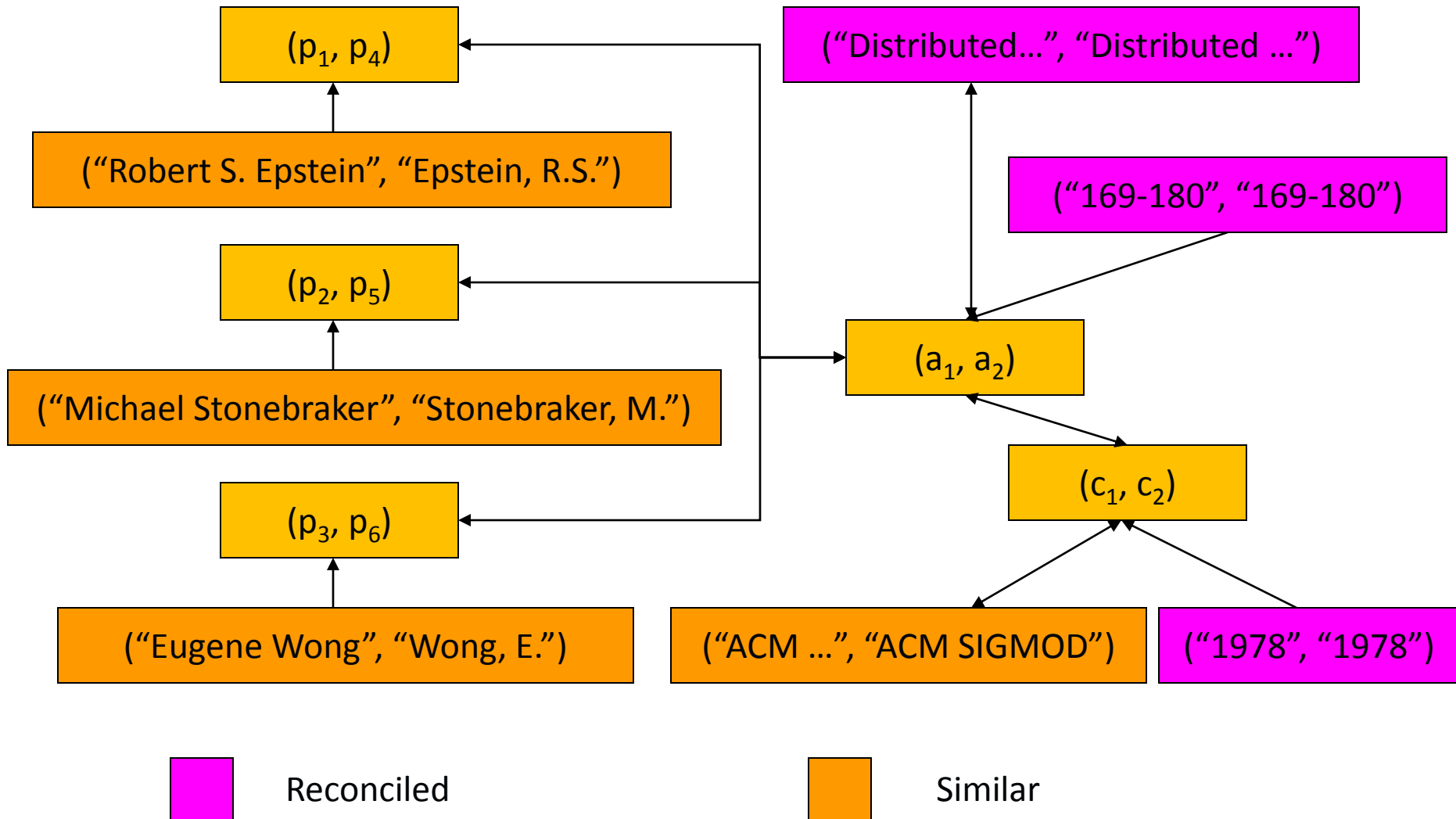
Dependency Graph Example II



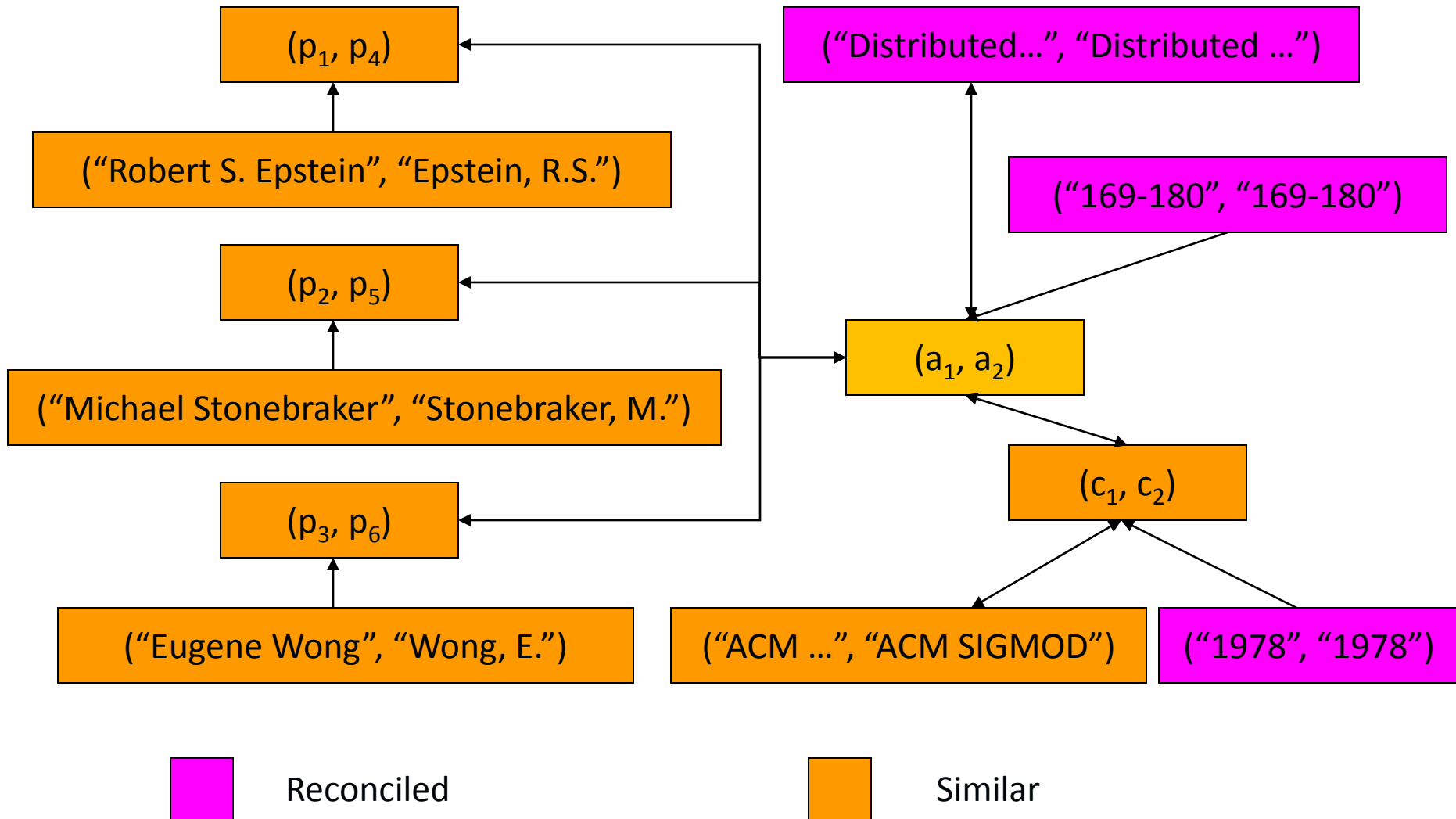
Exploit the Dependency Graph



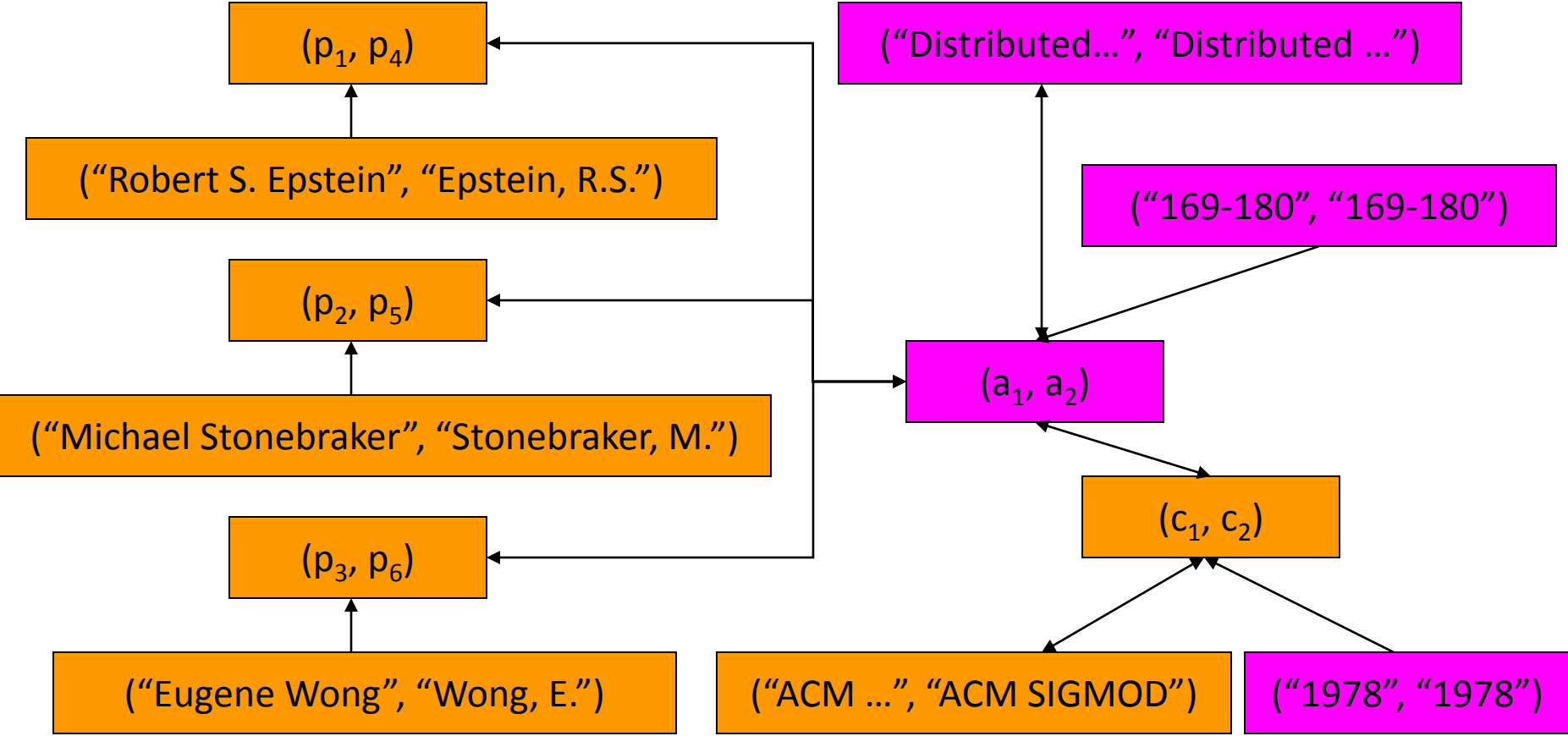
Exploit the Dependency Graph



Exploit the Dependency Graph



Exploit the Dependency Graph

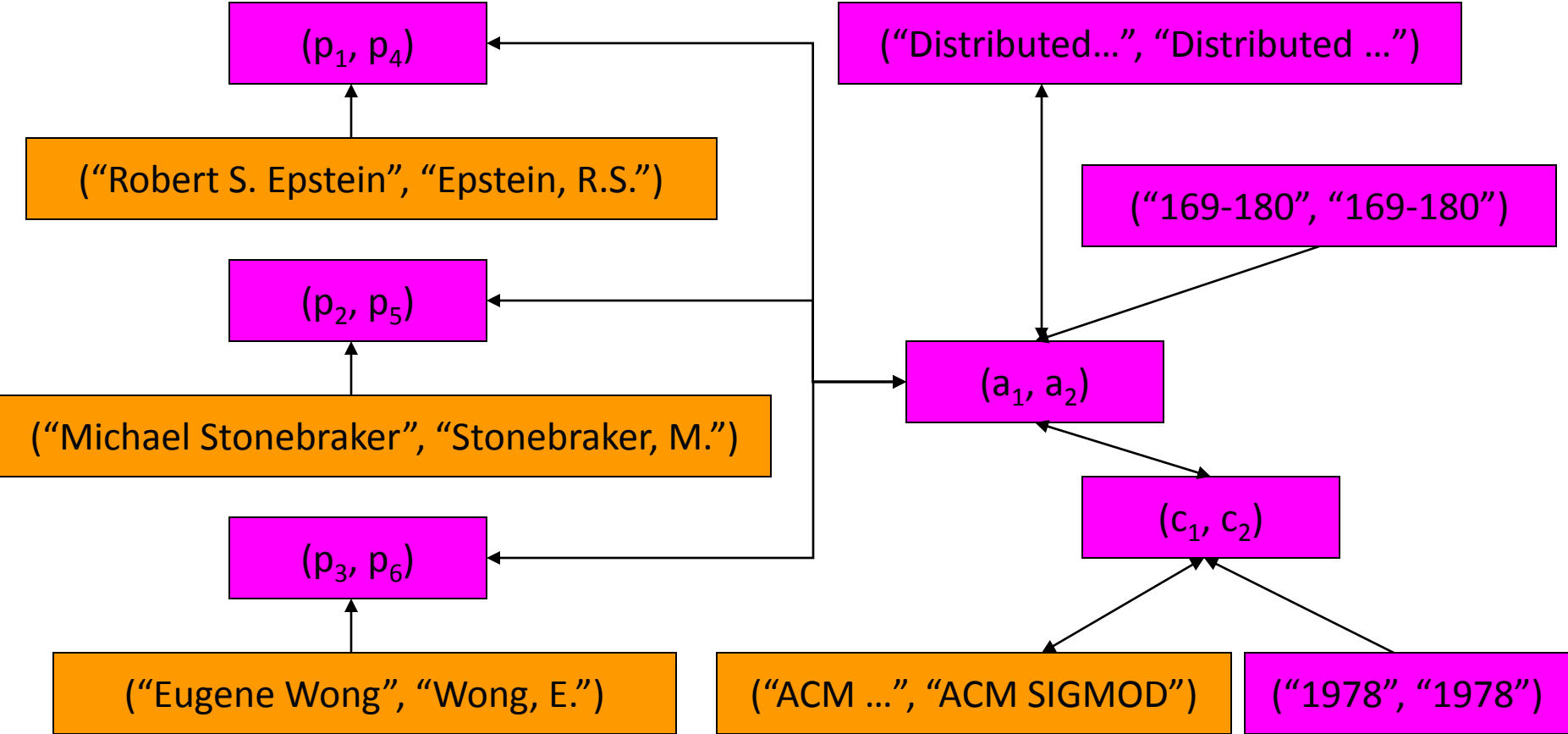


Reconciled



Similar

Exploit the Dependency Graph

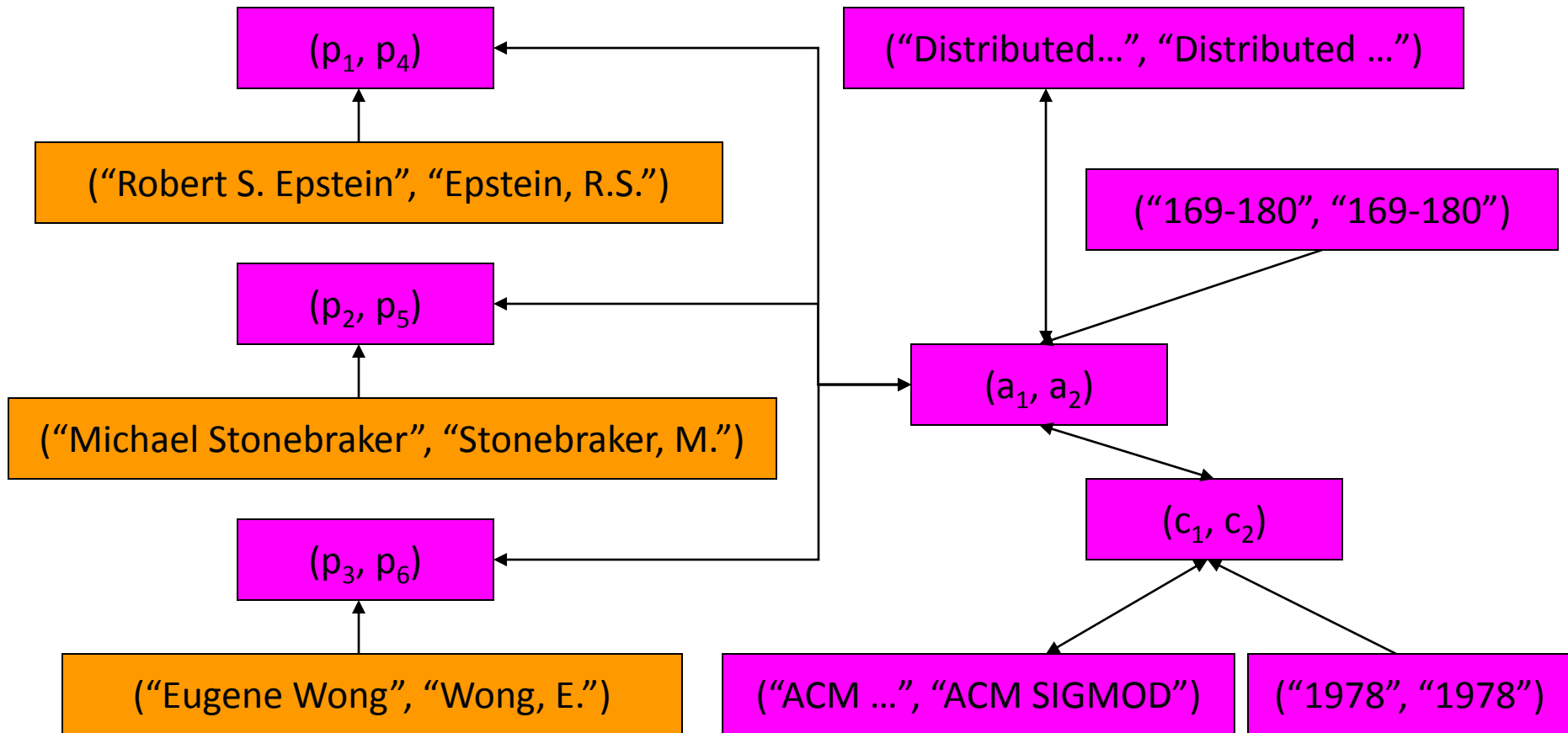


Reconciled



Similar

Exploit the Dependency Graph

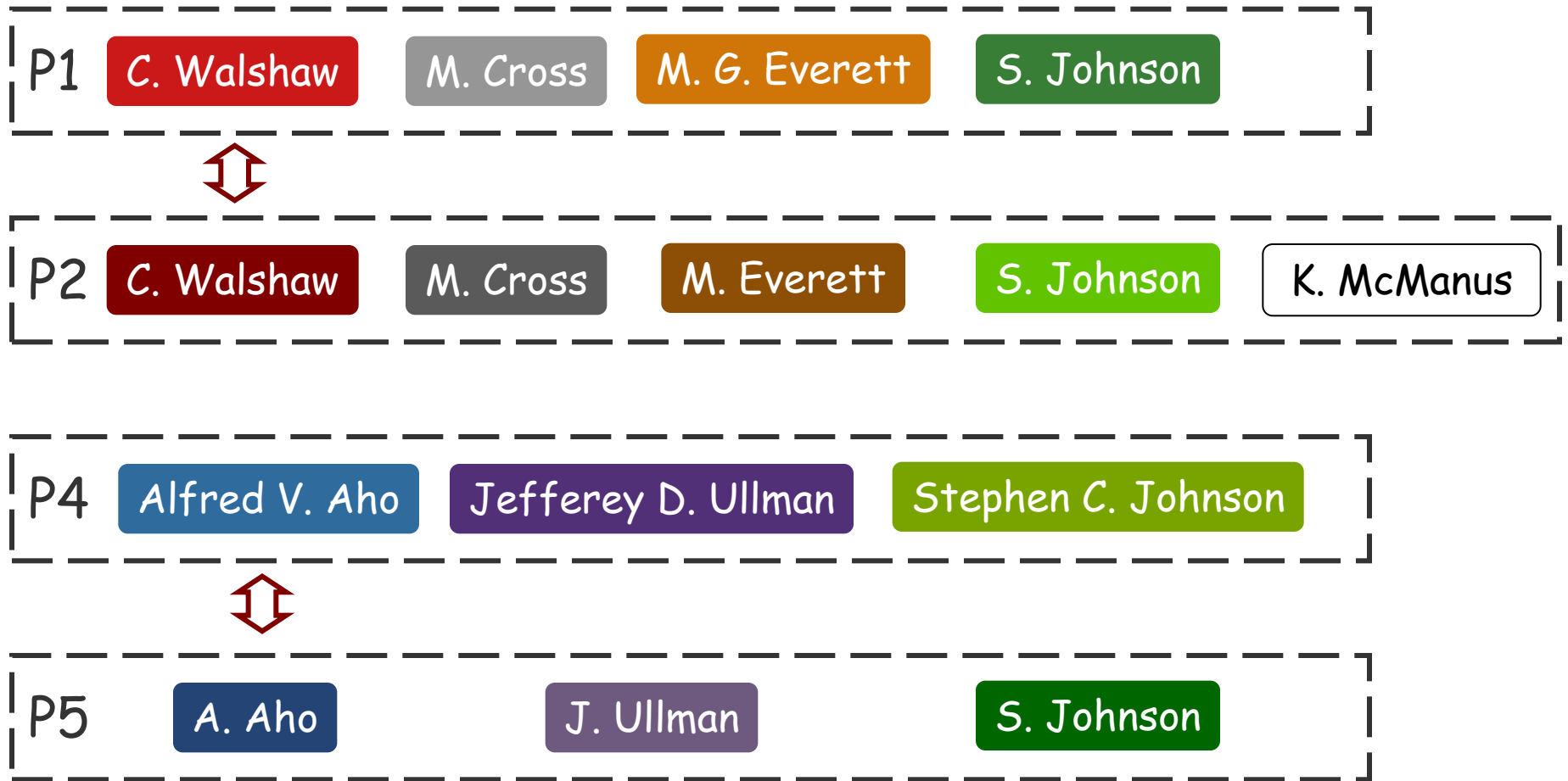


Reconciled

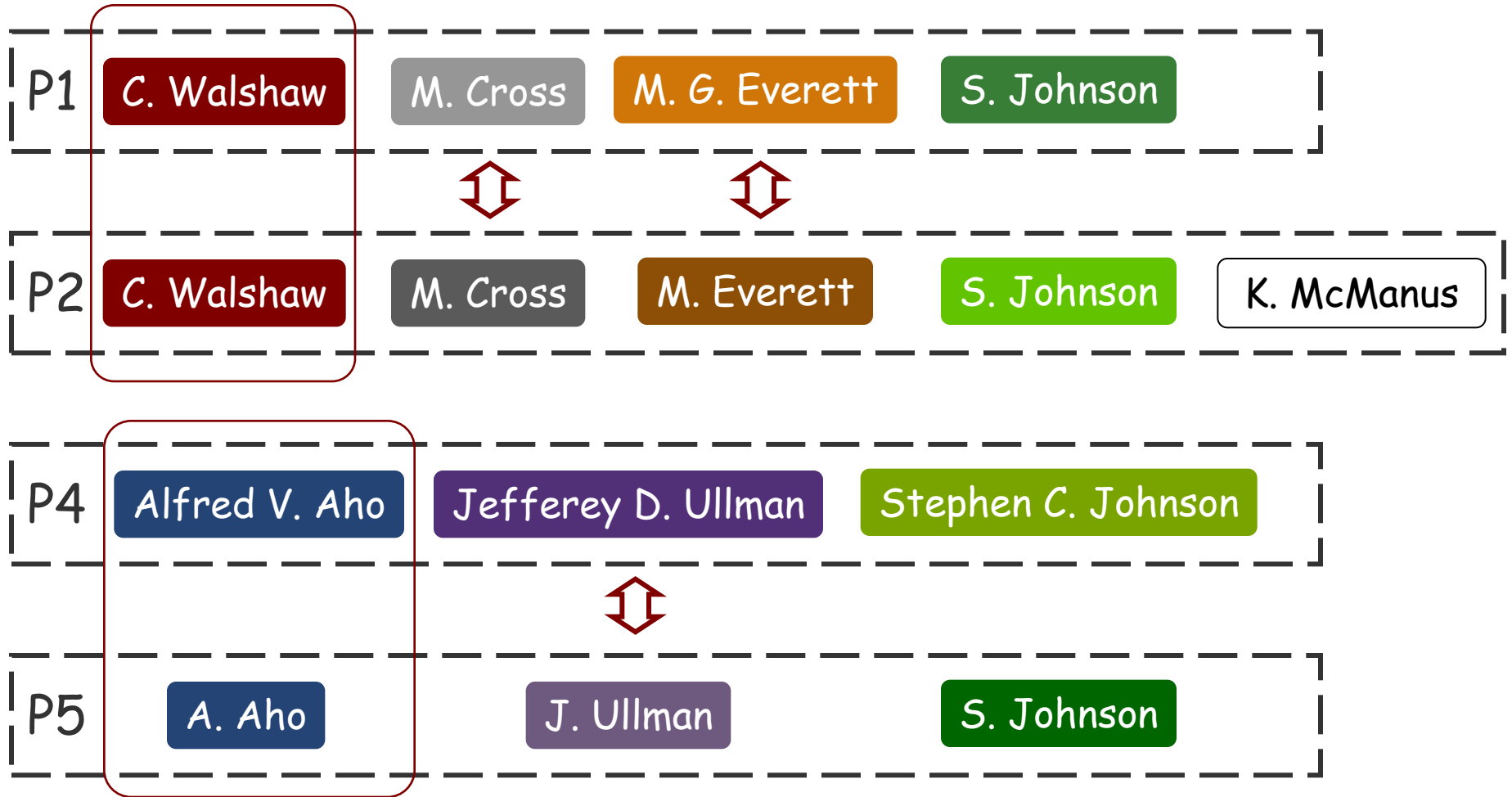


Similar

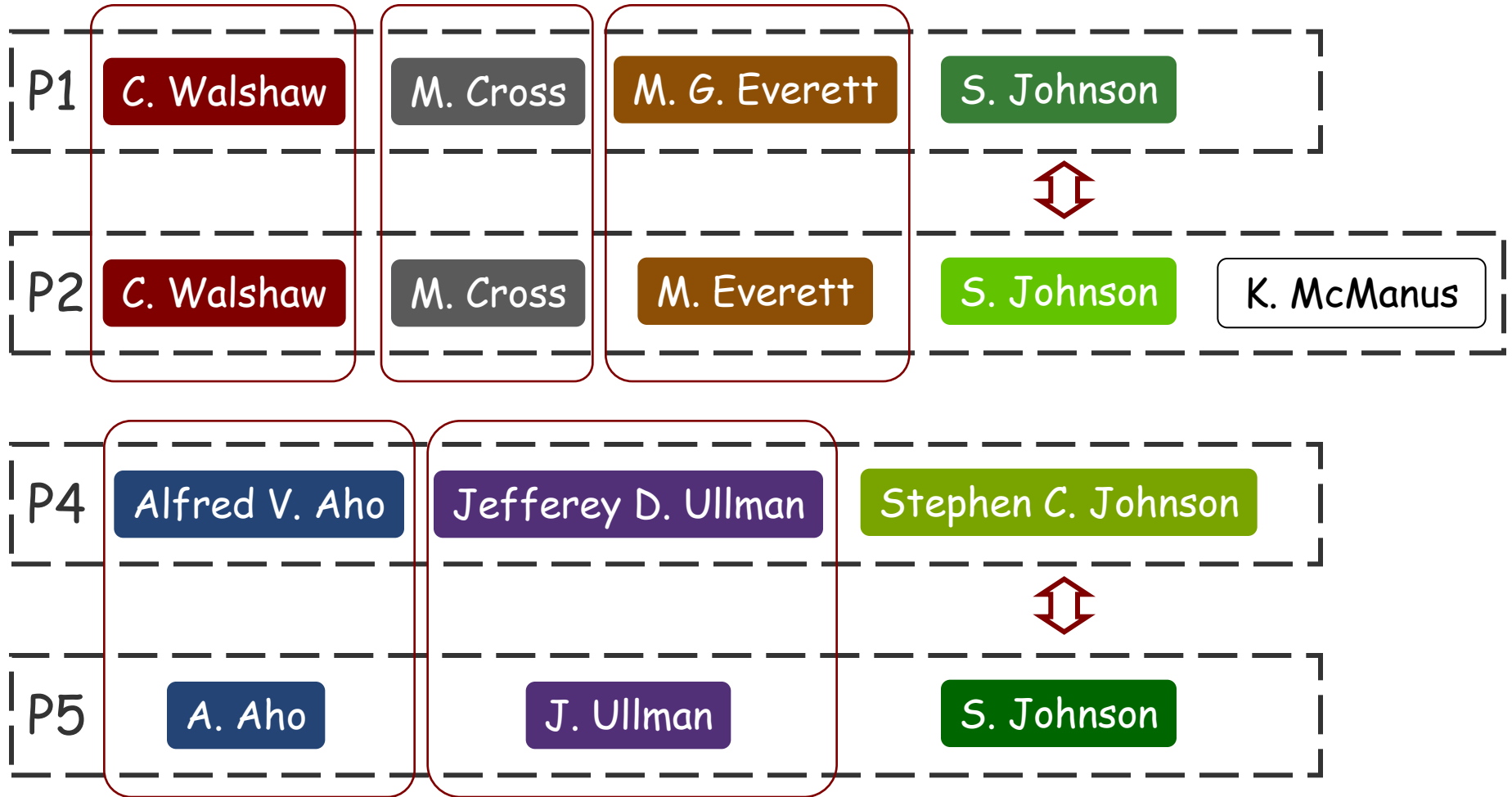
Relational Clustering for ER (RC-ER)



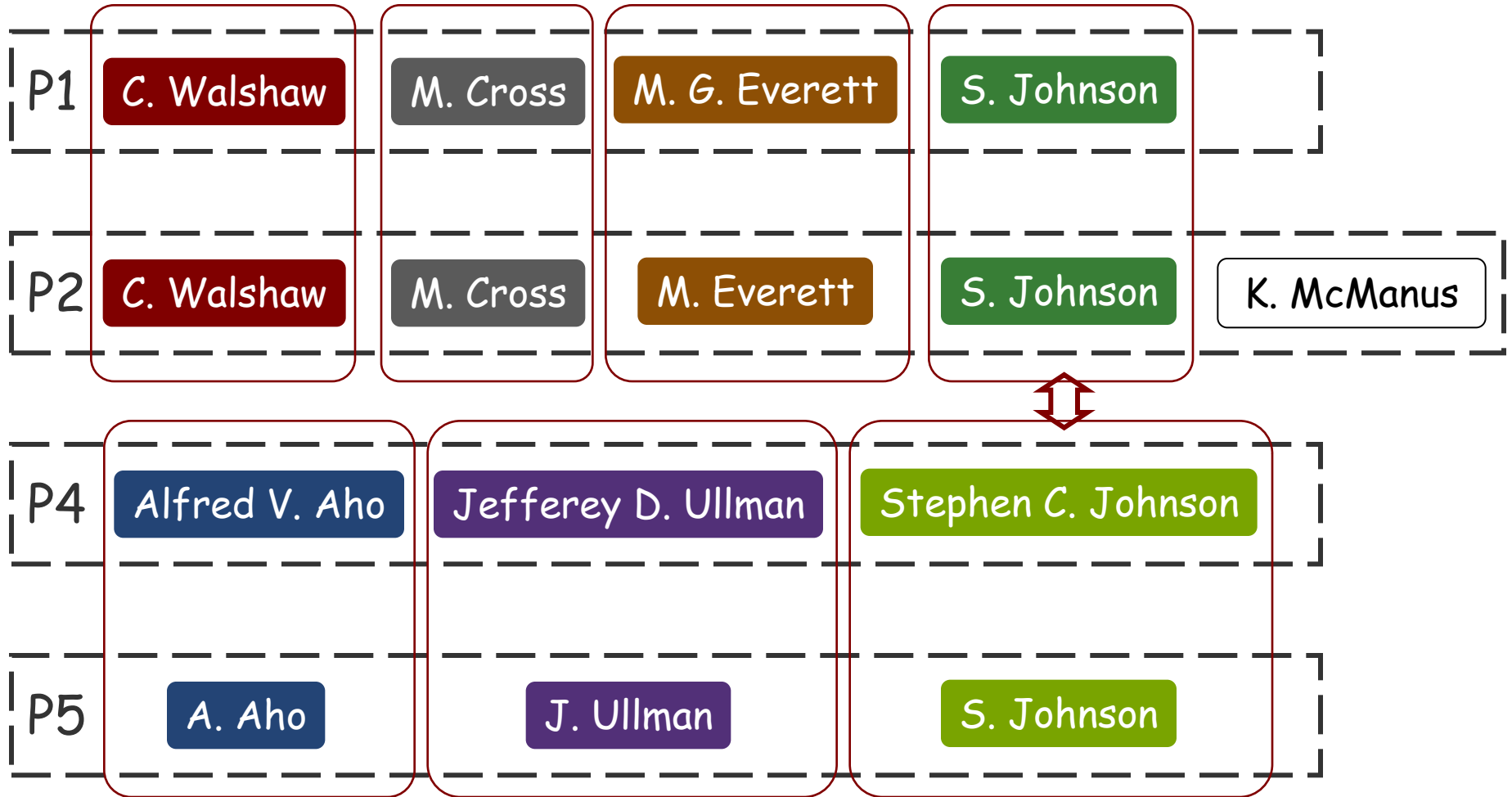
Relational Clustering for ER (RC-ER)



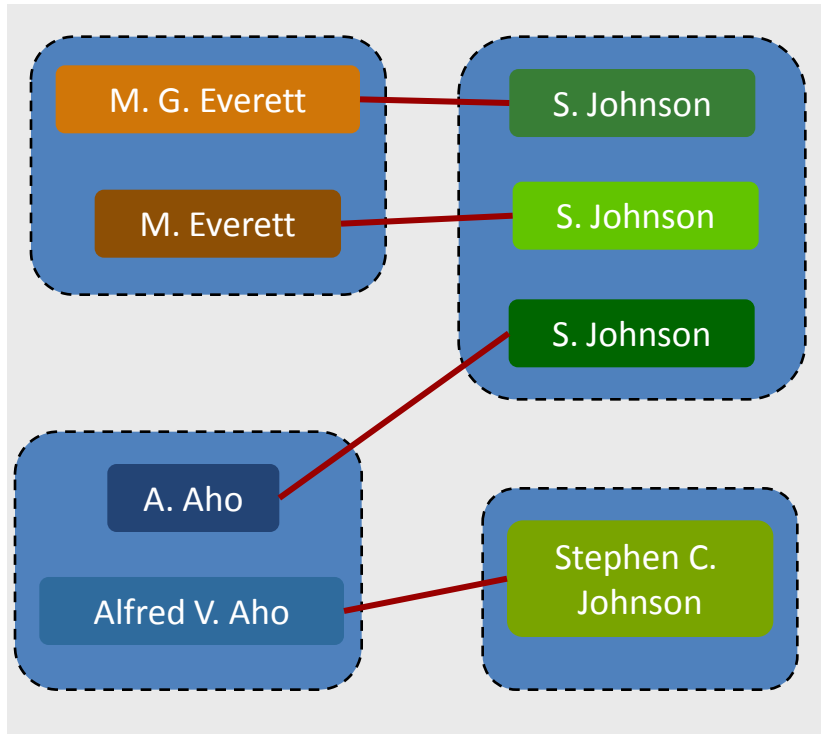
Relational Clustering for ER (RC-ER)



Relational Clustering for ER (RC-ER)



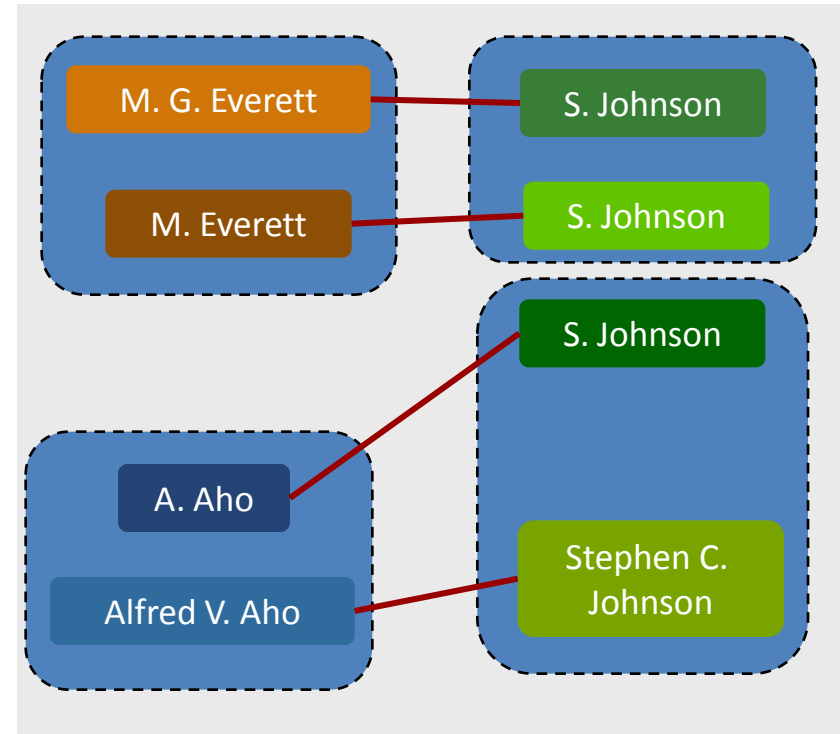
Collective Relational Clustering: Motivation



Good separation of attributes

Many cluster-cluster relationships

- Aho-Johnson1, Aho-Johnson2, Everett-Johnson1



Worse in terms of attributes

Fewer cluster-cluster relationships

- Aho-Johnson1, Everett-Johnson2

Objective Function

- Minimize:

$$\sum_i \sum_j w_A sim_A(c_i, c_j) + w_R sim_R(c_i, c_j)$$

weight for
attributes

similarity of
attributes

weight for
relations

Similarity based on relational edges
between c_i and c_j

- Greedy clustering algorithm: merge cluster pair with max reduction in objective function

$$\Delta(c_i, c_j) = w_A sim_A(c_i, c_j) + w_R (|N(c_i) \cap N(c_j)|)$$

Similarity of attributes

Common cluster neighborhood

Similarity Measures

- Attribute Similarity
 - Use best available measure for each attribute
 - Name Strings: *Soft TF-IDF, Levestein, Jaro*
 - Textual Attributes: *TF-IDF*
- Aggregate to find similarity between clusters
 - Single link, Average link, Complete link
 - Cluster representative
- Relational Similarity
 - Measures of set similarity
 - Higher order similarity: **Consider nbrs of nbrs**
 - Can also consider neighborhood as multi-set

Relational Clustering Algorithm

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

Similarity-propagation Approaches

	Method	Notes	Constraints	Evaluation
ReIDC [Kalashnikov et al, TODS06]	Reference disambiguation using using Relationship-based data cleaning (ReIDC)	Model choice nodes identified using feature-based similarity	Context attraction measures the relational similarity	Accuracy and runtime for Author resolution and director resolution in Movie database
Reference Reconciliation [Dong et al, SIGMOD05]	Dependency Graph for propagating similarities + enforce non-match constraints	Reference enrichment Explicitly handle missing values Parameters set by hand	Both positive and negative constraints	Precision/Recall, F1 on personal information management data (PIM), Cora dataset
Collective Relational Clustering [Bhattacharya & Getoor, TKDD07]	Modified hierarchical agglomerative clustering approach	Constructs canonical entity as merges are made	Focus on coauthor resolution and propagation	F1 on three bibliographic datasets: CiteSeer, ArXiv, and BioBase

PART 2-d

**CONSTRAINT OPTIMIZATION
APPROACHES**

Constraint-based Approaches

- Constraint-based approaches explicitly encode relational constraints
 - They can be formulated as hybrid of constraints and probabilistic models
 - Or as constraint optimization problem
- Examples
 - Constraint-based Entity Matching [Shen, Li & Doan, AAAI05]
 - Dedupalog [Arasu, Re, Suciu, ICDE09]

CME

- Two layer model:
 - Layer 1: Generative model for data sets that satisfy constraints; builds on (Li, Morie, & Roth, AI Mag 2004).
 - Layer 2: EM algorithm and the relaxation labeling algorithm to perform matching. Matching process is carried out in multiple iterations. In each iteration, use EM to estimate parameters of the generative model and a matching assignment, then employs relaxation labeling to exploit the constraints
- First layer clusters mentions into groups (such that all matching mentions belong to the same group) and exploits constraints at the *group level*. *Once this is done*, the second layer exploits additional constraints at the level of *individual matching mention pairs*.

Clustering with Dedupalog

PaperRef(id, title, conference, publisher, year)
Wrote(id, authorName, Position)

Data to be
deduplicated

TitleSimilar(title1,title2)
AuthorSimilar(author1,author2)

(Thresholded) Fuzzy-
Join Output

Step (0) Create Fuzzy Matches; this is input to Dedupalog.

Step (1) Declare the entities

“Cluster Papers, Publishers, & Authors”

Paper!(id) :- PaperRef(id,-,-,-)
Publisher!(p) :- PaperRef(-,-,-,p,-)
Author!(a) :- Wrote(-,a,-)

Dedupalog is *flexible*:
Unique Names Assumption (UNA**)**

Publishers (UNA) and Papers (NOT UNA)

Step (2) Declare Clusters

Input in the DB

PaperRef(id, title, conference, publisher, year)
Wrote(id, authorName, Position)

*“Cluster papers,
publishers, and authors”*

TitleSimilar(title1,title2)
AuthorSimilar(author1,author2)

Paper!(id) :- PaperRef(id,-,-,-)
Publisher!(p) :- PaperRef(-,-,-,p,-)
Author!(a) :- Wrote(-,a,-)

Clusters are *declared* using * (like IDBs or Views): These are output

Author*(a_1, a_2) \leftrightarrow AuthorSimilar(a_1, a_2)

Author1	Author2
AA	Arvind Arasu
Arvind A	Arvind Arasu

“Cluster authors with similar names”

*IDBs are **equivalence relations**:
Symmetric, Reflexive , & Transitively-
Closed Relations: i.e., *Clusters*

A Dedupalog program is a
set of datalog-like rules

Simple Constraints

“Papers with similar titles should likely be clustered together”

Paper*(id₁,id₂) \leftrightarrow PaperRef(id₁,t₁,-), PaperRef(id₂,t₂,-), TitleSimilar(t₁,t₂)

Author*(a₁,a₂) \leftrightarrow AuthorSimilar(a₁,a₂)

(\leftrightarrow) Soft-constraints:
Pay a cost if violated.

Paper*(id₁,id₂) \leq PaperEq(id₁,id₂)

\neg **Paper***(id₁,id₂) \leq PaperNeq(id₁,id₂)

(\leq) Hard-constraints: *Any clustering must satisfy these*

*“Papers in PaperEQ **must** be clustered together, those in PaperNEQ **must not** be clustered together”*

Hard constraints
are challenging!

1. PaperEQ, PaperNEQ are relations (EDBS)
2. \neg denotes Negation here.

Advanced Constraints

“Clustering two papers, then must cluster their first authors”

Author*(a₁,a₂) <= **Paper***(id₁,id₂), Wrote(id₁,a₁,1), Wrote(id₂,a₂,1)

“Clustering two papers makes it likely we should cluster their publisher”

Publisher*(x,y) <- Publishes(x,p₁), Publishes(x,p₂), **Paper***(p₁,p₂)

“if two authors do not share coauthors, then do not cluster them”

\neg **Author***(x, y) <- \neg (Wrote(x, p₁, -), Wrote(y, p₂, -), Wrote(z, p₁, -),
Wrote(z, p₂, -), **Author***(x, y))


Dedupalog via CC

Semantics: Translate a Dedupalog Program to a set of graphs

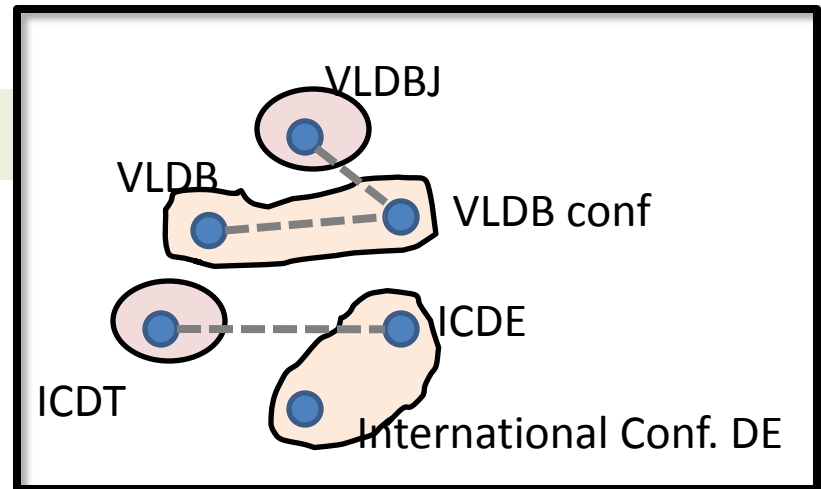
Nodes are references (in the ! Relation)

Entity References: Conference!(c)

Conference^{*}(c₁,c₂) <-> ConfSim(c₁,c₂)

 Positive edges

 Negative edges are implicit



For a single graph w.o. hard constraints
we can reuse prior work for O(1) apx.

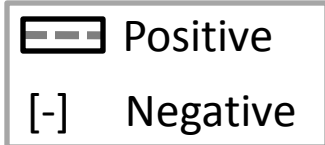
Correlation Clustering

Conference $^*(c_1, c_2) \leftarrow \text{ConfSim}(c_1, c_2)$

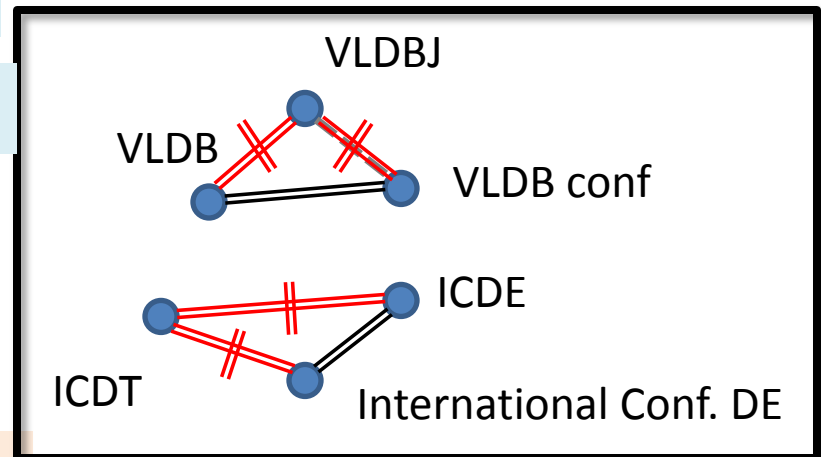
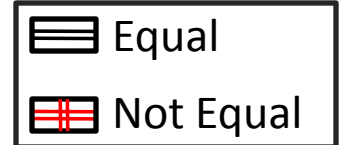
Conference $^*(c_1, c_2) \leq \text{ConfEQ}(c_1, c_2)$

Conference $^*(c_1, c_2) \leq \text{ConfNEQ}(c_1, c_2)$

Soft



Hard



1. Pick a random order of edges
2. **While** there is a soft edge **do**
 1. Pick first soft edge in order
 2. If turn into
 3. Else is [-] turn into
 4. Deduce labels
3. **Return** Transitively closed subsets

Simple, Combinatorial
algorithm is easy to scale!

Thm: This is a 3-*apx*!

Voting

Extend algorithm to **whole** language via *voting technique*.
Support many entities, recursive programs, etc.

Many dedupalog programs
have an $O(1)$ -apx

Thm: A recursive-hard
constraints no $O(1)$ apx

Thm: All “soft” programs $O(1)$

Expert: multiway-cut hard

System properties:

- (1) Streaming algorithm
- (2) linear in # of matches (not n^2)
- (3) User interaction

Features: Support for weights, reference tables
(partially), and corresponding hardness results.

PART 4-d

**PROBABILISTIC MODELS:
GENERATIVE APPROACHES**

Generative Probabilistic Approaches

- Probabilistic semantics based on Directed Models
 - Advantage: generative semantics, can “generate” new instances
 - Disadvantage: acyclicity requirement
- Variety of approaches
 - Based on Bayesian Network semantics, Latent Dirichlet Allocation, etc.
- Examples
 - Latent Dirichlet Allocation [Bhattacharya & Getoor, SDM07]
 - Probabilistic Relational Models [Pasula et al, NIPS02]

LDA-ER Probabilistic Generative Model

- Model how entity references co-occur in data
 1. Generation of references from entities
 2. Relationships between underlying entities
 - Groups of entities instead of pair-wise relations

Discovering Groups from Relations

Stephen P Johnson

Chris Walshaw

Kevin McManus

Mark Cross

Martin Everett

Parallel Processing Research Group



P1: C. Walshaw, M. Cross, M. G. Everett,
S. Johnson

P2: C. Walshaw, M. Cross, M. G. Everett,
S. Johnson, K. McManus

P3: C. Walshaw, M. Cross, M. G. Everett

Stephen C Johnson

Alfred V Aho

Ravi Sethi

Jeffrey D Ullman

Bell Labs Group

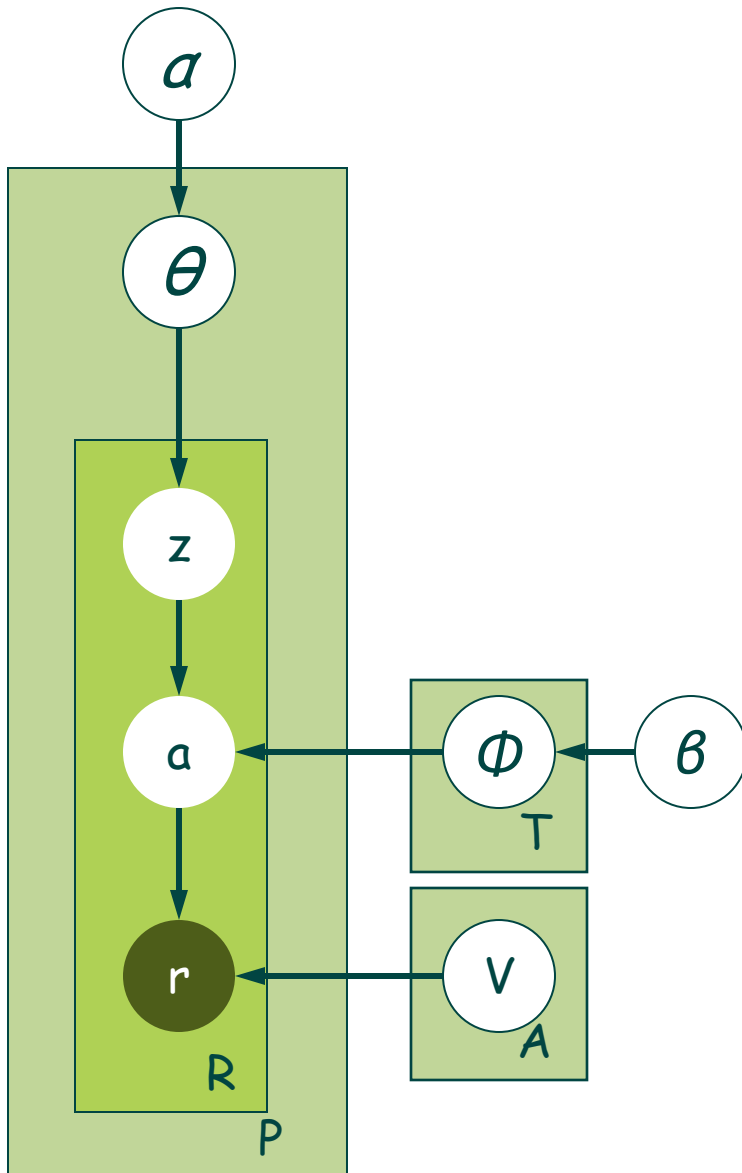


P4: Alfred V. Aho, Stephen C. Johnson,
Jefferey D. Ullman

P5: A. Aho, S. Johnson, J. Ullman

P6: A. Aho, R. Sethi, J. Ullman

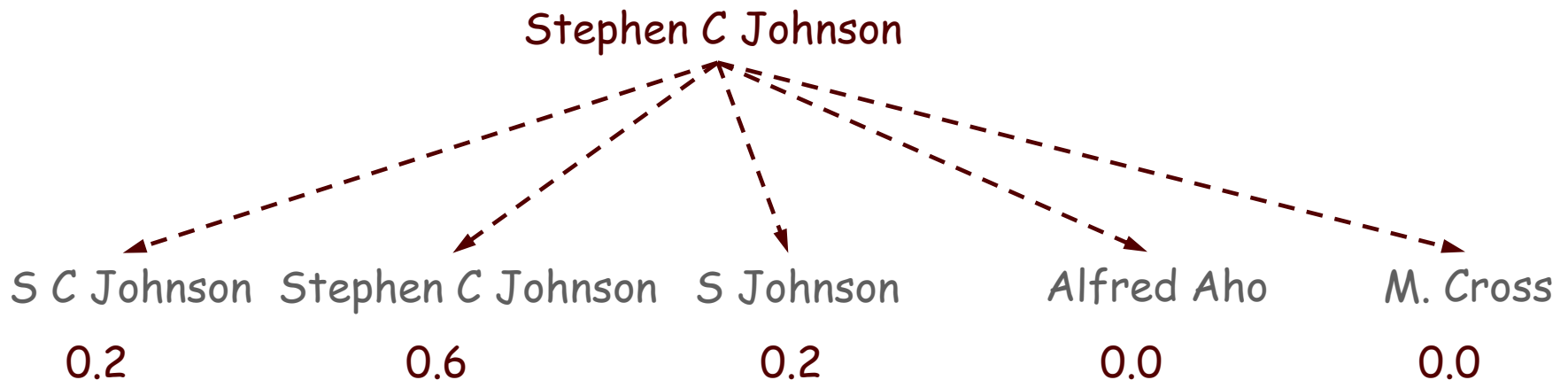
LDA-ER Model



- Entity label a and group label z for each reference r
- θ : 'mixture' of groups for each co-occurrence
- Φ_z : multinomial for choosing entity a for each group z
- V_a : multinomial for choosing reference r from entity a
- Dirichlet priors with α and β

Generating References from Entities

- Entities are not directly observed
 1. Hidden attribute for each entity
 2. Similarity measure for pairs of attributes
- A distribution over attributes for each entity



Approx. Inference Using Gibbs Sampling

- Conditional distribution over labels for each ref.
- Sample next labels from conditional distribution
- Repeat over all references until convergence

$$P(z_i=t|\mathbf{z}_{-i},\mathbf{a},\mathbf{r}) \propto \frac{n_{d_i,t}^{DT} + \alpha/T}{n_{d_i,*}^{DT} + \alpha} \times \frac{n_{a_i,t}^{AT} + \beta/A}{n_{*,t}^{AT} + \beta}$$

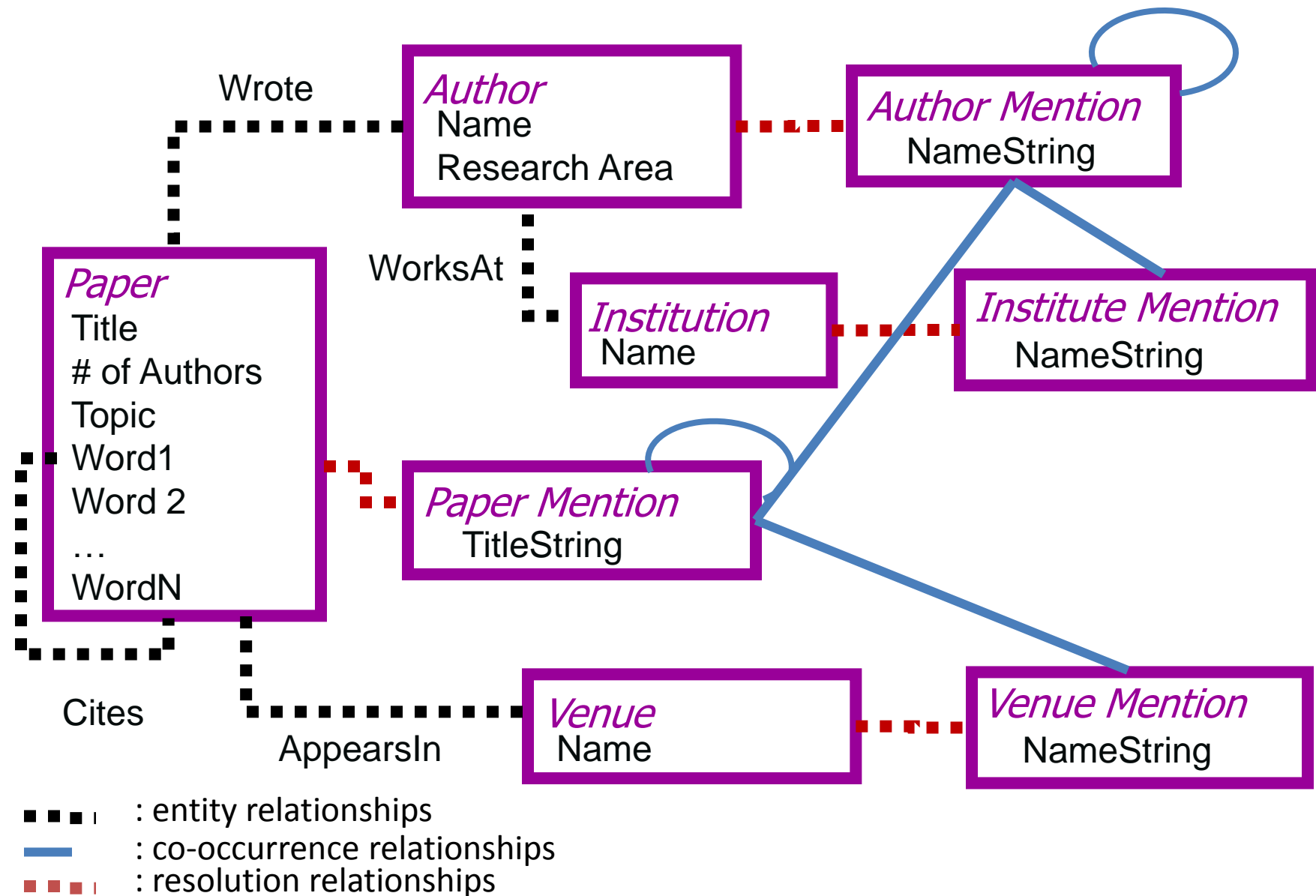
$$P(a_i=a|\mathbf{z},\mathbf{a}_{-i},\mathbf{r}) \propto \frac{n_{a_i,t}^{AT} + \beta/A}{n_{*,t}^{AT} + \beta} \times \text{Sim}(r_i, v_a)$$

- Converges to most likely number of entities

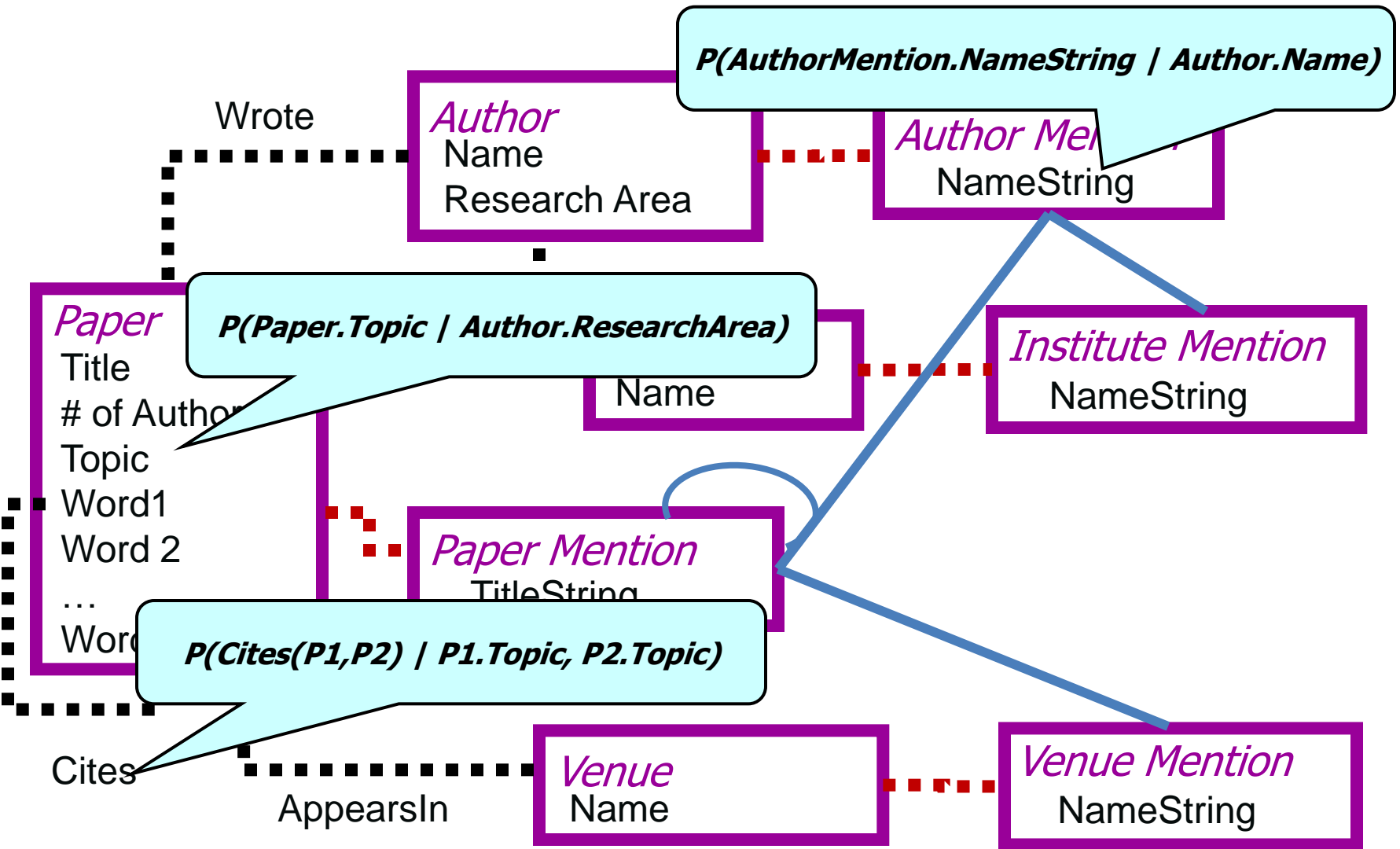
Faster Inference: Split-Merge Sampling

- Naïve strategy reassigns references individually
- Alternative: allow entities to merge or split
- For entity a_i , find conditional probabilities for
 1. Merging with existing entity a_j
 2. Splitting back to last merged entities
 3. Remaining unchanged
- Sample next state for a_i from distribution
- $O(n g + e)$ time per iteration compared to $O(n g + n e)$

Probabilistic Relational Models for ER

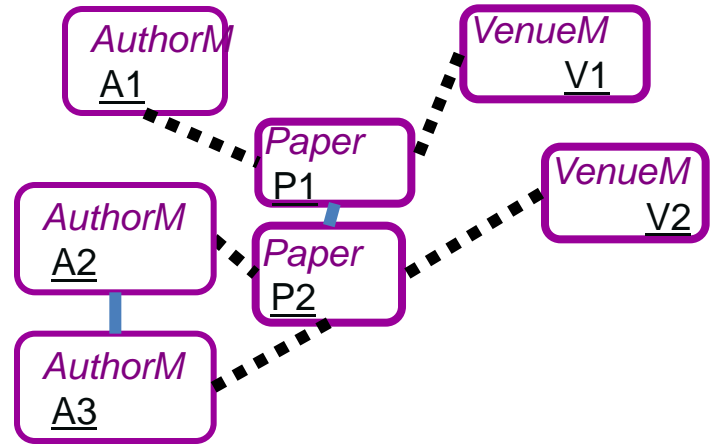
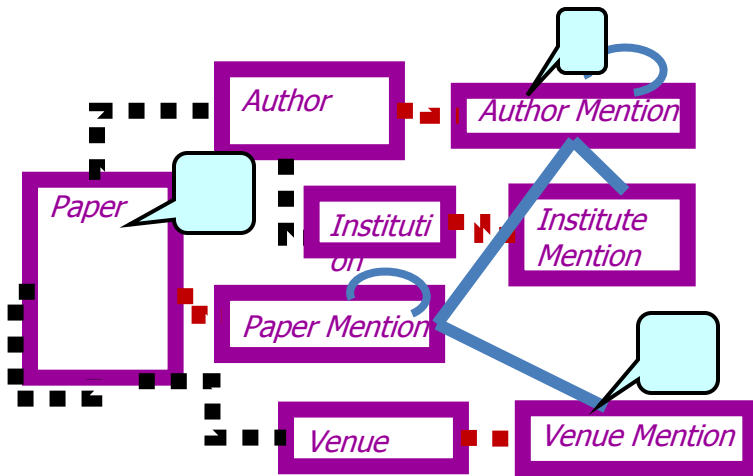


Probabilistic Relational Models



- ■ ■ : entity relationships
- : co-occurrence relationships
- - - : resolution relationships

PRM Semantics



PRM + relational skeleton σ =

probability distribution over completions I :

$$P(I \mid \sigma, \mathcal{S}, \Theta) = \prod_{x \in \sigma} \prod_{x.A} P(x.A \mid \text{parents}_{\mathcal{S}, \sigma}(x.A))$$

↑ ↑
 Objects Attributes

Inference in PRMs for Citation Matching

[Pasula *et al.*, NIPS 2002]

- Parameter estimation
 - Priors for names, titles, citation formats learned offline from labeled data
 - String corruption parameters learned with Monte Carlo EM
- Inference
 - MCMC with cluster recombination proposals
 - Guided by “canopies” of similar citations
 - Accuracy stabilizes after ~20 minutes

Generative Approaches

	Method	Learning/Inference Method	Evaluation
[Li, Morie, & Roth, AAAI 04]	Generative model for mentions in documents	Truncated EM to learn parameters and MAP inference for entities (unsupervised)	F1 on person names, locations and organizations in TREC dataset
Probabilistic Relational Models [Pasula et al., NIPS03]	Probabilistic Relational Models	Parameters learned on separated corpora, inference done using MCMC	% of correctly identified clusters on subsets of CiteSeer data
Latent Dirichlet Allocation [Bhattacharya & Getoor, SDM06]	Latent-Dirichlet Allocation Model	Blocked Gibbs Sampling	Precision/Recall /F1 on CiteSeer and HEP data

PART 4-e

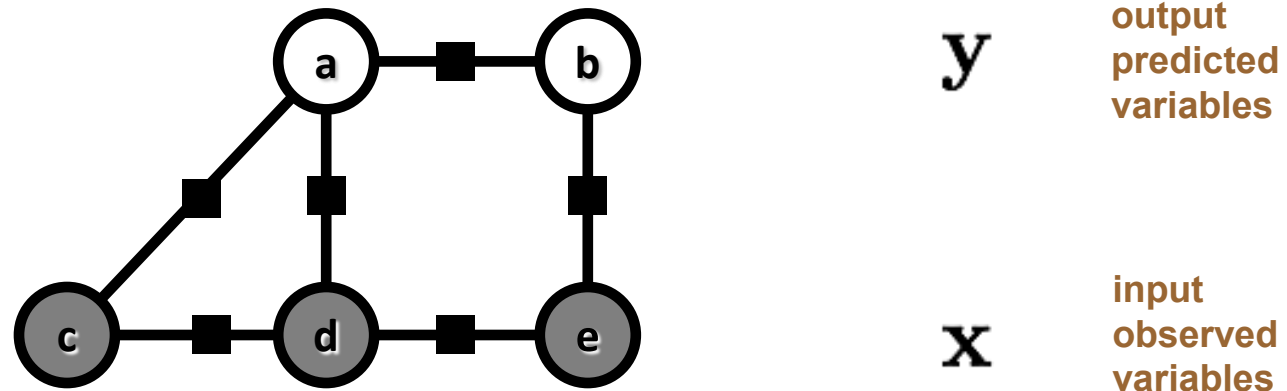
**PROBABILISTIC MODELS:
UNDIRECTED APPROACHES**

Undirected Probabilistic Approaches

- Probabilistic semantics based on Markov Networks
 - Advantage: no acyclicity requirements
- In some cases, syntax based on first-order logic
 - Advantage: declarative
- Examples
 - Conditional Random Fields (CRFs) [McCallum & Wellner, NIPS04]
 - Markov Logic Networks (MLNs) [Singla & Domingos, ICDM06]
 - Probabilistic Similarity Logic [Broecheler & Getoor, UAI10]

Conditional Random Field (CRF)

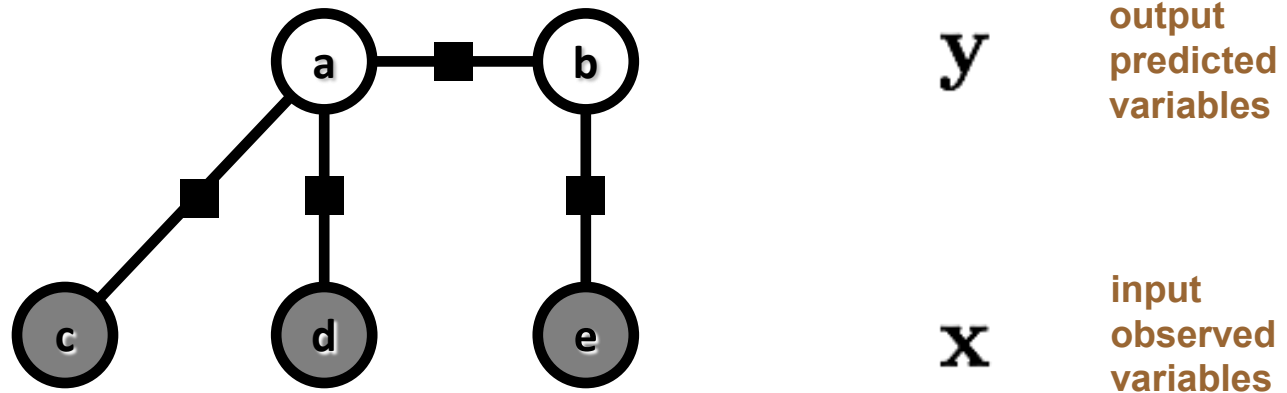
Undirected graphical model, conditioned on some data variables



$$p(\mathbf{y}|\mathbf{x}) = \frac{1}{Z_{\mathbf{x}}} \prod_f \phi(\mathbf{x}_{\in f}, \mathbf{y}_{\in f})$$

Conditional Random Field (CRF)

Undirected graphical model, conditioned on some data variables



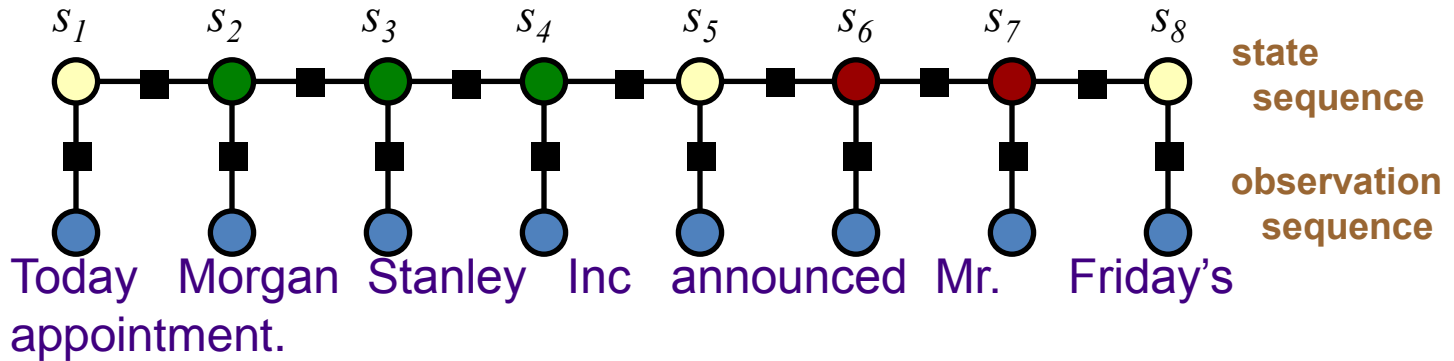
$$p(\mathbf{y}|\mathbf{x}) = \frac{1}{Z_{\mathbf{x}}} \prod_f \phi(\mathbf{x}_{\in f}, \mathbf{y}_{\in f})$$

- + Tremendous freedom to use arbitrary features of input.
- + Predict multiple dependent variables (“structured output”)

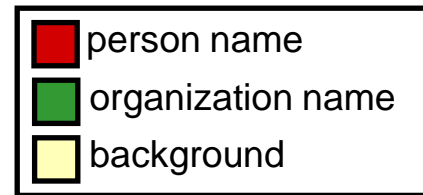
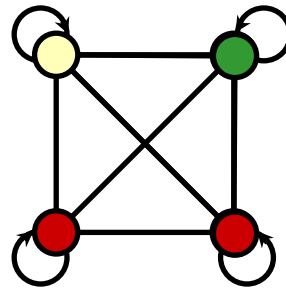
Information Extraction with Linear-chain CRFs

Logistic Regression analogue of a hidden Markov model

Graphical model



Finite state model



CRF for ER

[McCallum & Wellner, NIPS04]

- CRF with random variables for each mention pair
- Factors capture dependence among mentions assigned to the same cluster
- Show that inference in above CRF is equivalent to graph partitioning in graph where nodes are mentions and edges weights are log clique potentials over nodes
- Learn weights from training data; variety of weight learning approaches, here use voted perceptron
- Graph partitioning performed using correlation clustering

Markov Logic

- A logical KB is a set of **hard constraints** on the set of possible worlds
- Make them **soft constraints**; when a world violates a formula, it becomes less probable but not impossible
- Give each formula a **weight**
 - Higher weight \Rightarrow Stronger constraint

$$P(\text{world}) \propto \exp\left(\sum \text{weights of formulas it satisfies}\right)$$

Markov Logic

- A **Markov Logic Network (MLN)** is a set of pairs **(F, w)** where
 - **F** is a formula in first-order logic
 - **w** is a real number

$$P(X) = \frac{1}{Z} \exp \left(\sum_{i \in F} w_i n_i(x) \right)$$

The diagram illustrates the probability formula for a Markov Logic Network. The formula is $P(X) = \frac{1}{Z} \exp \left(\sum_{i \in F} w_i n_i(x) \right)$. Annotations include: a red box around the normalization constant Z with a label 'Normalization Constant' below it; a purple box around the summation index $i \in F$ with a label 'Iterate over all first-order MLN formulas' below it; and a blue box around the term $n_i(x)$ with a label '# true groundings of *ith* clause' above it.

Problem Formulation

- **Given**

- A database of records representing entities in the real world e.g. citations
- A set of fields e.g. author, title, venue
- Each record represented as a set of typed predicates e.g. *HasAuthor(citation,author), HasVenue(citation,venue)*

- **Goal**

- To determine which of the records/fields refer to the same underlying entity

Problem Formulation

- **Given**

- DB of mentions of entities in the real world, e.g. citations
- A set of fields, e.g. author, title, venue
- Each record represented as a set of typed predicates e.g. *HasAuthor(citation,author)*, *HasVenue(citation,venue)*

- Entities in the real world represented by one or more strings appearing in the DB, e.g. "**J. Cox**", "**Cox J.**"
- String constant for each record, e.g. "**C1**", "**C2**"
- **Goal:** for each pair of string constants $\langle x_1, x_2 \rangle$ of the same type, is $x_1 = x_2$?

Handling Equality

- Introduce *Equals(x,y)* or $x = y$
- Introduce the axioms of equality
 - Reflexivity: $x = x$
 - Symmetry: $x = y \Rightarrow y = x$
 - Transitivity: $x = y \wedge y = z \Rightarrow z = x$
 - Predicate Equivalence:

$$x_1 = x_2 \wedge y_1 \wedge y_2 \Rightarrow (R(x_1, y_1) \Leftrightarrow R(x_2, y_2))$$

Handling Equality

- Introduce **reverse predicate equivalence**
- Same relation with the same entity gives evidence about two entities being same

$$R(x_1, y_1) \wedge R(x_2, y_2) \wedge x_1 = x_2 \Rightarrow y_1 = y_2$$

- Not true logically, but gives useful information
- Example

$$\text{HasAuthor}(C1, J. Cox) \wedge \text{HasAuthor}(C2, Cox J.) \wedge (J. Cox = Cox J.) \Rightarrow C1 = C2$$

Model for Entity Resolution

- Model is in the form of an MLN
- Query predicate is *Equality*
- Evidence predicates are relations which hold according to the DB
- Introduce axioms of equality
- First-order rules for field comparison, Fellegi-Sunter model, relational models

Field Comparison

- Each field is a string composed of tokens
- Introduce *HasWord(field, word)*
- Use reverse predicate equivalence

$$\text{HasWord}(f_1, w_1) \wedge \text{HasWord}(f_2, w_2) \wedge w_1 = w_2 \Rightarrow f_1 = f_2$$

- Example

$$\text{HasWord}(J. Cox, Cox) \wedge \text{HasWord}(Cox J., Cox) \wedge (Cox = Cox) \Rightarrow (J. Cox = Cox J.)$$

- Different weight for each word : learnable similarity measure of Bilenko & Mooney [2003]

Two-level Similarity

- Individual words as units: Can't deal with spelling mistakes
- Break each word into ngrams: Introduce *HasNgram(word, ngram)*
- Use reverse predicate equivalence for word comparisons
- Gives a two level similarity measure as proposed by Cohen et al. [2003]

Fellegi-Sunter Model

- Uses Naïve Bayes for match decisions with field comparisons used as predictors
- Simplest Version: Field similarities measured by presence/absence of words in common

$$\text{HasWord}(f_1, w_1) \wedge \text{HasWord}(f_2, w_2) \wedge \text{HasField}(r_1, f_1) \wedge \text{HasField}(r_2, f_2) \wedge w_1 = w_2 \Rightarrow r_1 = r_2$$

- Example

$$\text{HasWord}(J. Cox, Cox) \wedge \text{HasWord}(Cox J., Cox) \wedge \text{HasAuthor}(C1, J. Cox) \wedge \text{HasAuthor}(C2, Cox J.) \wedge (Cox = Cox) \Rightarrow (C1 = C2)$$

Relational Models

- Fellegi-Sunter + transitivity [McCallum & Wellner 2005]

$$(f_1 = f_2) \wedge (f_2 = f_3) \Rightarrow (f_3 = f_1)$$

- Fellegi-Sunter + reverse predicate equivalence for records/fields [Singla & Domingos 2005]

$$\text{HasField}(r_1, f_1) \wedge \text{HasField}(r_2, f_2) \wedge f_1 = f_2 \Rightarrow r_1 = r_2$$

$$\text{HasAuthor}(C1, J. Cox) \wedge \text{HasAuthor}(C2, Cox J.) \wedge (J. Cox = Cox J.) \Rightarrow C1 = C2$$

Relational Models

- Co-authorship relation for entity resolution [Bhattacharya & Getoor, DMKD04]

$$\textit{HasAuthor}(c, a_1) \wedge \textit{HasAuthor}(c, a_2) \Rightarrow \textit{Coauthor}(a_1, a_2)$$

$$\textit{Coauthor}(a_1, a_2) \wedge \textit{Coauthor}(a_3, a_4) \wedge a_1 = a_3 \Rightarrow a_2 = a_4$$

Scalability

- $O(n^2)$ number of match decisions - too big even for small databases
- Use cheap heuristics (e.g. TFIDF based similarity) to identify plausible pairs
- Used the canopy approach [McCallum et al., KDD00]
- Inference/learning over plausible pairs

Probabilistic Soft Logic

- Declarative language for defining **constrained continuous Markov random field** (CCMRF) using first-order logic (FOL)
- Soft logic: truth values in $[0,1]$
- Logical operators relaxed using Lukasiewicz t-norms
- Mechanisms for incorporating similarity functions, and reasoning about sets
- MAP inference is a **convex optimization**
- Efficient sampling method for marginal inference

Predicates and Atoms

- Predicates
 - Describe relations
 - Combined with arguments to make atoms
- Atoms
 - Lifted: contains variables, e.g., Friends(X, Y)
 - Ground: no variables, e.g., AuthorOf(author1, paper1)
- Each ground atom can have a truth value in $[0,1]$
- PSL programs define distributions over the truth values of ground atoms

Weighted Rules

- A PSL program is a set of weighted, logical rules
- For example,

$$\text{authorName}(A1,N1) \wedge \text{authorName}(A2,N2) \wedge \text{similarString}(N1,N2) \\ \Rightarrow \text{sameAuthor}(A1,A2) : 1.0$$

- Variable substitution produces a set of weighted ground rules for a particular data set

Soft Logic Relaxation

- PSL uses the Lukasiewicz t-norm to relax hard logic operators to work on soft truth values

$$a \tilde{\wedge} b = \max\{0, a + b - 1\},$$

$$a \tilde{\vee} b = \min\{a + b, 1\},$$

$$\tilde{\neg} a = 1 - a,$$

- PSL converts rules to logical statements using above operators

$$X \tilde{\Rightarrow} Y \equiv \tilde{\neg} X \tilde{\vee} Y.$$

FOL to CCMRF

- PSL converts a weighted rule into potential functions by penalizing its **distance to satisfaction**, $d(g, x) = (1 - t_g(x))$,
- $t_g(x)$ is the truth value of ground rule g under interpretation x
- The distribution over truth values is

$$\Pr(x) = \frac{1}{Z} \exp \left(\sum_{r \in P} \sum_{g \in G(r)} w_r d(g, x) \right)$$

w_r : weight of rule r

$G(r)$: all groundings of rule r

P : PSL program

PSL Inference

- PSL finds the most likely state by solving

$$\operatorname{argmax}_x P(x) = \operatorname{argmax}_x \sum_{r \in P} \sum_{g \in G(r)} w_r d(g, x)$$

- The t-norms defining $t_g(x)$ form linear constraints on x , making inference a **linear program**
- PSL uses **lazy activation** to ground rules, thus reducing the number of active variables and increasing efficiency
- Other distance metrics (e.g., Euclidean) for distance to satisfaction produce other types of convex objectives (e.g., quadratic programs)

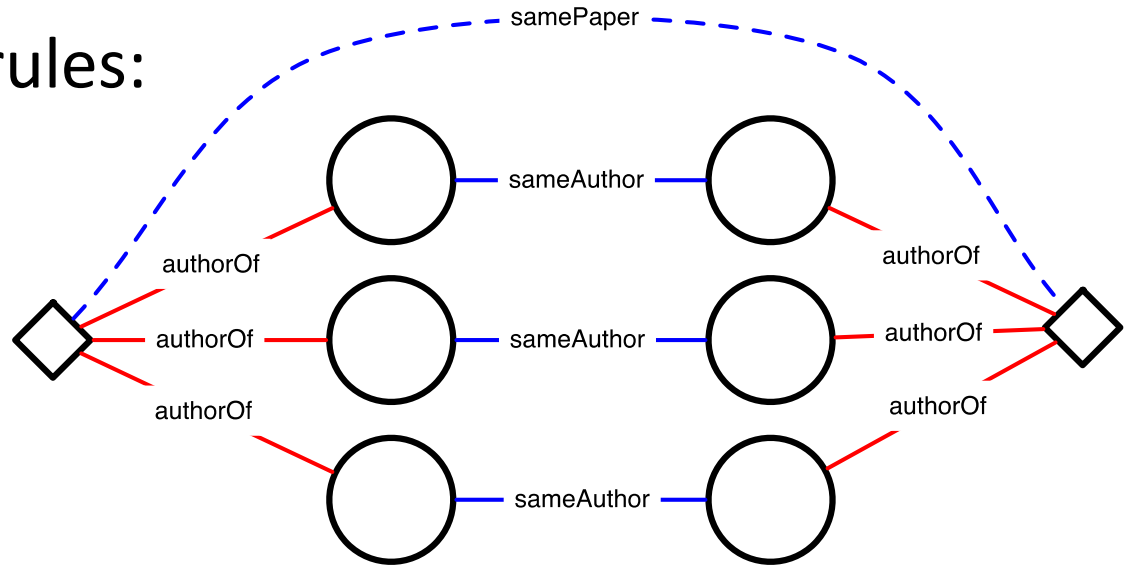
CiteSeer Example

- Citation listings collected from CiteSeer:
 - Pearl J. Probabilistic reasoning in intelligent systems.
Pearl, Judea. Probabilistic Reasoning in Intelligent Systems: Networks of Plausible Inference.
- Duplicate authors and papers
- Base model: Levenstein string similarity
 - $\text{authorName}(A1, N1) \wedge \text{authorName}(A2, N2) \wedge \text{similarString}(N1, N2)$
 $\Rightarrow \text{sameAuthor}(A1, A2)$
 - $\text{paperTitle}(P1, T1) \wedge \text{paperTitle}(P2, T2) \wedge \text{similarString}(T1, T2)$
 $\Rightarrow \text{samePaper}(P1, P2)$
- Only activate rule on pairs with similarity > 0.5

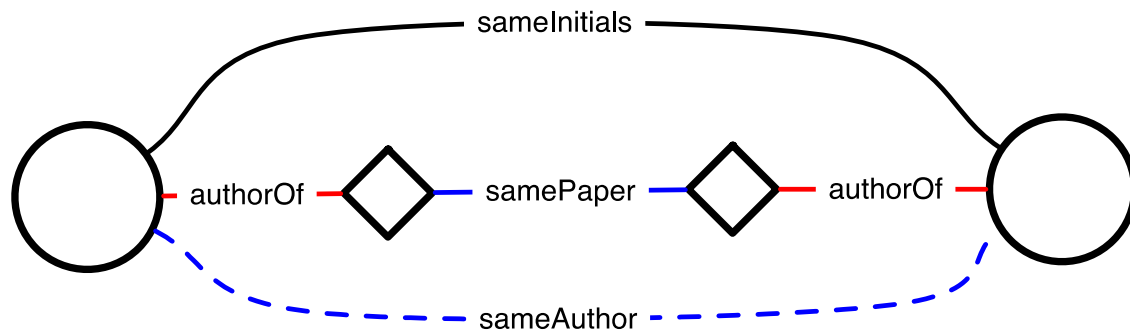
Reasoning about Sets

- Multi-Relational rules:

- $\text{sameAuthorSet}(P1,P2)$
 $\Rightarrow \text{samePaper}(P1,P2)$



- $\text{samePaper}(P1,P2) \wedge \text{authorOf}(A1,P1) \wedge \text{authorOf}(A2,P2) \wedge \text{authorName}(A1,N1) \wedge \text{authorName}(A2,N2) \wedge \text{sameInitials}(N1,N2) \Rightarrow \text{sameAuthor}(A1,A2)$



Undirected Approaches

	Method	Learning/Inference Method	Evaluation
[McCallum & Wellner, NIPS04]	Conditional Random Fields (CRFs) capturing transitivity constraints	Graph partitioning (Boykov et al. 1999), performed via correlation clustering	F1 on DARPA MUC & ACE datasets
[Singla & Domingos, ICDM06]	Markov Logic Networks (MLNs)	Supervised learning and inference using MaxWalkSAT & MCMC	Conditional Log-likelihood and AUC on Cora and BibServ data
[Broecheler & Getoor, UAI10]	Probabilistic Similarity Logic (PSL)	Supervised learning and inference using continuous optimization	Precision/Recall /F1 Ontology Alignment

Summary: Collective Approaches

- Decisions for cluster-membership depends on other clusters
 - Non-probabilistic approaches
 - Similarity propagation approaches
 - Constraint-based approaches
 - Probabilistic Models
 - Generative Models
 - Undirected Models
- Advantages of non-probabilistic approaches is they often scale better than generative probabilistic approaches
- Undirected Models are often easier to specify
- Scaling undirected models active area of research

PART 3

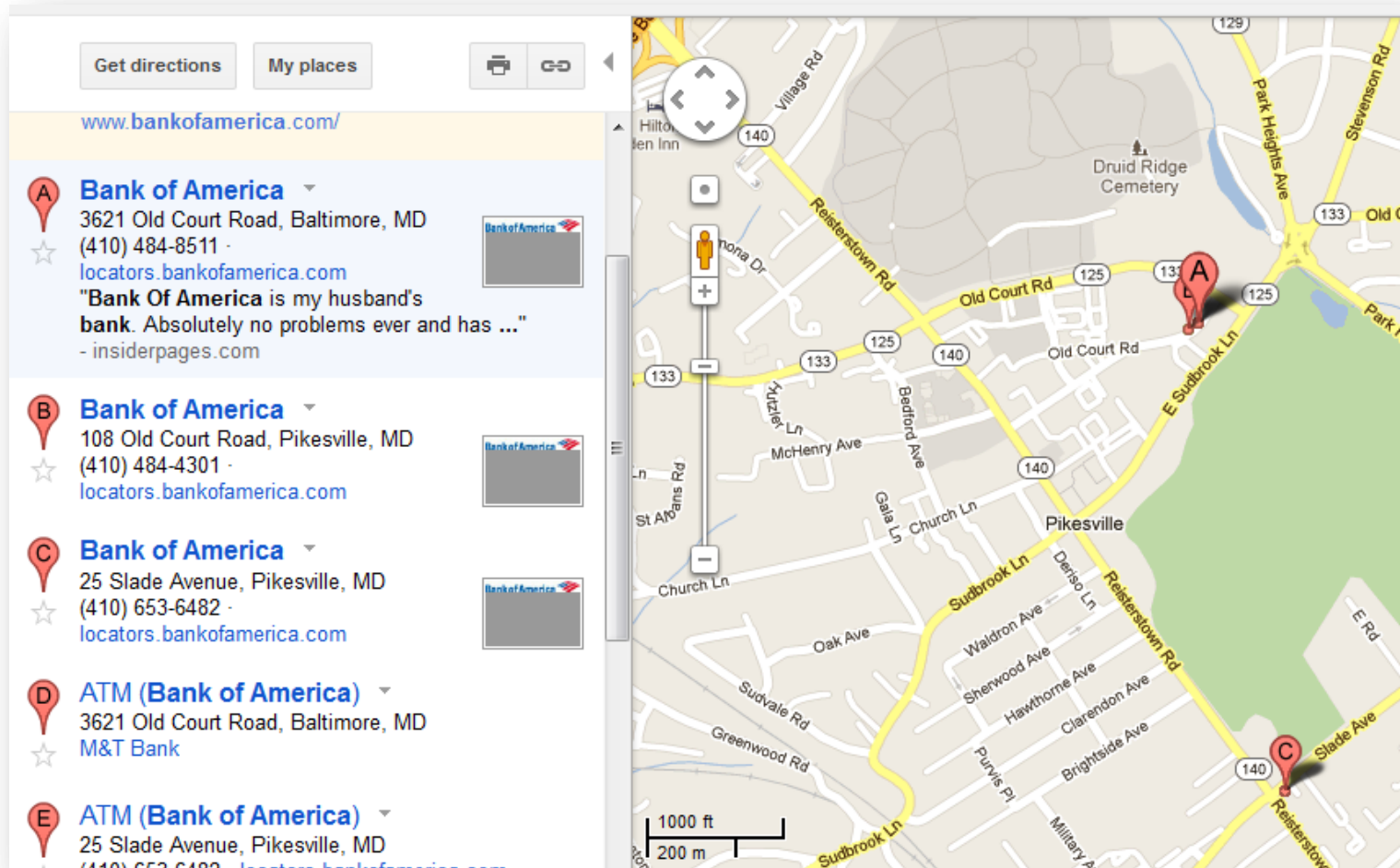
BLOCKING/CANOPY GENERATION

Blocking: Motivation

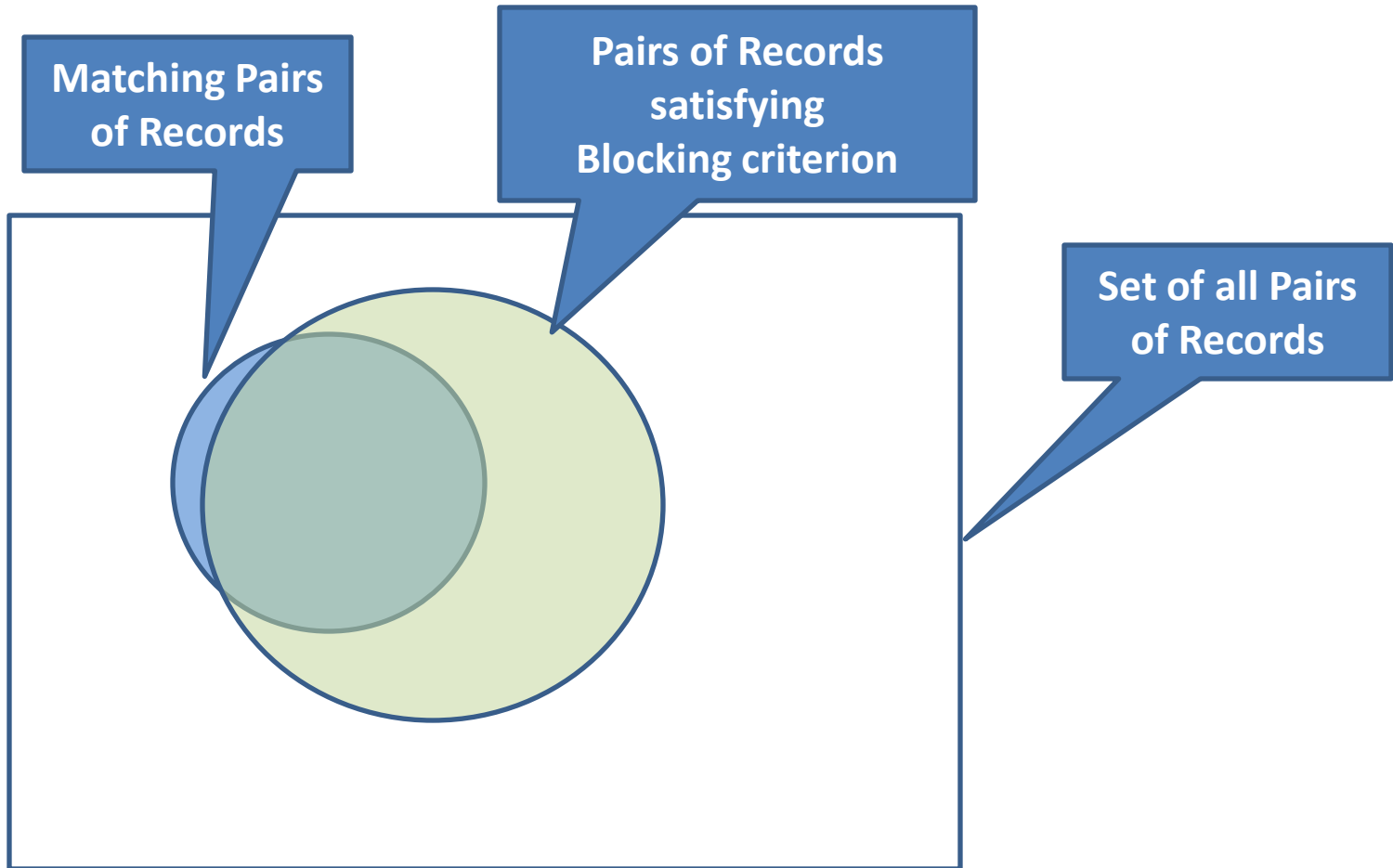
- Naïve pairwise: $|R|^2$ pairwise comparisons
 - 1000 business listings each from 1,000 different cities across the world
 - **1 trillion** comparisons
 - **11.6 days** (if each comparison is $1 \mu\text{s}$)
- Mentions from different cities are unlikely to be matches
 - **Blocking Criterion: City**
 - **10 million** comparisons
 - **10 seconds** (if each comparison is $1 \mu\text{s}$)

Blocking: Motivation

- Mentions from different cities are unlikely to be matches
 - May miss potential matches



Blocking: Motivation



Blocking: Problem Statement

Input: Set of records R

Output: Set of *blocks/canopies*

$$\{C_1, C_2, \dots, C_k\}, \text{ where } \forall_i C_i \subset R \text{ and } \bigcup_i C_i = R$$

Variants:

- *Disjoint Blocking:* Each mention appears in one block.

$$\forall_{i,j} C_i \cap C_j = \emptyset$$

- *Non-disjoint Blocking:* Mentions can appear in more than one block.

Blocking: Problem Statement

$\{C_1, C_2, \dots, C_k\}$, where $\forall_i C_i \subset R$ and $\bigcup_i C_i = R$

Metrics:

- **Efficiency** (or reduction ratio) :
$$\frac{\text{number of pairs compared}}{\text{total number of pairs in } R \times R}$$
$$= \frac{|\{(x, y) \mid \exists i C_i, s. t. x, y \in C_i\}|}{r(r-1)/2}$$
- **Recall*** (or pairs completeness) :
$$\frac{\text{number of true matches compared}}{\text{number of true matches in } R \times R}$$

**Need to know ground truth in order to compute this metric*

Blocking: Problem Statement

Metrics:

- Efficiency (or reduction ratio) : $\frac{\text{number of pairs compared}}{\text{total number of pairs in } R \times R}$
- Recall* (or pairs completeness) : $\frac{\text{number of true matches compared}}{\text{number of true matches in } R \times R}$
- Precision* (or pairs quality) : $\frac{\text{number of true matches compared}}{\text{number of matches compared}}$
- Max Canopy Size: $\max_i |C_i|$

**Need to know ground truth in order to compute this metric*

Blocking Algorithms 1

- Hash based blocking
 - Each block C_i is associated with a hash key h_i .
 - Mention x is hashed to C_i if $hash(x) = h_i$.
 - Within a block, all pairs are compared.
 - Each hash function results in disjoint blocks.
- What *hash* function?
 - Deterministic function of attribute values
 - Boolean Functions over attribute values
[Bilenko et al ICDM'06, Michelson et al AAAI'06, Das Sarma et al CIKM '12]
 - **minHash** (min-wise independent permutations)
[Broder et al STOC'98]

Blocking Algorithms 2

- Pairwise Similarity/Neighborhood based blocking
 - Nearby nodes according to a similarity metric are clustered together
 - Results in non-disjoint canopies.
- Techniques
 - Sorted Neighborhood Approach [Hernandez et al SIGMOD'95]
 - Canopy Clustering [McCallum et al KDD'00]

Simple Blocking: Inverted Index on a Key

Examples of blocking keys:

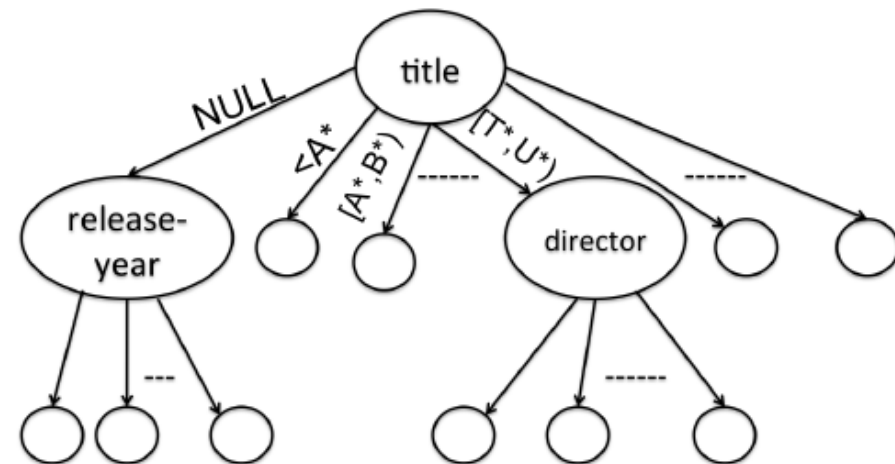
- First three characters of last name
- City + State + Zip
- Character or Token n-grams
- Minimum infrequent n-grams

Learning Optimal Blocking Functions

- Using one or more blocking keys may be insufficient
 - 2,376,206 American's shared the surname Smith in the 2000 US
 - NULL values may create large blocks.
- Solution: Construct blocking functions by combining simple functions

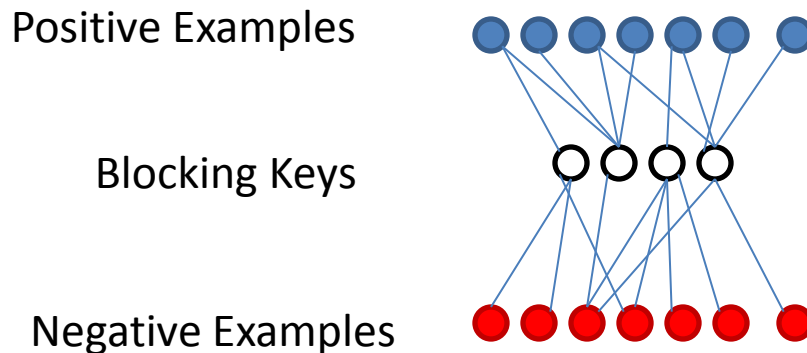
Complex Blocking Functions

- Conjunction of functions [Michelson et al AAAI'06, Bilenko et al ICDM'06]
 - {City} AND {last four digits of phone}
- Chain-trees [Das Sarma et al CIKM'12]
 - **If** ({City} = NULL or LA) **then** {last four digits of phone} AND {area code} **else** {last four digits of phone} AND {City}
- BlkTrees [Das Sarma et al CIKM'12]



Learning an Optimal function [Bilenko et al ICDM '06]

- Find k blocking functions that eliminate the most non-matches, while retaining almost all matches.
 - Need a training set of positive and negative pairs
- Algorithm Idea: Red-Blue Set Cover

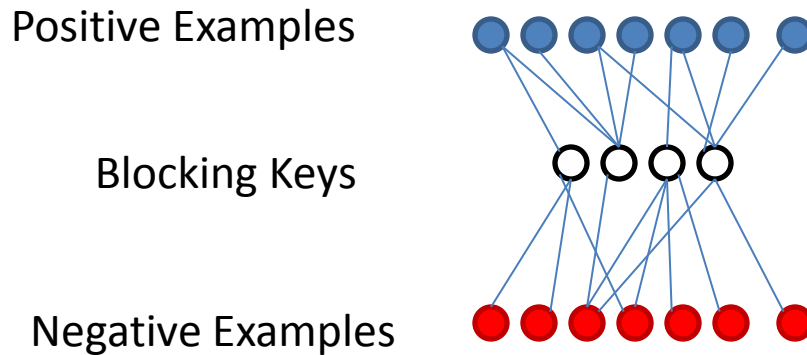


Pick k Blocking keys such that

- At most ϵ blue nodes are not covered
- Number of red nodes covered is minimized

Learning an Optimal function [Bilenko et al ICDM '06]

- Algorithm Idea: Red-Blue Set Cover



Pick k Blocking keys such that

- (a) At most ϵ blue nodes are not covered
- (b) Number of red nodes covered is minimized

- Greedy Algorithm:

- Construct “good” conjunctions of blocking keys $\{p_1, p_2, \dots\}$.
- Pick k conjunctions $\{p_{i_1}, p_{i_2}, \dots, p_{i_k}\}$, such that the following is minimized

$$\frac{\text{number of new blue nodes covered by } p_{i_j}}{\text{number of red nodes covered by } p_{i_j}}$$

minHash (Minwise Independent Permutations)

- Let F_x be a set of features for mention x
 - (functions of) attribute values
 - character ngrams
 - optimal blocking functions ...
- Let π be a random permutation of features in F_x
 - E.g., order imposed by a random hash function
- $\text{minHash}(x) = \text{minimum element in } F_x \text{ according to } \pi$

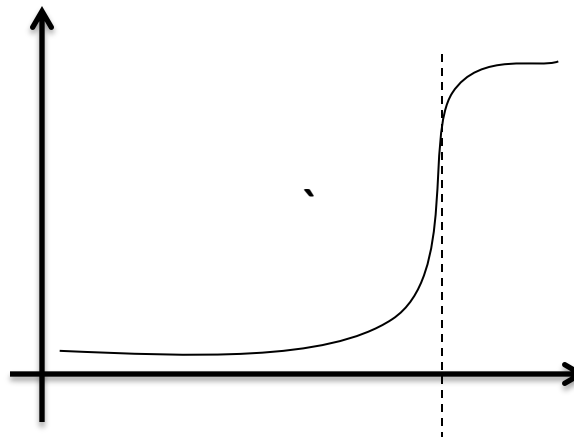
Why minHash works?

Surprising property: For a random permutation π ,

$$P(\text{minHash}(x) = \text{minhash}(y)) = \frac{F_x \cap F_y}{F_x \cup F_y}$$

How to build a blocking scheme such that only pairs with Jacquard similarity $> s$ fall in the same block (with high prob)?

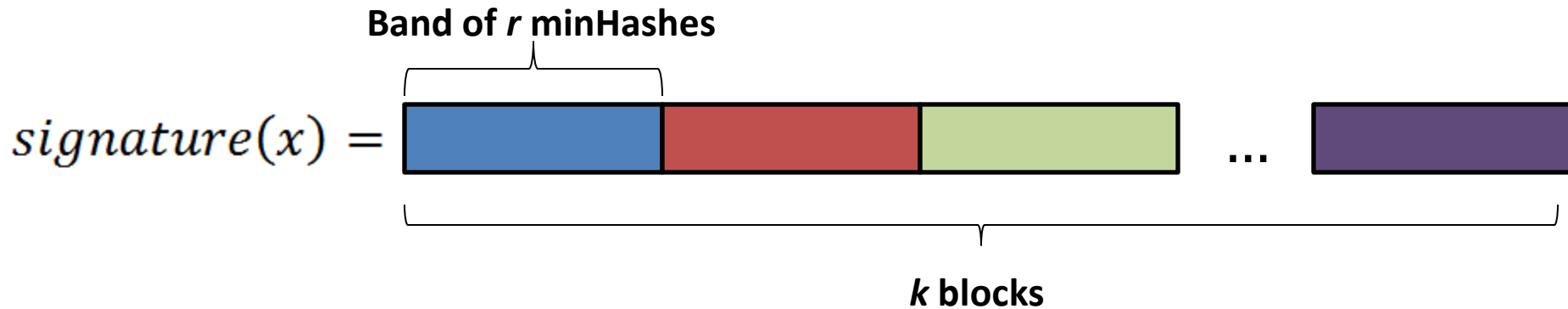
Probability that
(x,y) mentions are
blocked together



Similarity(x,y)

Blocking using minHashes

- Compute minHashes using $r * k$ permutations (hash functions)



- Signature's that match on **1 out of k** bands, go to the same block.

minHash Analysis

False Negatives: (missing matches)

P(pair x,y not in the same block
with Jacquard sim = s) = $(1 - s^r)^k$

should be very low for high similarity pairs

False Positives: (blocking non-matches)

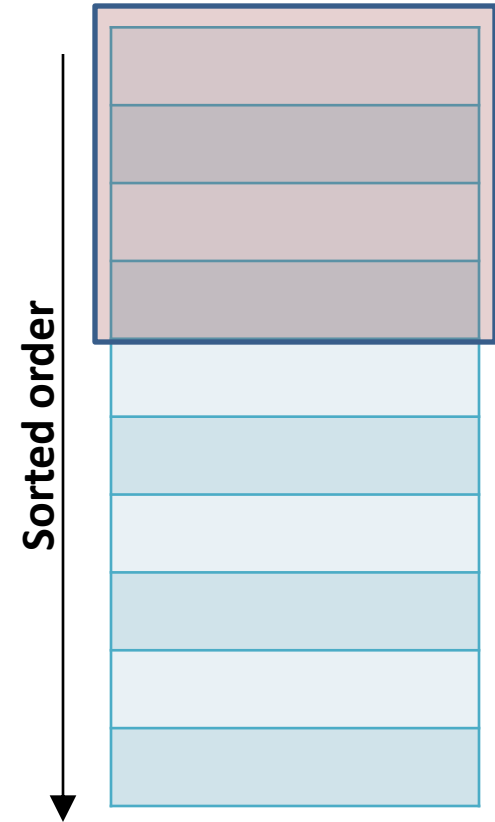
P(pair x,y in the same block
with Jacquard sim = s) = $k \times s^r$

$r = 5, k = 20$

Sim(s)	P(not same block)
0.9	10^{-8}
0.8	0.00035
0.7	0.025
0.6	0.2
0.5	0.52
0.4	0.81
0.3	0.95
0.2	0.994
0.1	0.9998

Sorted Neighborhood [Hernandez et al SIGMOD'95]

- Compute a **Key** for each mention.
- **Sort** the mentions based on the key.
- **Merge**: Check whether a record matches with $(w-1)$ previous records.
 - Efficient implementation using *Sort Merge Band Join* [DeWitt et al VLDB'91]
- Perform multiple passes with different keys

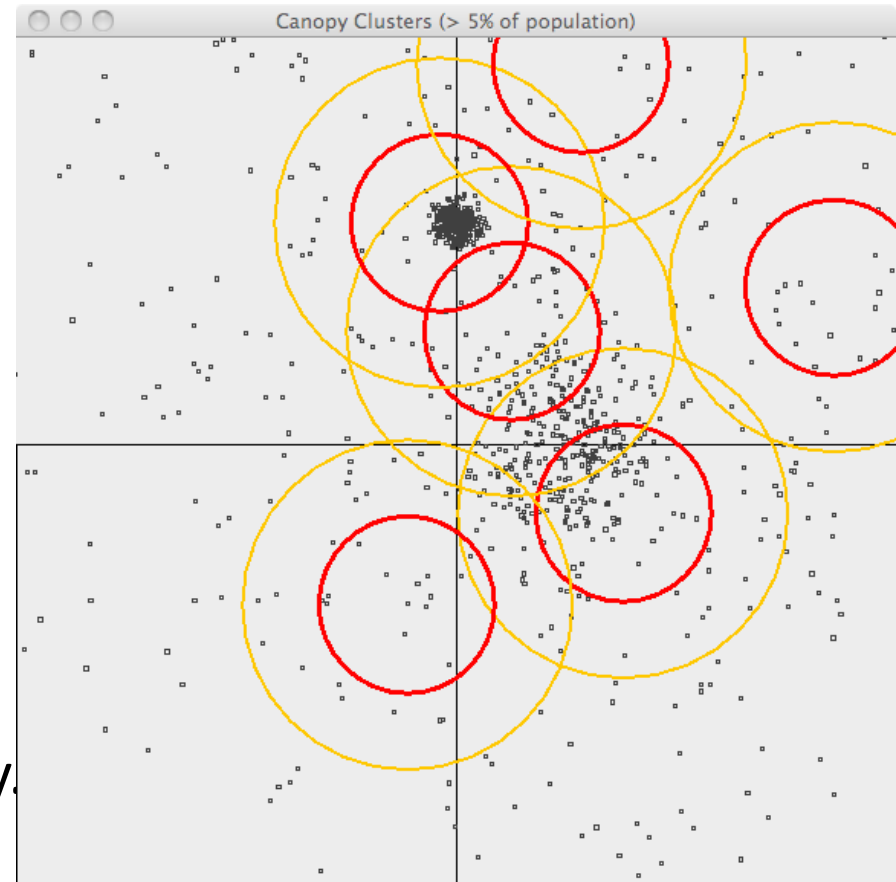


Canopy Clustering [McCallum et al KDD'00]

Input: Mentions M ,
 $d(x,y)$, a distance metric,
thresholds $T_1 > T_2$

Algorithm:

1. Pick a random element x from M
2. Create new canopy C_x using mentions y s.t. $d(x,y) < T_1$
3. Delete all mentions y from M s.t. $d(x,y) < T_2$
4. Return to Step 1 if M is not empty.



SCALING COLLECTIVE ER

Scaling Collective ER [Rastogi et al VLDB11]

Current state-of-the-art: **Collective Entity Matching**

(+) High *accuracy*

(-) Often scale only to a few 1000 entities [SD06],
since runtime is quadratic in the number of pairs.

Example: Dedup papers and authors

Id	Author-1	Author-2	Paper
A ₁	John Smith	Richard Johnson	Indices and Views
A ₂	J Smith	R Johnson	SQL Queries
A ₃	Dr. Smyth	R Johnson	Indices and Views

Slides adapted from [Rastogi et al VLDB11] talk

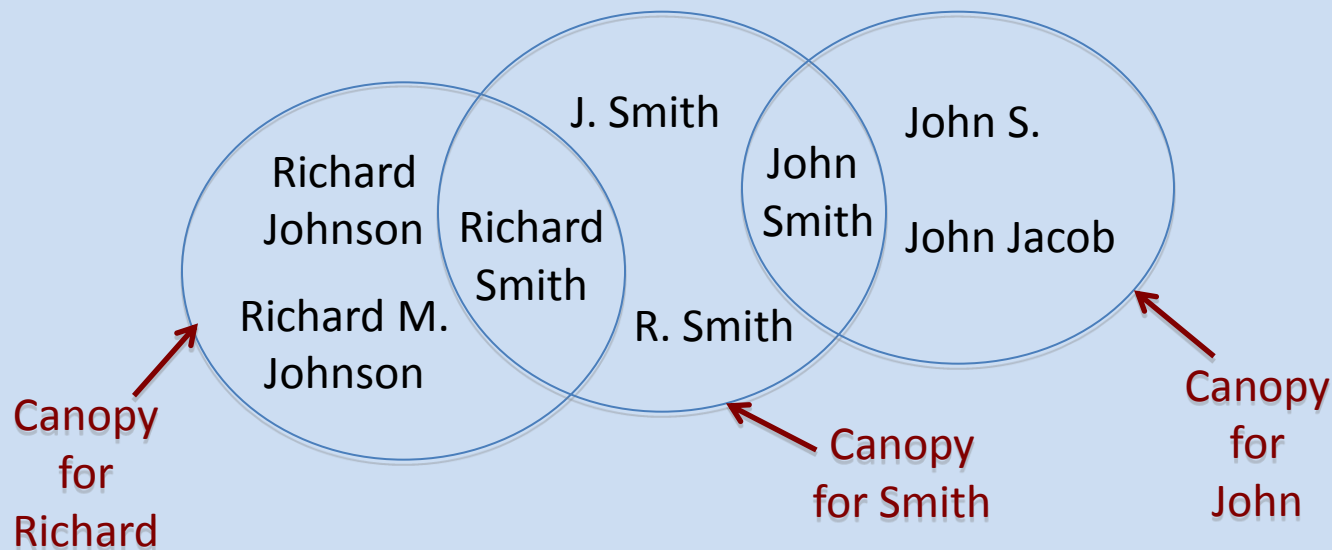
Algorithm

- Generates overlapping canopies (e.g., Canopy clustering)
- Run collective matcher on each canopy

Efficiency: Use Canopies [McCallum et al KDD 00]

Reduces # of candidate pairs from:

$O(|\text{Mentions}|^2)$ to $|\text{Candidate pairs}|$



Pair-wise approach becomes efficient: $O(|\text{Candidate pairs}|)$

Efficiency of Collective approach

Collective methods still not efficient: $\Omega(|\text{Candidate pairs}|^2)$

Example for Collective methods_[SD06]

- $|\text{References}| = 1000$, $|\text{Candidate pairs}| = 15,000$,
 - Time \sim 5 minutes
- $|\text{References}| = 50,000$, $|\text{Candidate pairs}| = 10$ million
 - Time required = 2,500 hours \sim 3 months

Distribute

Run collective entity-matching in each canopy separately

Example for Collective methods_[SD06]

- |References| = 1000, |Candidates| = 15,000,
 - Time = 5 minutes
- One canopy: |References| = 100, |Candidates| ~ 1000,
 - Time ~ 10 Seconds
- |References| = 50,000, # of canopies ~ 13k
 - Time ~ 20 hours << 3 months!

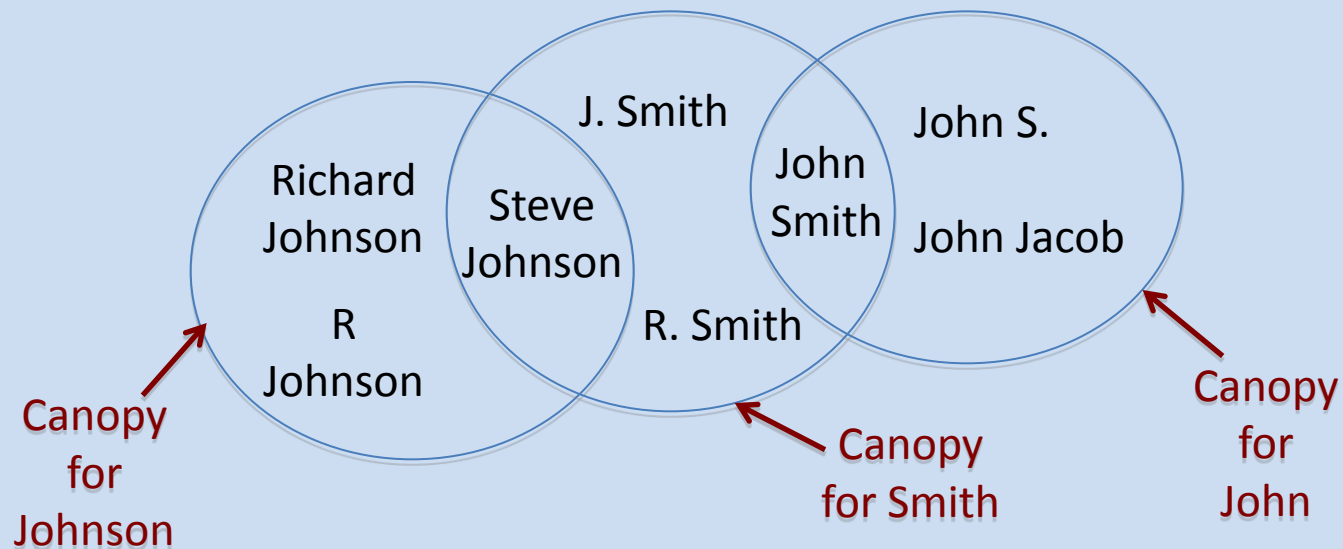
Partitioning into smaller chunks helps!

Problem: Correlations across canopies will be lost

$\text{CoAuthor}(A_1, B_1) \wedge \text{CoAuthor}(A_2, B_2) \wedge \text{match}(B_1, B_2) \rightarrow \text{match}(A_1, A_2)$

Example: CoAuthor rule grounds to the correlation

$\text{match}(\text{Richard Johnson}, \text{R Johnson}) \Rightarrow \text{match}(\text{J. Smith}, \text{John Smith})$



Message Passing

Simple Message Passing (SMP)

1. Run entity matcher **M** locally in each canopy
2. If **M** finds a **match**(r_1, r_2) in some canopy, **pass** it as evidence to all canopies
3. Rerun **M** within each canopy using **new evidence**
4. Repeat until no new matches found in each canopy

Runtime: $O(k^2 f(k) c)$

- k : maximum size of a canopy
- $f(k)$: Time taken by ER on canopy of size k
- c : number of canopies

Formal Properties

for a well behaved ER method ...

Convergence: No. of steps \leq no. of matches

Consistency: Output independent of the canopy order

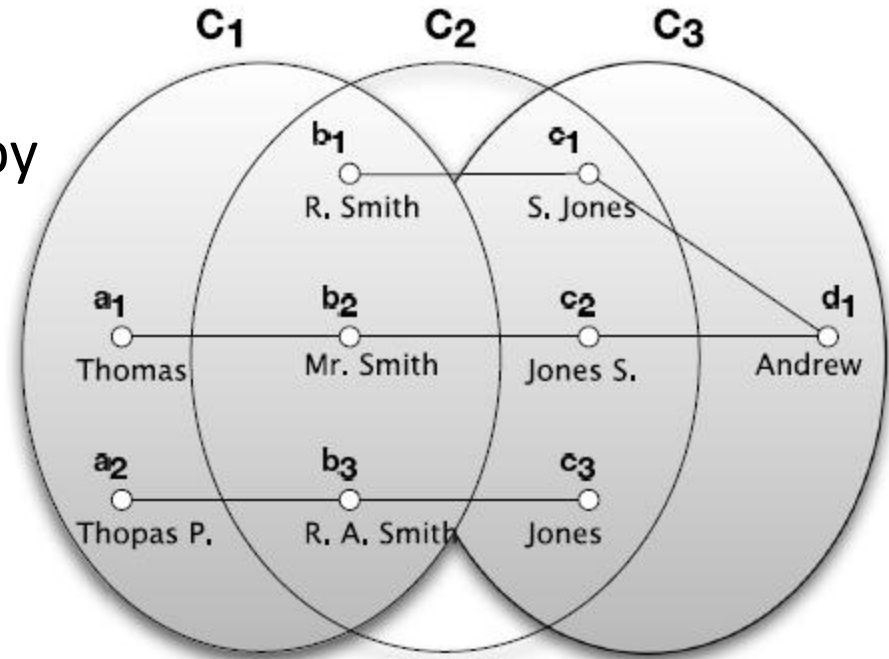
Soundness: Each output match is actually a true match

~~**Completeness:** Each true match is also a output match~~

Completeness

Papers 2 and 3 match only if a canopy knows that

- match(a1,a2)
- match(b2,b3)
- match(c2,c3)



Simple message passing will not find any matches

- thus, no messages are passed, no progress

Solution: Maximal message passing

- **Send a message if there is a potential for match**

Summary of Blocking

- $O(|R|^2)$ pairwise computations can be prohibitive.
 - Blocking eliminates comparisons on a large fraction of non-matches.
- Equality-based Blocking:
 - Construct (one or more) blocking keys from features
 - Records not matching on any key are not compared.
- Similarity based Blocking:
 - Form overlapping canopies of records based on similarity.
 - Only compare records within a cluster.
- Message Passing + blocking can help scale collective ER.

Part 4

**CHALLENGES AND FUTURE
DIRECTIONS**

Challenges

- So far, we have viewed ER as a one-time process applied to entire database; none of these hold in real world.
- Temporal ER
 - ER algorithms need to account for change in real world
 - Reasoning about multiple sources [Pal & M et al. WWW 12]
 - Model transitions [Li et al VLDB11]
- Reasoning about source quality
 - Sources are not independent
 - Copying Problem [Dong et al VLDB09]
- Query Time ER
 - How do we selectively determine the smallest number of records to resolve, so we get accurate results for a particular query?
 - Collective resolution for queries [Bhattacharya & Getoor JAIR07]

Open Issues

- ER & User-generated data
 - Deduplicated entities interact with users in the real world
 - Users tag/associate photos/reviews with businesses on Google / Yahoo
 - What should be done to support interactions?
- ER is often part of bigger inference problem
 - Pipelined approaches and joint approaches to information extraction and graph identification
 - How can we characterize how ER errors affect overall quality of results?
- ER Theory
 - Need better support for theory which can give relational learning bounds
- ER & Privacy
 - ER enables record re-identification
 - How do we develop a theory of privacy-preserving ER?

Summary

- Growing omnipresence of massive linked data, and the need for creating knowledge bases from text and unstructured data motivate a number of challenges in ER
- Especially interesting challenges and opportunities for ER and social media data
- As data, noise, and knowledge grows, greater needs & opportunities for intelligent reasoning about entity resolution
- Many other challenges
 - Large scale identity management
 - Understanding theoretical potentials & limits of ER

THANK YOU!

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