



Data Profiling

An ICDE 2016 Tutorial

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If we just have a bunch of data sets in a repository, it is unlikely anyone will ever be able to find, let alone reuse, any of this data. With adequate metadata, there is some hope, but even so, challenges will remain.. ,



[D. Agrawal, P. Bernstein, E. Bertino, S. Davidson, U. Dayal, M. Franklin, J. Gehrke, L. Haas, A. Halevy, J. Han, H. V. Jagadish, A. Labrinidis, S. Madden, Y. Papakonstantinou, J. M. Patel, R. Ramakrishnan, K. Ross, C. Shahabi, D. Suciu, S. Vaithyanathan, and J. Widom. Challenges and opportunities with Big Data. Technical report, Computing Community Consortium, <http://cra.org/ccc/docs/init/bigdatawhitepaper.pdf>, 2012.]

Profiling relational data: a survey

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Abstract Profiling data to determine metadata about a given dataset is an important and frequent activity of any IT professional and researcher and is necessary for various use-cases. It encompasses a vast array of methods to examine datasets and produce metadata. Among the simpler results are statistics, such as the number of null values and distinct values in a column, its data type, or the most frequent patterns of its data values. Metadata that are more difficult to compute involve multiple columns, namely correlations, unique column combinations, functional dependencies, and inclusion dependencies. Further techniques detect condi-

1 Data profiling: finding metadata

Data profiling is the set of activities and processes to determine the metadata about a given dataset. Profiling data is an important and frequent activity of any IT professional and researcher. We can safely assume that any reader of this article has engaged in the activity of data profiling, at least by eye-balling spreadsheets, database tables, XML files, etc. Possibly, more advanced techniques were used, such as keyword searching in datasets, writing structured queries, or even using dedicated data profiling tools.

Tutorial Overview

- Motivation
 - Task classification
 - Use cases
- Tools
 - Research and industry
 - Shortcomings
- Single and Multiple Column Analysis
 - Cardinalities and datatypes
 - Co-occurrences and summaries
- Dependencies
 - UCCs, INDs, FDs
 - and their discover algorithms
- Outlook
 - Functionality
 - Semantics



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12		1		ALAMANCE		9121656 A		ACTIVE		AV		VERIFIED		ABADIE		JACK		EDWARD JR		612 SIDEVIEW ST		GRAHAM		NC		27253 612 SIDEVIEW ST		GRAHAM NC		27253 336 212 8140		W		NL		UNA	
13		1		ALAMANCE		9118154 I		INACTIVE		IU		CONFIRMATIABADIE		COLLEEN		MIASHEL		1097 IVEY RD #C		GRAHAM		NC		27253 1097 IVEY RD #C		GRAHAM NC		27217 336 212 8140		W		NL		UNA			
14		1		ALAMANCE		9131788 A		ACTIVE		AV		VERIFIED		ABBAS		FALISA		507 SUMMIT RIDGE MEbane		NC		27302 707 SUMMIT RIDGE RD #		MEbane		NC		27302 919 568 9001		B		UN		DEM			
15		1		ALAMANCE		9068460 A		ACTIVE		AV		VERIFIED		ABBAS		RAFAT		514 WESTRIDGE D BURLINGTON		NC		27215 514 WESTRIDGE DR		BURLINGTON NC		27215 A		W		UN		DEM					
16		1		ALAMANCE		9049573 A		ACTIVE		AV		VERIFIED		ABBATECOL/RONALD		JOSEPH JR		504 BROOKFIELD E GIBSONVILLE		NC		27249 504 BROOKFIELD DR		GIBSONVILLE NC		27249 W		NL		UNA		DEM					
17		1		ALAMANCE		9033877 A		ACTIVE		AV		VERIFIED		ABBOTT		BOONE		504 BROOKFIELD E GIBSONVILLE		NC		27249 504 BROOKFIELD DR		GIBSONVILLE NC		27249 W		NL		DEM							
18		1		ALAMANCE		9083557 I		INACTIVE		IU		CONFIRMATIABBET		DAWN		LEANN		3900 JOHNS CREEK DR		NC		27249 3900 JOHNS CREEK DR		GIBSONVILLE NC		27249 336 584 3319		W		NL		DEM					
19		1		ALAMANCE		9027554 A		ACTIVE		AV		VERIFIED		ABBEY		BRENT		3304 GOLDEN OAK GRAHAM		NC		27253 3304 GOLDEN OAKS DR		GRAHAM		NC		27253 919 682 6873		W		NL		REP			
20		1		ALAMANCE		9029477 A		ACTIVE		AV		VERIFIED		ABBEY		DEMETRA		AINSWORTH		3304 GOLDEN OAK GRAHAM		NC		27253 3													

904055	A	ACTIVE	AV	VERIFIED	HAWKINS	DEBORAH	A	307 N SEVENTH ST	MEBANE	NC	27302	307 N SEVENTH ST	MEBANE
9115545	A	ACTIVE	AV	VERIFIED	HAWKINS	DERRICK	JEROME	106 TADWORTH CT	MEBANE	NC	27302	106 TADWORTH CT	MEBANE
9060012	A	ACTIVE	AV	VERIFIED	HAWKINS	DIANA	LEE	424 MEADOWOOD	BURLINGTON	NC	27215	424 MEADOWOOD DR	BURLINGTON
9118697	A	ACTIVE	AV	VERIFIED	HAWKINS	DOMINIQUE	DEVON	3 SHERRY DR	BURLINGTON	NC	27215	3 SHERRY DR	BURLINGTON
2848800	R	REMOVED	RD	DECEASED	HAWKINS	DONALD	LEE	2847 SNUG HARBOR	BURLINGTON	NC	27217	2847 SNUG HARBOR RD	BURLINGTON
9025486	I	INACTIVE	IN	CONFIRMATI	HAWKINS	DONNA	KAYE	859 ROSS ST	BURLINGTON	NC	27217	859 ROSS ST	BURLINGTON
9134349	A	ACTIVE	AV	VERIFIED	HAWKINS	ELAINE	TERESA	779 WOODY DR	GRAHAM	NC	27253	779 WOODY DR	GRAHAM
9081107	A	ACTIVE	AV	VERIFIED	HAWKINS	ERIC	THOMAS	1720 OLD ST MARK'S CHURCH	BURLINGTON	NC	27215	1720 OLD ST MARK'S CHURCH	BURLINGTON
9110146	A	ACTIVE	AV	VERIFIED	HAWKINS	ERIC	MICHAEL	5828 ANDOVER DR	GRAHAM	NC	27253	5828 ANDOVER DR	GRAHAM
9018277	A	ACTIVE	AV	VERIFIED	HAWKINS	ERIK	CHRISTIAN	2420 US HWY 70	MEBANE	NC	27302	2428 US HWY 70	MEBANE
9010269	A	ACTIVE	AV	VERIFIED	HAWKINS	ERIK	CHRISTIAN	307 N EVENTH ST	MEBANE	NC	27302	307 N EVENTH ST	MEBANE
9072769	A	ACTIVE	AV	VERIFIED	HAWKINS	HEATHER	ANN	7439 COBLE MILL FS	SNOW CAMP	NC	27349	7439 COBLE MILL RD	SNOW CAMP
2850000	A	ACTIVE	AV	VERIFIED	HAWKINS	IRIS	WATKINS	2912 MARLBOROUGH	BURLINGTON	NC	27215	2912 MARLBOROUGH RD	BURLINGTON
9139873	A	ACTIVE	AV	VERIFIED	HAWKINS	ISAIAH	FORRIESHE	726 DAILEY ST	BURLINGTON	NC	27217	726 DAILEY ST	BURLINGTON
9102693	A	ACTIVE	AV	VERIFIED	HAWKINS	JACQUELINE	ISLEY	2111 FAIRWIND DR	GRAHAM	NC	27253	2111 FAIRWIND DR	GRAHAM
2850100	A	ACTIVE	AV	VERIFIED	HAWKINS	JACQUELINE	ISLEY	859 ROSS ST	BURLINGTON	NC	27217	859 ROSS ST	BURLINGTON
9131359	A	ACTIVE	AV	VERIFIED	HAWKINS	JAIJUAN	DEBRADSFER	203 EDWARD CT	MEBANE	NC	27302	203 EDWARD CT	MEBANE
2850401	A	ACTIVE	AV	VERIFIED	HAWKINS	JAMES	D	30 GRANITE CT	GIBSONVILLE	NC	27249	30 GRANITE CT	GIBSONVILLE
9034990	A	ACTIVE	AV	VERIFIED	HAWKINS	JAMES	EDWARD	1107 SOUTHERN H	BURLINGTON	NC	27215	1107 SOUTHERN HIGH SCH	BURLINGTON
9102435	A	ACTIVE	AV	VERIFIED	HAWKINS	JAMES	EDWARD	1107 SOUTHERN H	BURLINGTON	NC	27215	1107 SOUTHERN HIGH SCH	BURLINGTON
9083219	A	ACTIVE	AV	VERIFIED	HAWKINS	JERMANE	KENDRICK	109 SLADE ST	ELON	NC	27244	109 SLADE ST	ELON
9013096	A	ACTIVE	AV	VERIFIED	HAWKINS	JERRY	MICHAEL	2730 BELLEMONT-B	BURLINGTON	NC	27215	2730 BELLEMONT-ALAMA	BURLINGTON
9110147	A	ACTIVE	AV	VERIFIED	HAWKINS	KOELLE	ROELLE	5828 ANDOVER DR	GRAHAM	NC	27253	5828 ANDOVER DR	GRAHAM
9119019	A	ACTIVE	AV	VERIFIED	HAWKINS	JOHN	MATSON	3314 N NC HWY 62	BURLINGTON	NC	27217	3314 N NC HWY 62	BURLINGTON
2851100	A	ACTIVE	AV	VERIFIED	HAWKINS	JOHN	THOMAS	613 N FOURTH ST	MEBANE	NC	27302	613 N FOURTH ST	MEBANE
9029983	A	ACTIVE	AV	VERIFIED	HAWKINS	KIRK	THOMAS	232 MONROE LN	ELON	NC	27244	232 MONROE LN	ELON
9001801	R	REMOVED	RL	MOVED-OUT	HAWKINS	JOHN	DANIEL	862 ROSS ST	BURLINGTON	NC	27217	862 ROSS ST	BURLINGTON
9008655	R	REMOVED	RL	MOVED-OUT	HAWKINS	JOHN	DANIEL	862 ROSS ST	BURLINGTON	NC	27217	862 ROSS ST	BURLINGTON
9109154	I	INACTIVE	IN	CONFIRMATI	HAWKINS	JUSTIN	ANDREW	2111 FAIRWIND DR	GRAHAM	NC	27253	2111 FAIRWIND DR	GRAHAM
9063027	A	ACTIVE	AV	VERIFIED	HAWKINS	KAREN	COOK	1717 DURHAM ST	BURLINGTON	NC	27217	1717 DURHAM ST #61	BURLINGTON
9014773	A	ACTIVE	AV	VERIFIED	HAWKINS	KATHY	ROGERS	716 S WILLIAMSON	ELON	NC	27244	716 S WILLIAMSON AVE	ELON
2851300	A	ACTIVE	AV	VERIFIED	HAWKINS	KENNETH	WESLEY	485 PARKVIEW DR	BURLINGTON	NC	27215	485 PARKVIEW DR	BURLINGTON
9115548	A	ACTIVE	AV	VERIFIED	HAWKINS	KIAIR	JESSICA-SHA	114 W SEBASTIAN	MEBANE	NC	27215	1107 SOUTHERN HIGH SCH	BURLINGTON
9059505	D	DENIED	DI	UNAVAILABLE	HAWKINS	KATRINA	NICOLE	2430 MARION CT	BURLINGTON	NC	27215	2430 MARION CT	BURLINGTON
9135064	A	ACTIVE	AV	VERIFIED	HAWKINS	KENNETH	WESLEY	3165 WILLIAMS LN	GRAHAM	NC	27302	114 W SEBASTIAN CT	MEBANE
9133012	A	ACTIVE	AV	VERIFIED	HAWKINS	KIAR	JESSICA-SHA	618 CENTER AVE	BURLINGTON	NC	27253	3165 WILLIAMS LN	GRAHAM
9124536	I	INACTIVE	IN	CONFIRMATI	HAWKINS	LADARIS	CHONDELLE	801 TROLLINGWOOD	MEBANE	NC	27215	618 CENTER AVE #C	BURLINGTON
9109155	A	ACTIVE	AV	VERIFIED	HAWKINS	LADONNA	EDWINA	114 W SEBASTIAN	MEBANE	NC	27302	801 TROLLINGWOOD-HA	MEBANE
9135065	A	ACTIVE	AV	VERIFIED	HAWKINS	LIZA	LYNN	114 W SEBASTIAN	MEBANE	NC	27302	114 W SEBASTIAN CT	MEBANE
9079866	A	ACTIVE	AV	VERIFIED	HAWKINS	LORA	ROBERT	406-410 1ST CT	BURLINGTON	NC	27217	1288 ELWOOD CT	BURLINGTON
9120114	D	DENIED	DU	VERIFICATIO	HAWKINS	LORETTA	ANNE	408 HOOD ST	BURLINGTON	NC	27217	408 HOOD ST	BURLINGTON
2851600	R	REMOVED	RD	DECEASED	HAWKINS	MAE	PITTMAN	2730 BELLEMONT-B	BURLINGTON	NC	27215	2730 BELLEMONT-ALAMA	BURLINGTON

Many interesting questions remain

- What are possible keys and foreign keys?
 - Phone
 - firstname, lastname, street
- Are there any functional dependencies?
 - zip -> city
 - race -> voting behavior
- Which columns correlate?
 - Date-of-Birth and first name
 - State and last name
- What are frequent patterns in a column?
 - dddd
 - dd aaaa St

Definition Data Profiling

- Data profiling is the process of examining the data available in an existing data source [...] and collecting statistics and information about that data.

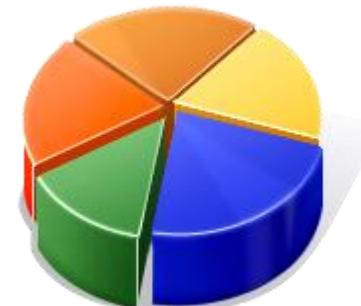
[Wikipedia 04/2016]

- Data profiling refers to the activity of creating small but informative summaries of a database.

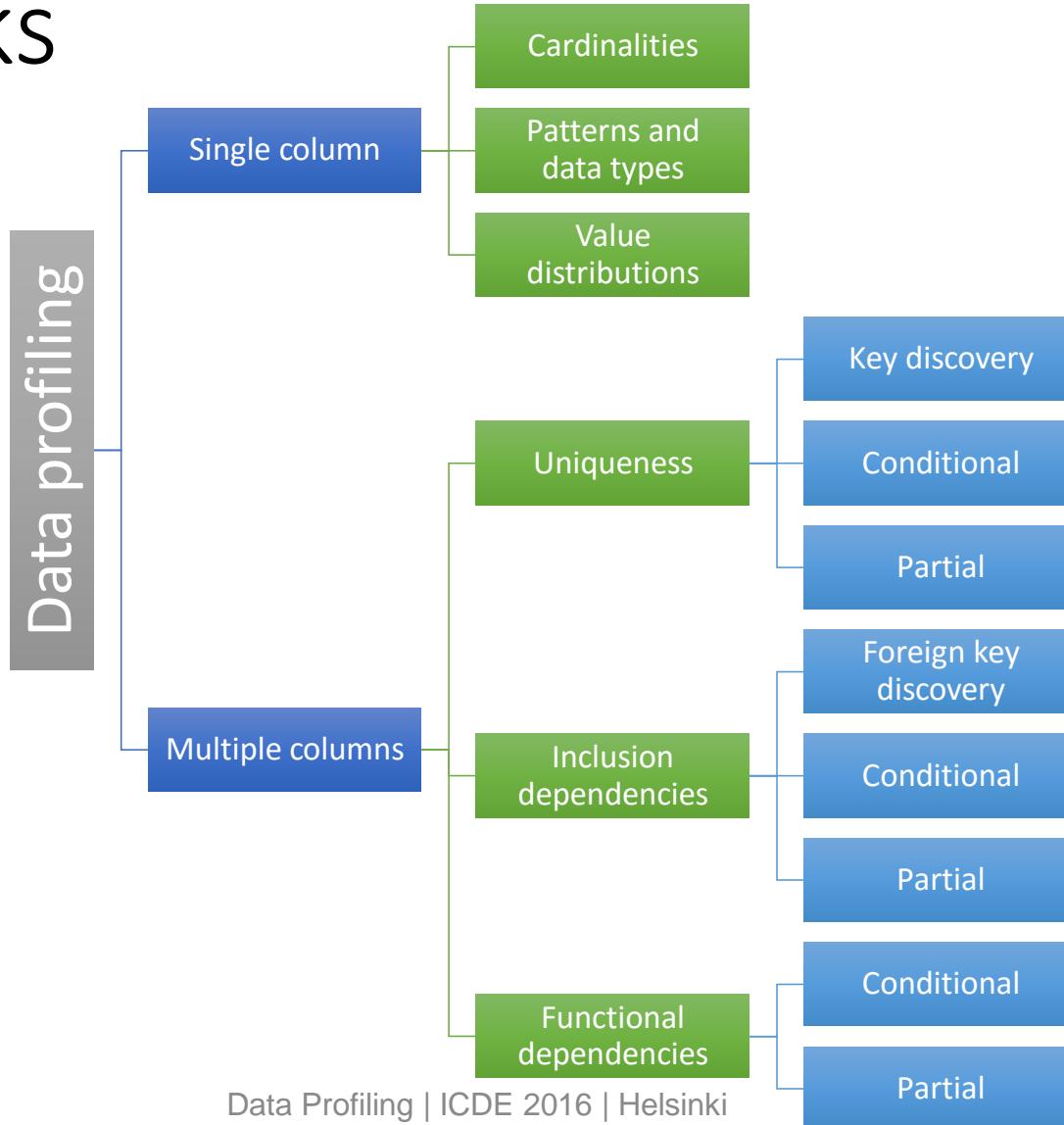
[Ted Johnson, Data Profiling, Encyclopedia of Database Systems, 2009]

- Data profiling is the set of activities and processes to determine the metadata about a given dataset.

- A fixed set of data profiling tasks / results



Classification of Traditional Profiling Tasks



Data Profiling vs. Data Mining

- Data profiling gathers technical metadata to support data management
- Data mining and data analytics discovers non-obvious results to support business management
- Data profiling results: information about columns and column sets
- Data mining results: information about rows or row sets
 - clustering, summarization, association rules, ...
- Rahm and Do on data cleaning
 - Profiling: Individual attributes
 - Mining: Multiple attributes

[Rahm and Do, Data Cleaning: Problems and Current Approaches, IEEE DE Bulletin, 2000]

Challenges of (Big) Data Profiling

- Large search space
 - Number of rows AND number of columns (and column combinations)
 - “Small” table with 100 columns:
$$2^{100} - 1 = 1,267,650,600,228,229,401,496,703,205,375$$

= 1.3 nonillion column combinations
- Large solution space: Exponential number of dependencies
- New data types and new data models
- New requirements: User-oriented, interactive, streaming
- Solutions: Scale up, scale out, scale in
- Better: Intelligent enumeration and aggressive pruning

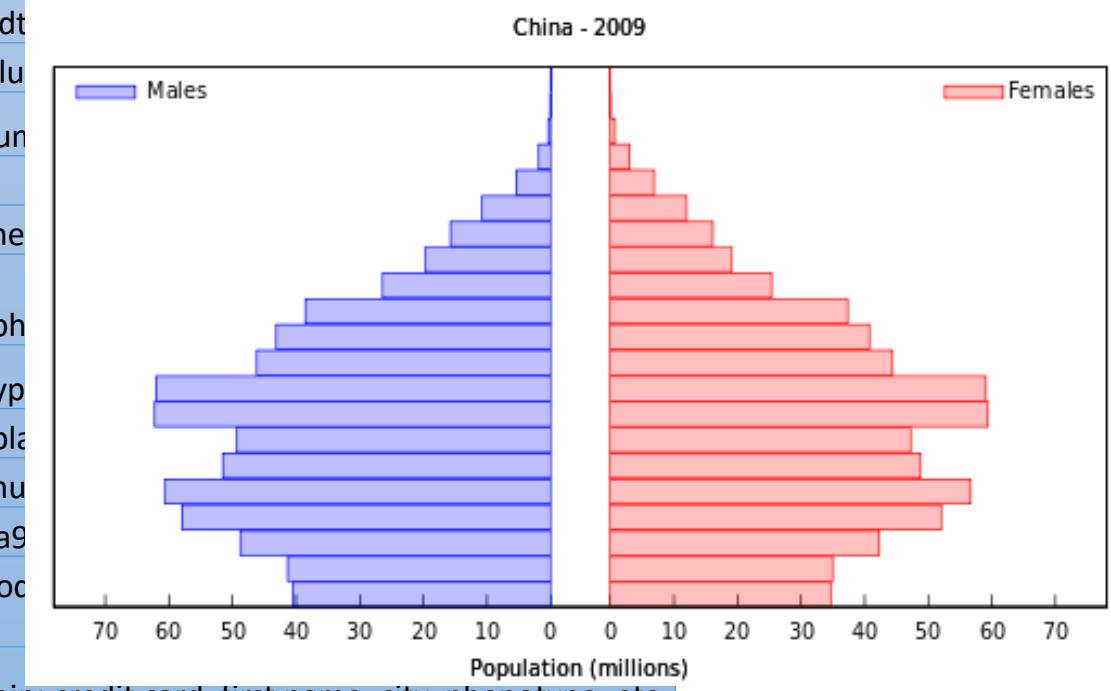
Use Cases for Profiling

- Query optimization
 - Counts and histograms
- Data cleansing
 - Patterns and violations
- Data integration
 - Cross-DB inclusion dependencies
- Scientific data management
 - Handle new datasets
- Data analytics
 - Profiling as preparation and for initial insights
 - Borderline to data mining
- Database reverse engineering

Basic Statistics

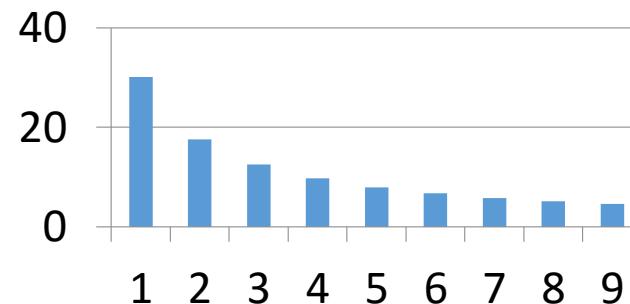
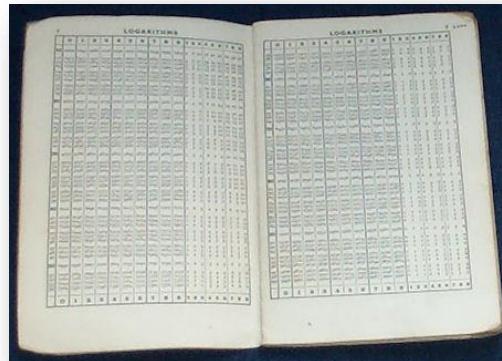


Cardinalities, Distributions, and Patterns



An Aside: Benford Law Frequency (“first digit law”)

- Statement about the distribution of first digits d in (many) naturally occurring numbers:
 - $P(d) = \log_{10}(d + 1) - \log_{10}(d) = \log_{10}(1 + 1/d)$



- Holds if $\log(x)$ is uniformly distributed



[Benford: The law of anomalous numbers". Proc. Am. Philos. Soc. 78 (4): 551–572, 1938]

Examples for Benford's Law

- Surface areas of 335 rivers
- Sizes of 3259 US populations
- 104 physical constants
- 1800 molecular weights
- 308 numbers contained in an issue of Reader's Digest
- Street addresses of the first 342 persons listed in American Men of Science

Heights of the 60 tallest structures

Leading digit	meters	
	Count	%
1	26	43.3%
2	7	11.7%
3	9	15.0%
4	6	10.0%
5	4	6.7%
6	1	1.7%
7	2	3.3%
8	5	8.3%
9	0	0.0%

In Benford's law
30.1%
17.6%
12.5%
9.7%
7.9%
6.7%
5.8%
5.1%
4.6%



Uses for Basic Statistics

- Traditional uses
 - Query optimization
 - Outlier/error detection
 - Visualize distribution
- Semantic uses
 - Categorization of attributes: Data types
 - Relevance of attributes: Completeness and quality
 - Semantics of attributes: Matching and cleansing

Unique Column Combinations



Unique Column Combinations

- Unique column
 - Only unique values
- Unique column combination
 - Only unique value combinations
 - Minimality: No subset is unique
- (Primary) key candidate
 - No null values
 - Uniqueness and non-null in one instance does not imply key: Only human can specify keys (and foreign keys)
- Meaning of NULL values?

Uses for UCCs

- Learn characteristics of a new data set
- Database management
 - Find a primary key
 - Find unique constraints
- Query optimization
 - Cardinality estimations for joins
- Find duplicates / data quality issues
 - If expected unique column combinations are not unique
 - Or with partial uniques

Inclusion Dependencies



Inclusion Dependencies

- $A \subseteq B$: All values in A are also present in B
- $A_1, \dots, A_i \subseteq B_1, \dots, B_j$:
All value combinations in A_1, \dots, A_i are also present in B_1, \dots, B_j
- Prerequisite for foreign key
 - Used across relations
 - Use across databases
 - But again: Discovery on a given instance, only user can specify for schema

Motivation for IND Discovery

- General insight into data
- Detect unknown foreign keys
- Example: PDB – Protein Data Bank
 - OpenMMS provides relational schema
 - 175 tables, 2705 attributes
 - Not a single foreign key constraint!
- Example: Ensembl – genome database
 - Shipped as MySQL dump files
 - More than 200 tables
 - Not a single foreign key constraint!
- Web tables: No schema, no constraints, but many connections

```
_pdbx_poly_seq_scheme.pdb_strand_id
_pdbx_poly_seq_scheme.pdb_ins_code
_pdbx_poly_seq_scheme.hetero
A 1 1 DC 1 1 1 DC C A . n
A 1 2 DC 2 2 2 DC C A . n
A 1 3 DG 3 3 3 DG G A . n
A 1 4 DT 4 4 4 DT T A . n
A 1 5 DA 5 5 5 DA A A . n
A 1 6 DC 6 6 6 DC C A . n
A 1 7 DG 7 7 7 DG G A . n
A 1 8 DT 8 8 8 DT T A . n
A 1 9 DA 9 9 9 DA A A . n
A 1 10 DC 10 10 10 DC C A . n
A 1 11 DG 11 11 11 DG G A . n
A 1 12 DG 12 12 12 DG G A . n
#
loop_
_refine_B_iso.class
_refine_B_iso.details
_refine_B_iso.treatment
_refine_B_iso.pdbx_refine_id
'ALL ATOMS' TR isotropic 'X-RAY DIFFRACTION'
'ALL WATERS' TR isotropic 'X-RAY DIFFRACTION'
#
loop_
_refine_occupancy.class
_refine_occupancy.treatment
_refine_occupancy.pdbx_refine_id
'ALL ATOMS' fix 'X-RAY DIFFRACTION'
'ALL WATERS' fix 'X-RAY DIFFRACTION'
#
loop_
_pdbx_version.entry_id
_pdbx_version.revision_date
_pdbx_version.major_version
_pdbx_version.minor_version
_pdbx_version.revision_type
_pdbx_version.details
116D 2008-05-22 3 2 'Version format compliant
116D 2011-07-13 4 0000 'Version format compliant
#
software_name          NHCLSO
```

Functional and other dependencies



Functional and Other Dependencies

- Functional dependency
 - „ $X \rightarrow A$ “: whenever two records have the same X values, they also have the same A values.
 - Multi-valued dependencies
 - Join dependencies
 - Order dependencies
 - `SELECT emp_name
FROM employees
ORDER BY rank, salary`
 - `SELECT emp_name
FROM employees
ORDER BY rank`
-
- salary
orders rank
- Remove
rank
- Replace with
salary (if index
only on salary)

emp_name	rank	salary
Smith	1	40k
Johnson	1	40k
Williams	1	45k
Brown	2	60k
Davis	2	60k
Miller	3	70k
Wilson	4	100k

Uses for FDs

- Schema design
 - Normalization
 - Keys
 - Data cleansing
 - Schema design and normalization
 - Key discovery
 - Data cleansing (especially partial/conditional FDs)
 - Anomaly detection
 - Data integrity constraints
 - Data curation rules
 - Query optimization:
Independence of column attributes
 - Index selection
- ... and genealogy research!

Functional Dependencies



Functional Dependencies

Person	Lineage	Hair	Religion	
			New gods	<p>Some Functional Dependencies:</p> <ol style="list-style-type: none"> 1. Person → Lineage 2. Person → Hair 3. Person → Religion 4. Lineage → Hair 5. Religion, Hair → Lineage 6. ...
			New Gods	<p>Ned Stark: „#4 looks like a reasonable quality constraint“</p>
			Old gods	<p>Ned Stark: „I believe Joffrey violates my database constraint.“</p>
			New gods	
			Old gods	

Properties of Dependencies



Partial Dependencies

- Aka. “approximate dependencies”
- INDs and FDs that do not perfectly hold
 - For all but 10 of the tuples
 - Only for 80% of the tuples
 - Only for 1% of the tuples
- Also for patterns, types, uniques, and other constraints
- Useful for: Data cleansing

Conditional Dependencies

- Given a partial IND or FD: For **which** part do they hold?
- Expressed as a condition over the attributes of the relation
- Problems:
 - Infinite possibilities of conditions
 - Interestingness:
 - Many distinct values: less interesting
 - Few distinct values: surprising condition – high coverage
- Useful for Integration
 - Cross-database cINDs

Other (Relaxed) Dependencies

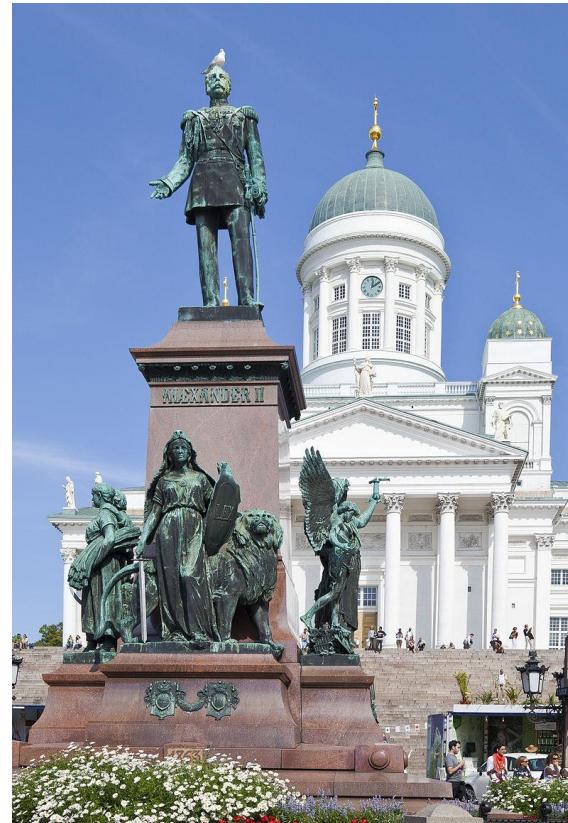
- Partial dependencies
- Approximate dependencies
- Conditional dependencies
- Matching dependencies
- Metric dependencies

RFD abbrev.	RFD name
ACOD	Approximate comparable dependency
ADD	Approximate differential dependency
AFD	Approximate functional dependency
COD	Comparable dependency
CFD	Conditional functional dependency
CFD ^p	CFD with built-in predicates
CFD ^c	CFD with cardinality constraints and synonym rules
CMD	Conditional matching dependency
CSD	Conditional sequential dependency
CD	Constrained functional dependency
DD	Differential dependency
eCFD	Extended conditional functional dependency
FFD	Fuzzy functional dependency
MD	Matching dependency
MFD	Metric functional dependency
ND	Neighborhood dependency
NuD	Numerical dependency
OD	Order dependency
OD _K	OD satisfied within bound k
ODEA	OD satisfied almost everywhere
OFD	Ordered functional dependency
PD	Partial determination
POD	Polarized order dependencies
prefD	Preference functional dependency
PAC	Probabilistic approximate constraint
pFD	Probabilistic functional dependency
PUD	Purity dependency
RUD	Roll-up dependency
SD	Sequential dependency
SFD	Similarity functional dependency
soft FD	Soft functional dependency
IMFD	Imprecise functional dependency
XCFD	XML conditional functional dependency
$\sigma\theta$ xFD	XML FD with σ and θ approximation

[Caruccio, Deufemia, Polese: Relaxed Functional Dependencies - A Survey of Approaches. TKDE '16]

Tutorial Overview

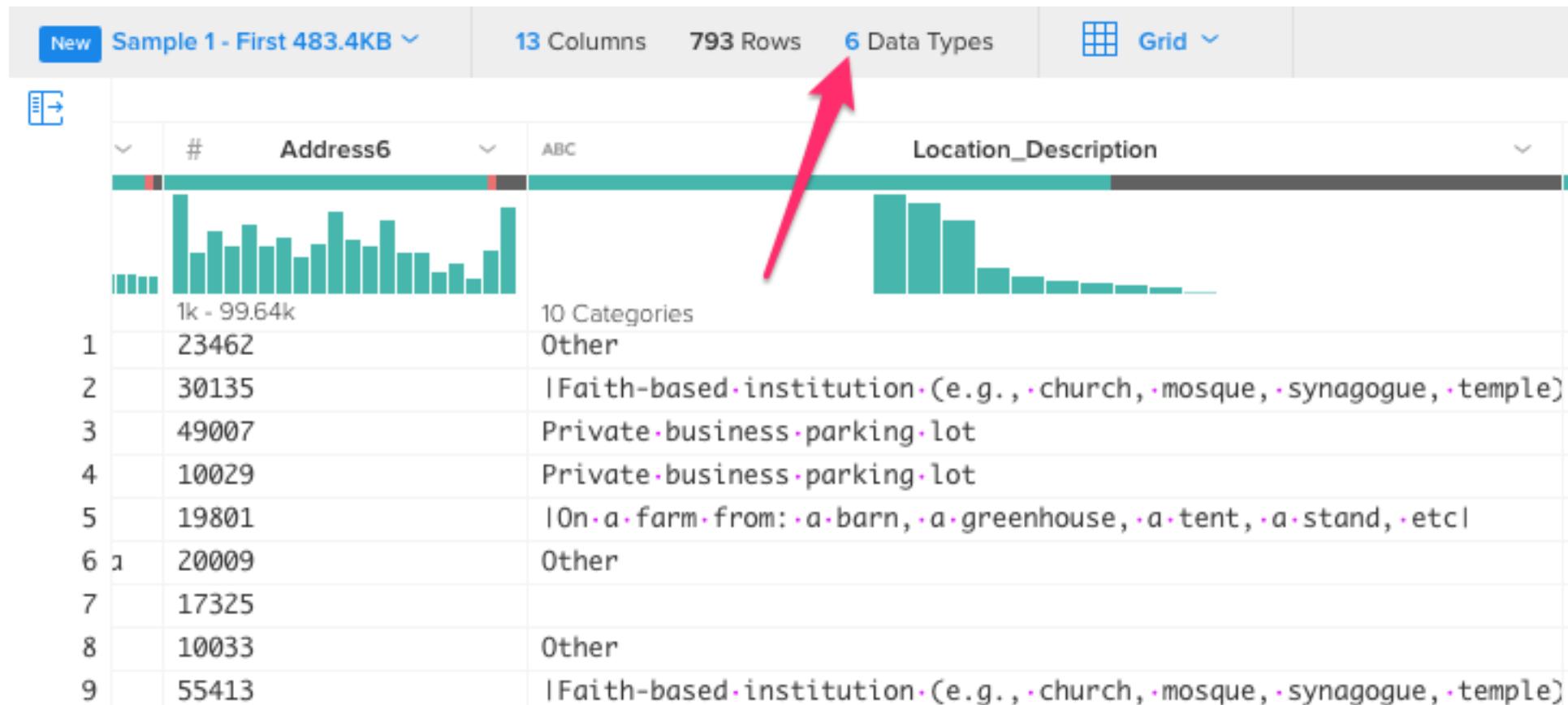
- Motivation
 - Task classification
 - Use cases
- Tools
 - Research and industry
 - Shortcomings
- Single and Multiple Column Analysis
 - Cardinalities and datatypes
 - Co-occurrences and summaries
- Dependencies
 - UCCs, INDs, FDs
 - and their discover algorithms
- Outlook
 - Functionality
 - Semantics



Tools in Industry



Trifacta



Open Refine

Google refine MGH / TeamSite Pages Export - Subset [Permalink](#)

Facet / Filter Undo / Redo 12

Refresh Reset All Remove All

LAST MODIFIED DATE change reset
2008-08-18 00:05:32 — 17:15:33

Author change
122 choices Sort by: name count Cluster
mk855 59
ks191 51
dp682 43
ea848 39

Subsection change
198 choices Sort by: name count Cluster
bhi 106
heartcenter 93
gastroenterology 89
geriatrics 83
transplant 81
nephrology 78
thoracicsurgery 75
palliativecare 73
imaging 70
digestive 69
regenmed 69
radiology 66

5679 rows

Show as: rows records Show: 5 10 25 50 rows « first < previous 1 - 50 next > last »

All	PAGE URL	DCT TYPE	Number of Versi	PAGE TITLE	Autho
1.	http://www.massgeneral.org/search.aspx	MGH_FacetedBrowse/fb_googleSearch	1		awb9
2.	http://www.massgeneral.org/_t.aspx	MGH_HomePages/hp_3illustration	1	Home	jy915
3.	http://www.massgeneral.org/partners.aspx	MGH_InteriorPages/ip_1_2	9	Partners HealthCare	jo860
4.	http://www.massgeneral.org/pngu_staff.aspx	MGH_InteriorPages/ip_1_2	1	Psychiatric & Neurodevelopment Genetics Unit (PNGU)	khs19
5.	http://www.massgeneral.org/FUS_TLS.aspx	MGH_InteriorPages/ip_3	1	FUS/TLS	mjr46
6.	http://www.massgeneral.org/TDP_43_TARDBP.aspx	MGH_InteriorPages/ip_3	1	TDP 43 TARDBP	mjr46
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8.	http://www.massgeneral.org/proto.aspx	MGH_InteriorPages/ip_1_2	10	Proto Magazine	nag16
9.	http://www.massgeneral.org/PCI_Newsletters.aspx	MGH_InteriorPages/ip_3	2	pci newsletters	sh550
10.	http://www.massgeneral.org/ip2c.aspx	MGH_InteriorPages/ip_2customflash	4	testing page again	jy915
11.	http://www.massgeneral.org/agenda_CSAA.aspx	MGH_InteriorPages/ip_3	5	HMS Seminar Agenda	ks191
12.	http://www.massgeneral.org/Magnet_recognition_notice.aspx	MGH_InteriorPages/ip_1_2	3	Mass General seeks feedback for Magnet recognition	vf045
13.	http://www.massgeneral.org/testing1235.aspx	MGH_InteriorPages/ip_3	1	asdf	jo860
14.	http://www.massgeneral.org/externallink.aspx	MGH_InteriorPages/ip_3	14	externallink class (IE) fix	jo860
15.	http://www.massgeneral.org/test.aspx	MGH_InteriorPages/ip_1_2	11	Weight Center Medical Management Program	jy915

Uses Cases Covered By Industrial Tools

Tool	Statistics	Patterns	Data types	Uniques	Restricted data types	Restricted number of columns
Attacama , DQ Analyzer	✓	✓		✓		
IBM , InfoSphere Information Analyzer	✓	✓		✓	✓	
Microsoft SQL Server Data Profiling Task	✓	✓				✓
Oracle Enterprise Data Quality	✓	✓				
Paxata Adaptive Preparation	✓					
SAP Information Steward	✓	✓	✓		✓	
Splunk Enterprise/Hunk			✓			✓
Talend Data Profiler	✓	✓			✓	
Trifecta	✓	✓	✓			
Tamr	✓			✓		
OpenRefine						

Tools in Research



RuleMiner

Dataset: Tax **Browse...**

Approximate Threshold: **0.01** Constant Frequency: **0**

Formula **Linguistics**

Coverage : **0.40** Filtering:
FDs

Succinctness: **0.60**

not(t1.areacode=t2.areacode & t1.phone=t2.phone)	<input checked="" type="checkbox"/> Yes	<input checked="" type="checkbox"/> No
not(t1.city!=t2.city & t1.zip=t2.zip)	<input checked="" type="checkbox"/> Yes	<input checked="" type="checkbox"/> No
There cannot exist two tuples t_1, t_2 in the dataset, such that they have different city ,and they have same zip		
not(t1.state=t2.state & t1.haschild=t2.haschild & t1.childexemp!=t2.childexemp)	<input checked="" type="checkbox"/> Yes	<input checked="" type="checkbox"/> No
not(t1.state=t2.state & t1.maritalstatus=t2.maritalstatus & t1.singleexemp!=t2.singleexemp)	<input checked="" type="checkbox"/> Yes	<input checked="" type="checkbox"/> No
not(t1.state=t2.state & t1.salary=t2.salary & t1.rate!=t2.rate)	<input checked="" type="checkbox"/> Yes	<input checked="" type="checkbox"/> No
not(t1.state=t2.state & t1.salary>t2.salary & t1.rate<t2.rate)	<input checked="" type="checkbox"/> Yes	<input checked="" type="checkbox"/> No
not(t1.phone=t2.phone)	<input checked="" type="checkbox"/> Yes	<input checked="" type="checkbox"/> No
not(t1.fname=t2.fname)	<input checked="" type="checkbox"/> Yes	<input checked="" type="checkbox"/> No

Data **Example**

Negative Example:

tid	fname	lname	areacode	phone	city	state	zip	maritalstatus	haschild	salary	rate	singleexemp
1	Mark	Ballin	304	2327667	Anthony	AR	25813	S	Y	5000	3	2000
8	Marcelino	Nuth	304	5404707	Kyle	WV	25813	M	N	10000	4	0

Positive Examples:

tid	fname	lname	areacode	phone	city	state	zip	maritalstatus	haschild	salary	rate	singleexemp
1	Mark	Ballin	304	2327667	Anthony	WV	25813	S	Y	5000	3	2000
8	Marcelino	Nuth	304	5404707	Kyle	WV	25813	M	N	10000	4	0

tid	fname	lname	areacode	phone	city	state	zip	maritalstatus	haschild	salary	rate	singleexemp
1	Mark	Ballin	304	2327667	Anthony	AR	25813	S	Y	5000	3	2000
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tid	fname	lname	areacode	phone	city	state	zip	maritalstatus	haschild	salary	rate	singleexemp
1	Mark	Ballin	304	2327667	Anthony	AR	10000	S	Y	5000	3	2000
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tid	fname	lname	areacode	phone	city	state	zip	maritalstatus	haschild	salary	rate	singleexemp
1	Mark	Ballin	304	2327667	Anthony	AR	25813	S	Y	5000	3	2000
8	Marcelino	Nuth	304	5404707	Kyle	WV	10000	M	N	10000	4	0

ProLOD++

ProLOD++

Overview Graph Analysis Properties Inverse Properties Association Rules Synonyms Key Discovery

Graphs / Pattern 1

- ▶ DailyMed (11,271)
- ▶ DBpedia (4,222,586)
- ▼ **Diseasome (9,047)**
 - ▼ ■ diseases (4,213)
 - ▼ ■ genes (9,743)
- ▶ DrugBank (19,694)
- ▶ LinkedMDB (631,003)

Pattern: 41

Nodes: 5

Edges: 5

Diameter: 2

Statistics:

Pattern:	41
Nodes:	5
Edges:	5
Diameter:	2

Class distribution:

Class	Count
genes	4
diseases	2
unknown	1

Tools in Research

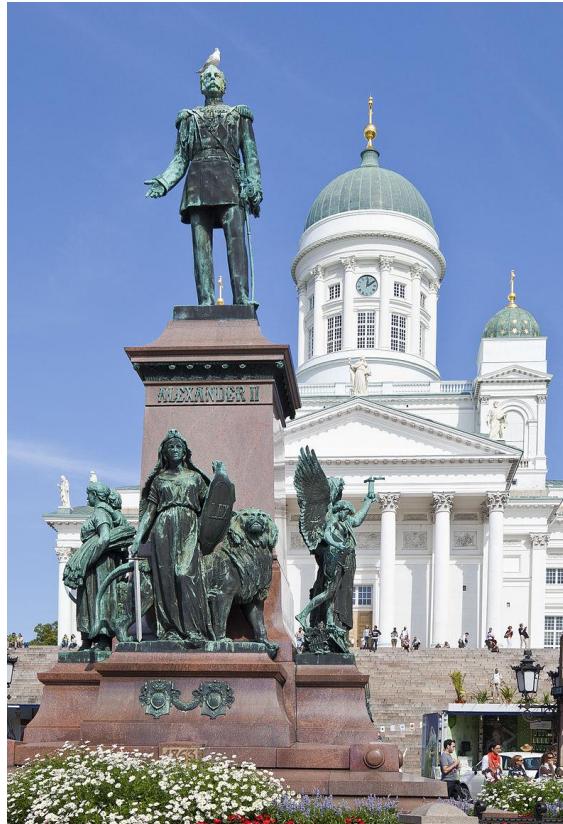
Tool	Main purpose	Statistics	Patterns	Data types	Uniques	Dependencies	Data Mining
Bellmann	Data quality browser	✓			✓		
Potter's Wheel	ETL tool	✓	✓				
Data Auditor	Rule discovery						
RuleMiner	Dependency discovery					✓	
MADLib	Machine learning	✓				✓	
Metanome	Data profiling	✓			✓		
ProLOD++	Profiling and Mining	✓	✓		✓	✓	✓

Shortcomings

- No “real” profiling tool
- Tools focus on “easy” problems:
 - Statistics
 - Single column or “few” column dependencies
 - Many industry tools use SQL instead of optimized algorithms
- No tool covers all types of meta-data
- Management of large meta-data results
 - Summarizing meta-data
 - Ranking meta-data based on relevance

Tutorial Overview

- Motivation
 - Task classification
 - Use cases
- Tools
 - Research and industry
 - Shortcomings
- **Single and Multiple Column Analysis**
 - Cardinalities and datatypes
 - Co-occurrences and summaries
- Dependencies
 - UCCs, INDs, FDs
 - and their discover algorithms
- Outlook
 - Functionality
 - Semantics



Single Column Analysis



Cardinalities and distributions

- Number of values
- Number of distinct values
- Number of NULLs
- MIN and MAX value



Count(*)
count(distinct X),
count (X) where X=null

For (value in column)
If (value>max)
max=value

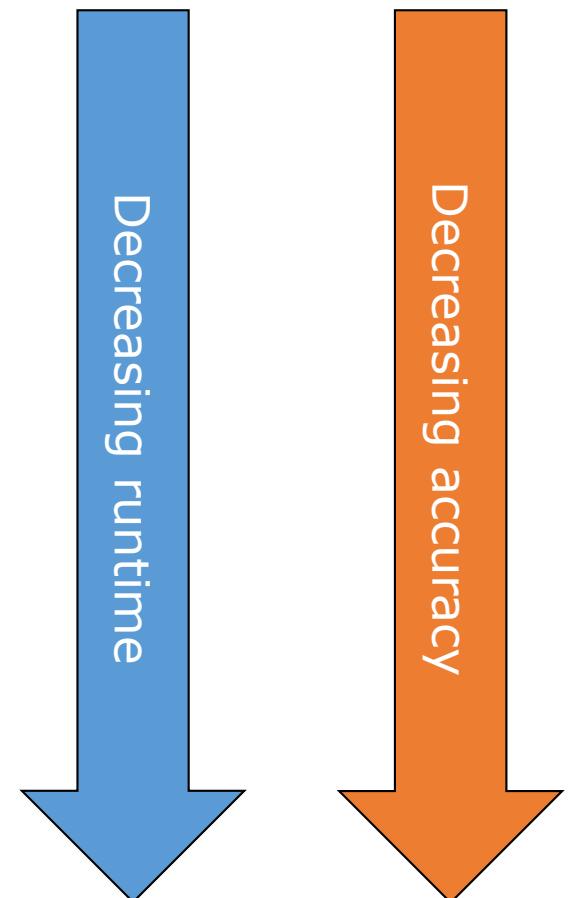
- Histograms
- Probability distribution for numeric values
- Detect whether data follows some well-known distribution



Bottleneck is sorting the data

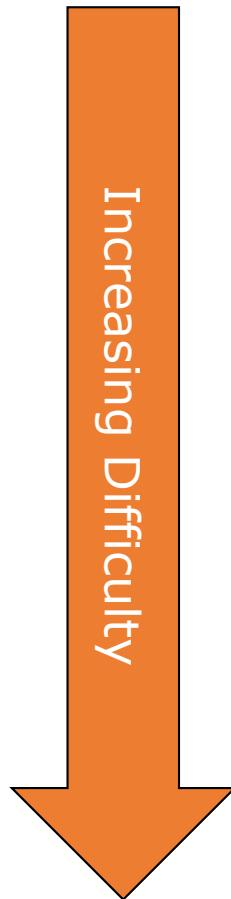
Count distinct in sublinear time and space?

- Linear Counting
 - [Whang, Vander-Zanden, Taylor: A linear-time probabilistic counting algorithm for database applications. TODS, 1990]
- Stochastic Averaging
 - [Flajolet, Martin: Probabilistic counting algorithms for data base applications. JCSS, 1985]
- Loglog Algorithm
 - [Durand, Flajolet: Loglog counting of large cardinalities. Algorithms-ESA, 2003]
- SuperLogLog Algorithm
 - [Durand, Flajolet: Loglog counting of large cardinalities. Algorithms-ESA, 2003]
- HyperLogLog Algorithm
 - [Flajolet, Fusy, Gandouet, Meunier: Hyperloglog: the analysis of a near-optimal cardinality estimation algorithm. DMTCS, 2008]



Data types and value patterns

- String vs. number
- String vs. number vs. date
- Categorical vs. continuous
 - Days of the week vs. measurements
- SQL data types
 - CHAR, INT, DECIMAL, TIMESTAMP, BIT, CLOB, ...
- Domains
 - VARCHAR(12) vs. VARCHAR (13)
- XML data types
 - More fine grained
- Regular expressions `(\d{3})-(\d{3})-(\d{4})-(\d+)`
- Semantic domains
 - Adress, phone, email, first name



Multi Column Analysis



Frequencies, Rules, Correlations

- Frequencies:
 - Which values co-occur with each other?
- Rules:
 - Which values depend on a specific value?
- Correlations:
 - Which values correlate?
 - Which values substitute each other?

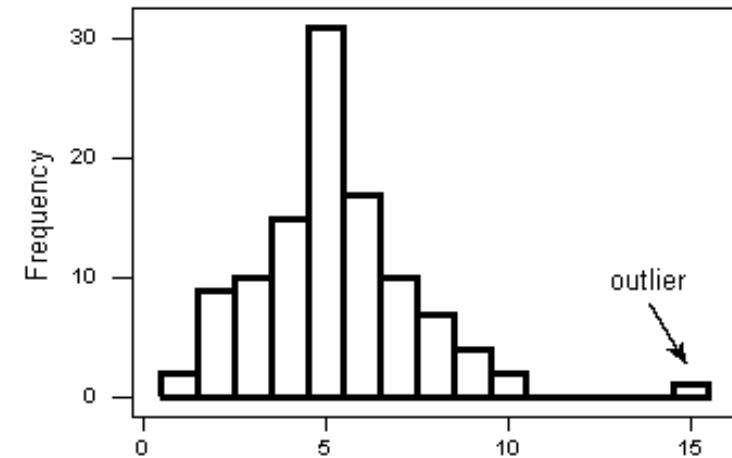


Core step: Frequent Itemset Mining

- Origin: Transactional Analysis
 - Which products have been bought together?
- Main step:
 - Find frequencies for all item combinations
- Optimization:
 - Find frequencies for all relevant item combinations, i.e., item combinations with minimum support
- Algorithms:
 - Apriori [\[Aggrawal, Srikant: fast Algorithms for Mining Association rules, VLDB'94\]](#)
 - FP-Growth [\[Han, Pei, Yin: Mining frequent patterns without candidate generation, SIGMOD'00\]](#)
 - ..
 - Survey: [\[Hipp, Guentzer, Nakhaeizadeh: Algorithms for Association Mining – A General Survey and Comparison, KDD'00\]](#)

Outlier detection

- Low-frequent values
- Structural outliers
 - Wrong value representations, e.g.:
 - CA instead of California
- Numerical outliers
 - E.g., according to Gaussian distribution
- Outlier combinations
 - Co-occurrence analysis
- Survey: [\[Hodge, Austin: A survey of outlier detection methodologies, AI'04\]](#)



Sketches and Summaries

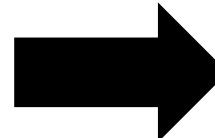
- Use cases:
 - Assess column similarity
 - Dimension reduction
 - Data stream samples
- Techniques:
 - Sampling
 - Hashing:
 - Minhash [Broder: Compression and Complexity of Sequences, 1997]
 - LSH [Gionis, Indyk, Motwani: Similarity search in high Dimensions via hashing, VLDB'99]
 - Sketches [Cormode, Garofalakis, Haas, Jermaine: Synopses for Massive Data:Samples, Histograms, Wavelets, Sketches, FTD'12]

Column Similarity:

$$\text{Jaccard}(C_1, C_2) = \text{intersect}(C_1, C_2) / \text{Union}(C_1, C_2)$$

- N^2 pairwise comparisons
- Reduce dimension through Minhash:
 - Find a hash function $h(\cdot)$ such that:
 - If $\text{sim}(C_1, C_2)$ is high, then with high prob. $h(C_1) = h(C_2)$
 - If $\text{sim}(C_1, C_2)$ is low, then with high prob. $h(C_1) \neq h(C_2)$
 - Estimate similarity by applying k different $h_i(\cdot)$
 - Transform table into a Boolean matrix

Residence (A)	Country (B)	Birthplace (C)
Helsinki	Finland	Oslo
Oslo	Germany	Copenhagen
Berlin	Denmark	Helsinki



Values	A	B	C
Helsinki	1	0	1
Oslo	1	0	1
Berlin	1	0	0
Finland	0	1	0
Germany	0	1	0
Denmark	0	1	0
Copenhagen	0	0	1

Minhash Example

- Simulate hash through permutation of row numbers
- Pick smallest row number where matrix value equals 1

Values	A	B	C
Helsinki	1	0	1
Oslo	1	0	1
Berlin	1	0	0
Finland	0	1	0
Germany	0	1	0
Denmark	0	1	0
Copenhagen	0	0	1

h1	h2	h3
1	7	5
2	4	6
3	1	7
4	5	2
5	3	3
6	6	4
7	2	1

Hash	A	B	C
h1	1	4	1
h2	1	3	2
h3	5	2	1

$$\text{sim}(A,B) = 0$$

$$\text{sim}(A,C) = 0.33$$

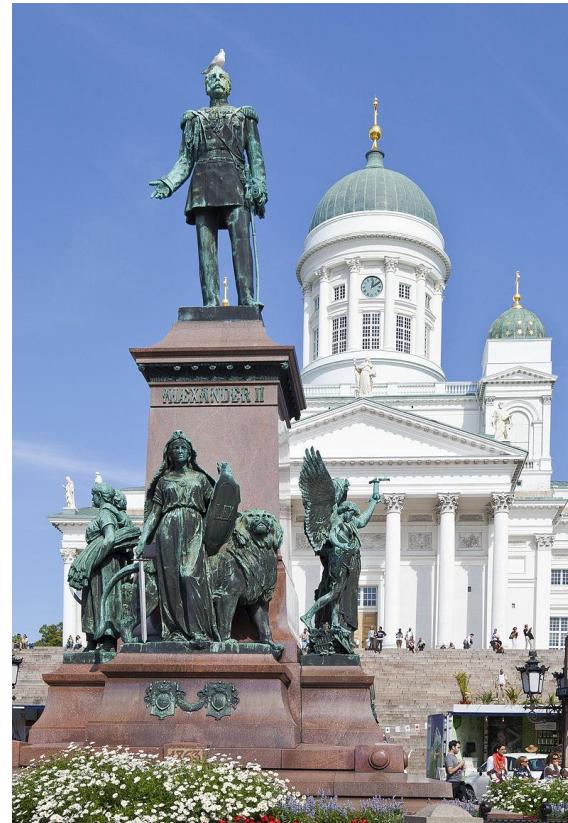
$$\text{sim}(B,C) = 0$$

Single & Multi-Column Analysis

- Cardinalities
- Data types
- Patterns
- Co-occurrences
- Sketches, summaries
-
- Strong overlap with data mining
- Most of them:
 - Not very complex but approximations needed on big data

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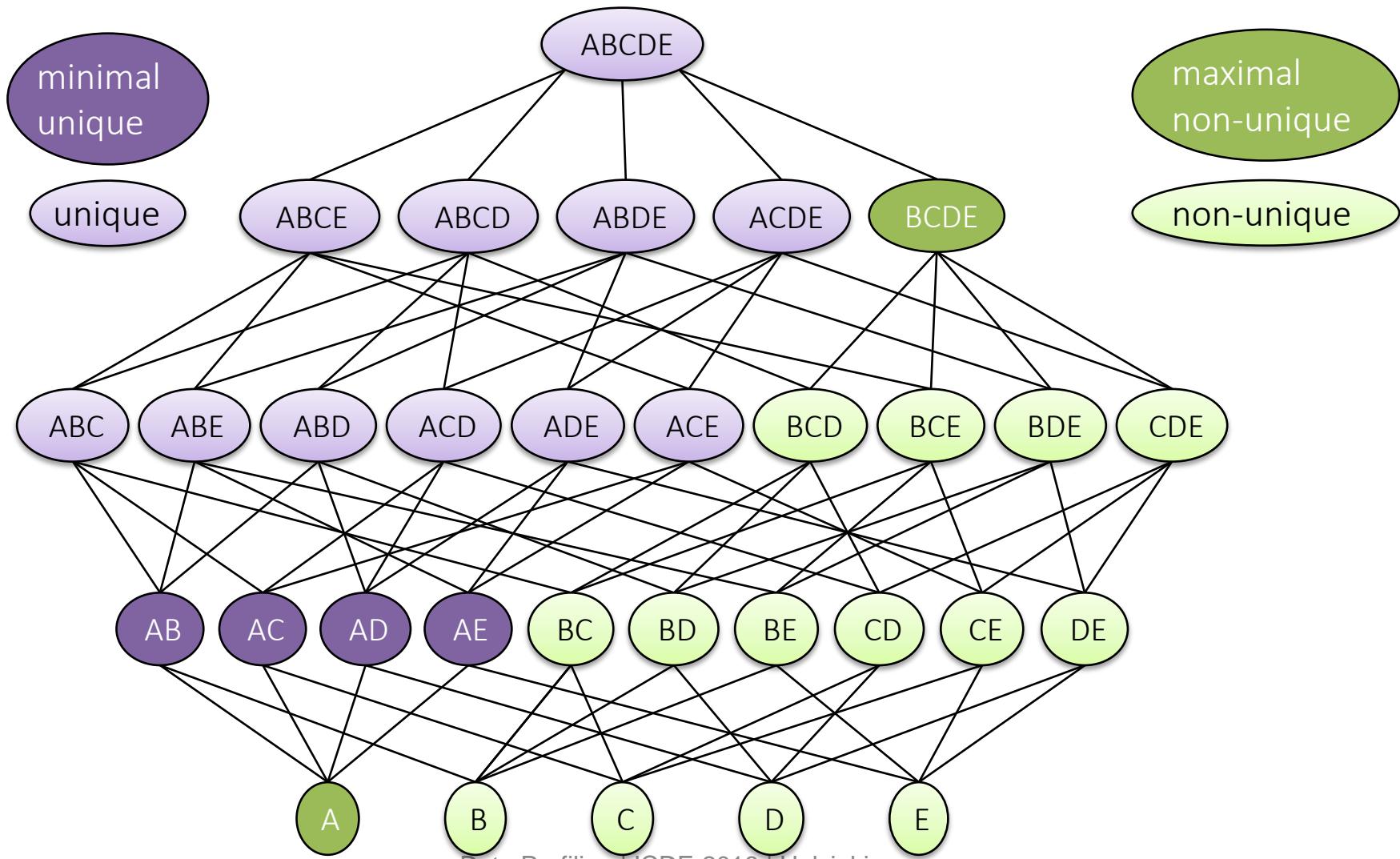


©2005 JESSICA AND JOHN WILLIAMS

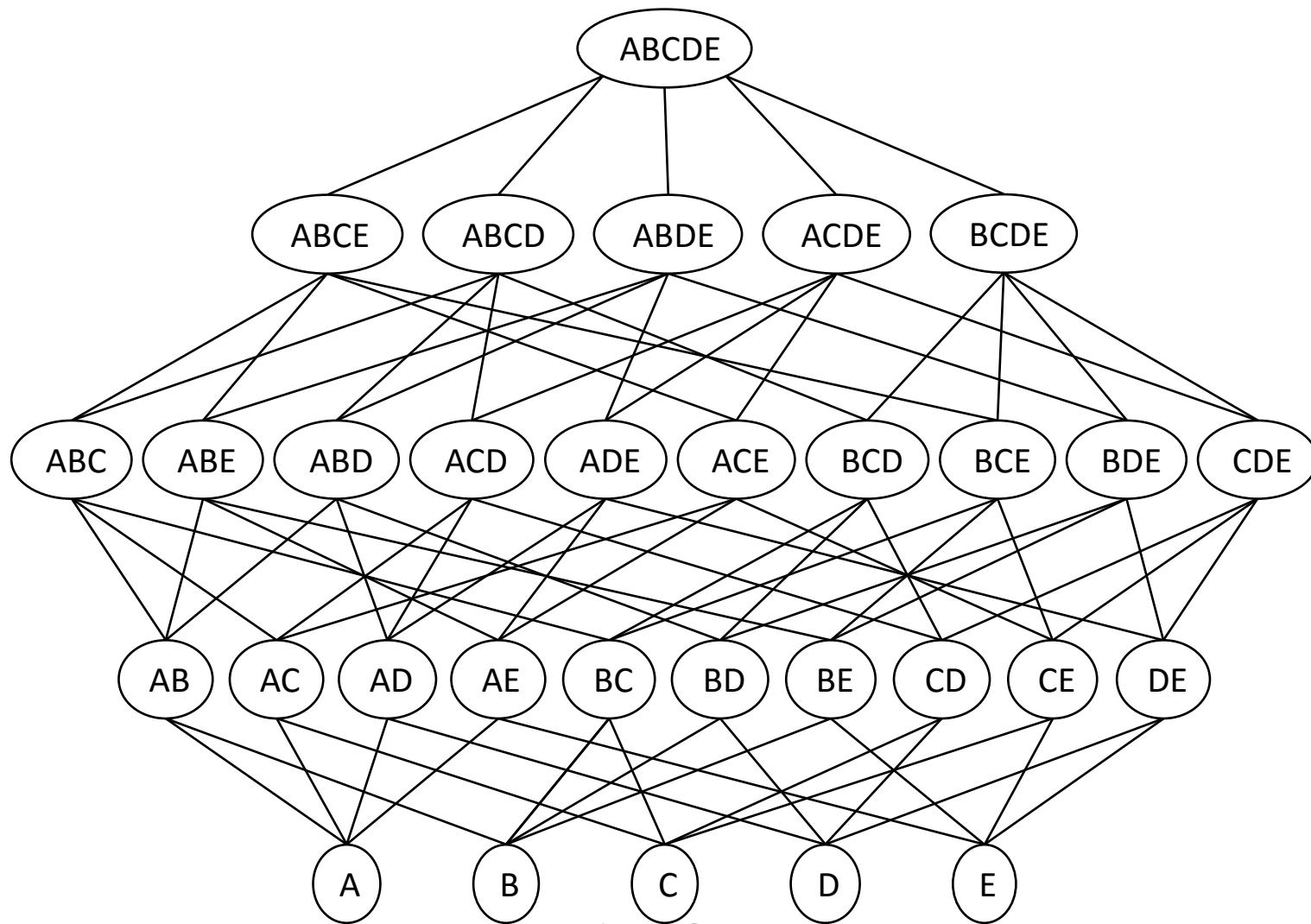
UNIQUE

JUST BECAUSE YOU ARE UNIQUE DOES NOT MEAN YOU ARE USEFUL

Result of algorithm



Challenge: Exponential search space



$$\binom{5}{5} = 1$$

$$\binom{5}{4} = 5$$

$$\binom{5}{3} = \frac{5 \cdot 4}{2}$$

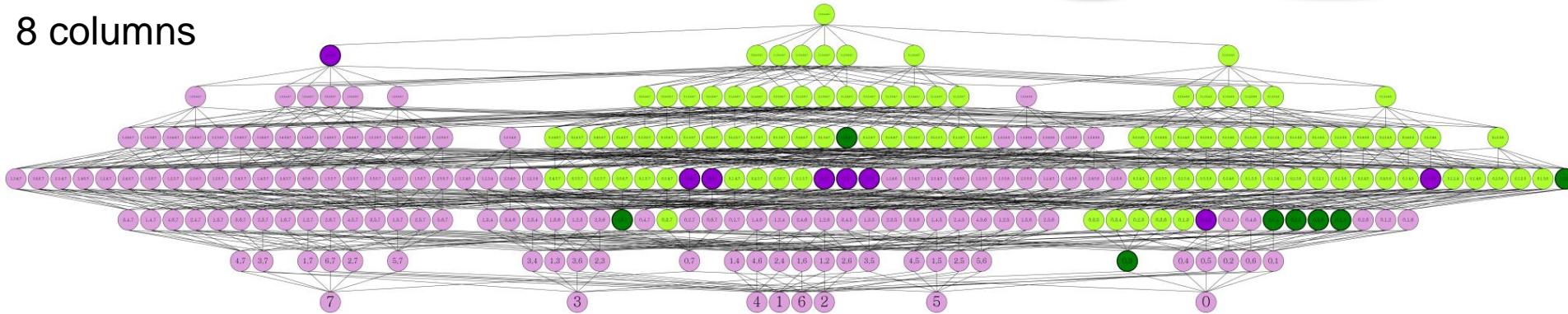
$$\binom{5}{2} = \frac{5 \cdot 4 \cdot 3}{2 \cdot 3}$$

$$\binom{5}{1} = \frac{5 \cdot 4 \cdot 3 \cdot 2}{2 \cdot 3 \cdot 4}$$

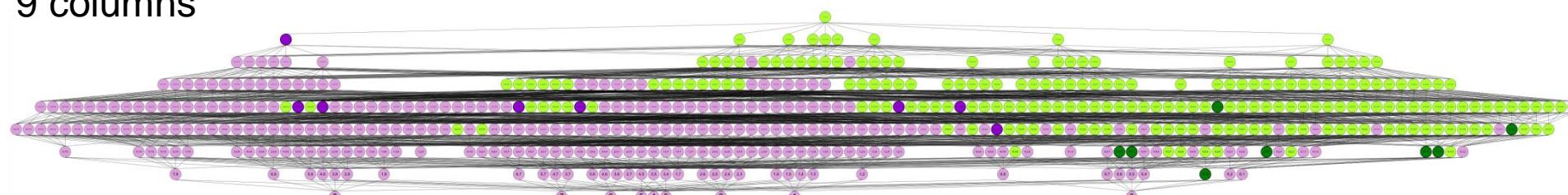
TPCH line item

unique non-unique

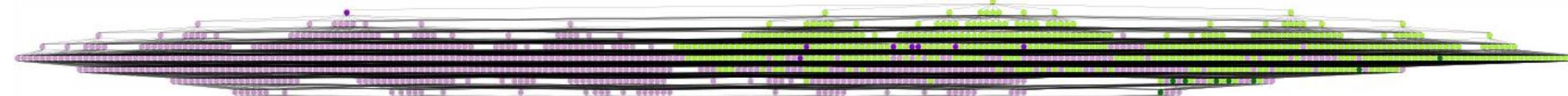
8 columns



9 columns



10 columns



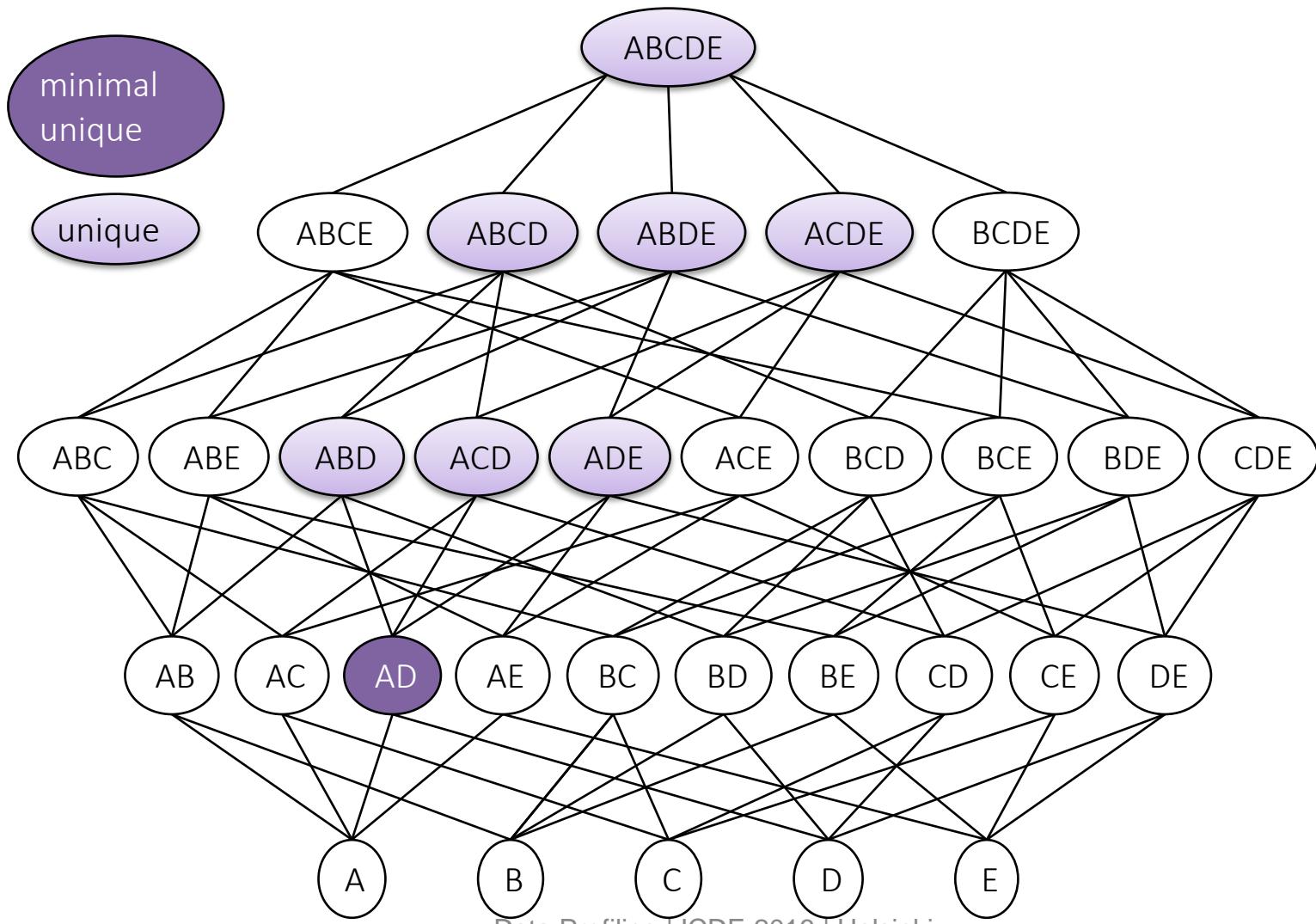
Computational feasibility

- For a lattice over n columns
 - $\binom{n}{k}$ combinations of size k
 - All combinations: $2^n - 1$ (let's ignore -1 for the remaining slides)
 - Largest solution set: $\binom{n}{n/2}$ minimal uniques are of size $\frac{n}{2}$
$$\binom{n}{k} \in \Theta(n^k) \Rightarrow \binom{n}{n/2} \in \Theta(n^n)$$
 - Verifying minimality, requires to check also all combinations of size $\frac{n}{2} - 1$
 - Adding a column doubles search space (and vice versa)

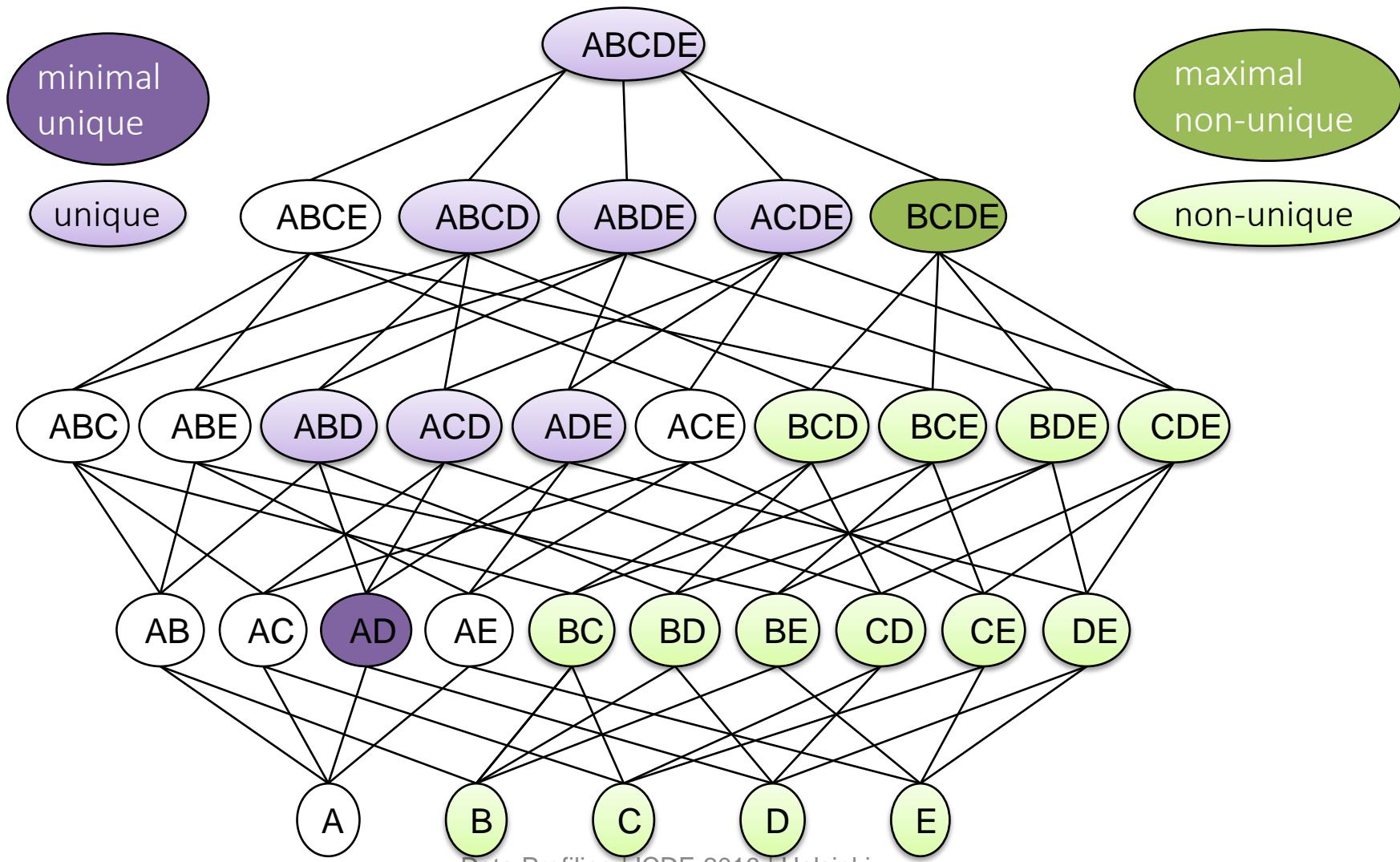
Pruning with uniques #2

- Pruning: inferring the type of a combination without actual verification
- If A is unique, supersets must be unique
- Finding a unique column prunes half of the lattice
 - Remove column from initial data set and restart
- Finding a unique column pair removes a quarter of the lattice
 - In general, the lattice over the combination is removed
- The pruning power of a combination is reduced by prior findings
 - AB prunes a quarter
 - BC additionally prunes only one eighth
 - ABC was already pruned by AB and constitutes already one eighth

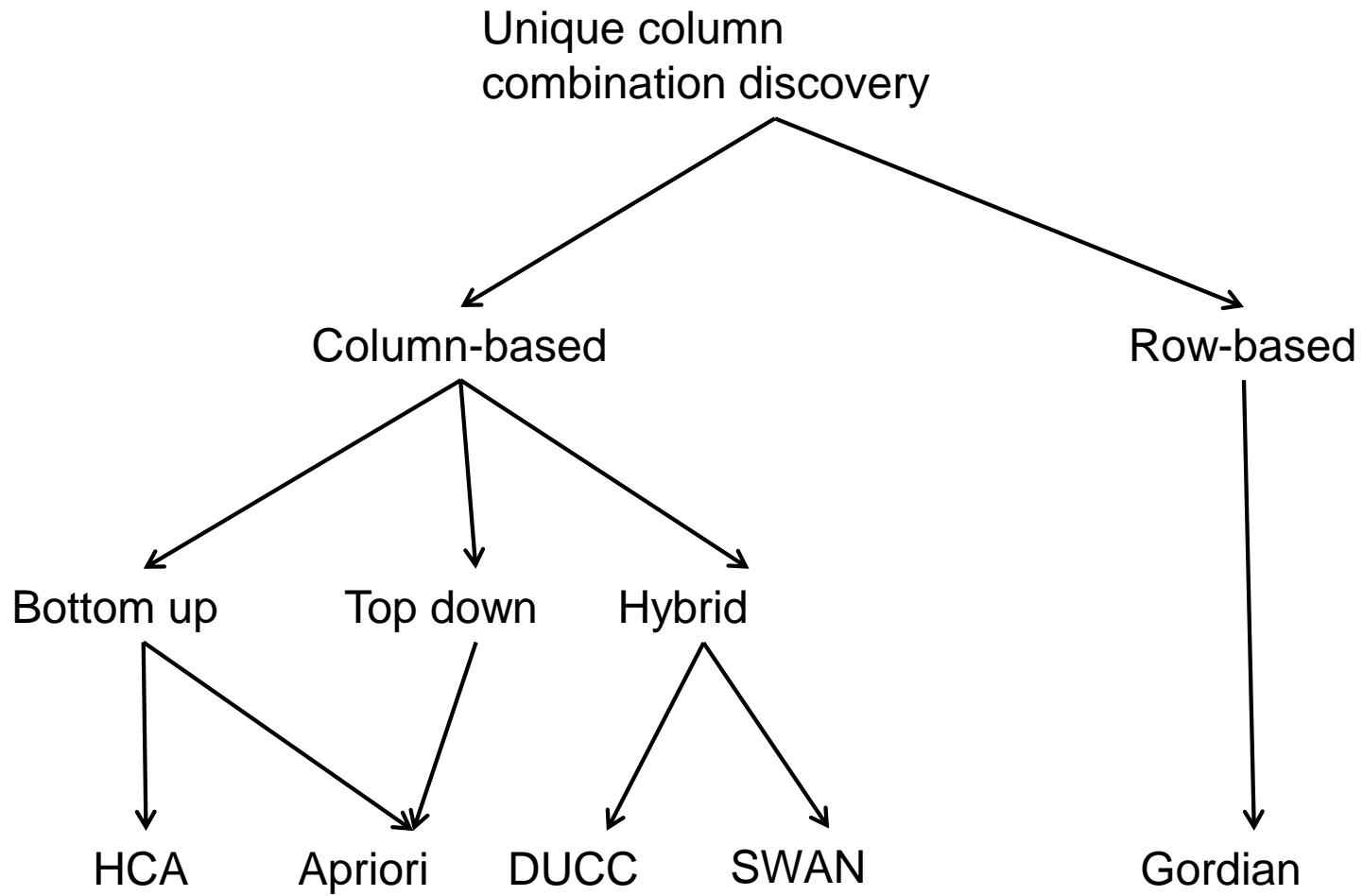
Pruning effect of a pair



Pruning both ways



Discovery Algorithms



Column-based algorithms

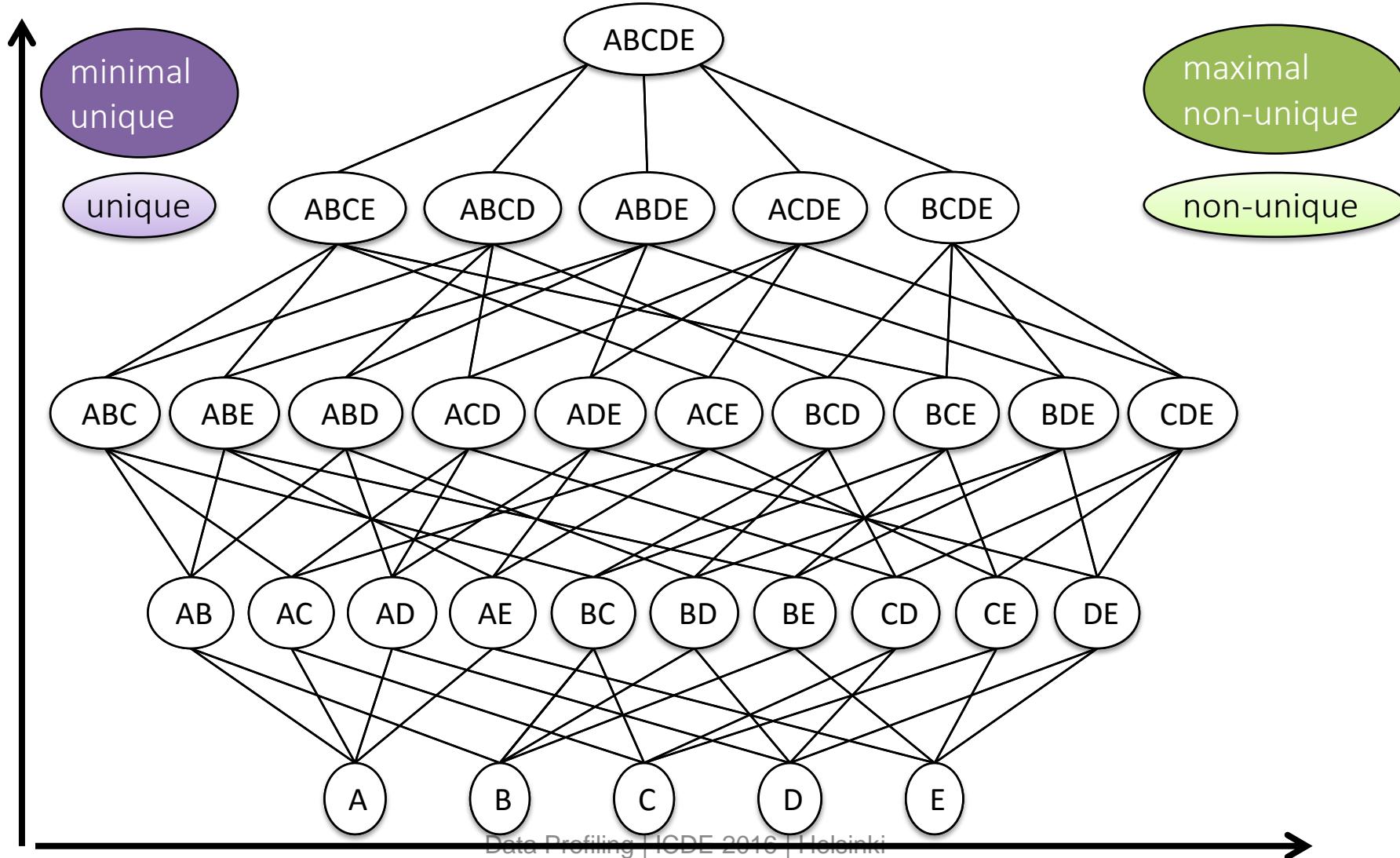
- Traverse through lattice
- Check for uniqueness
 - Different approaches possible
 - Use database backend and distinctness query
 - `SELECT COUNT(DISTINCT A, B, C) FROM R`
 - Compare with row-count
 - Position list indexes (explained later)
 - For now, check is blackbox
- Prune lattice accordingly

Apriori-based

[Giannella, Wyss: Finding minimal keys in a relation instance. (1999)]

- Basic idea:
 - Using the state of combinations of size k
 - We need to visit only unpruned combinations of size $k+1$
 - Add non-unique columns to combination of size k
- Start with individual columns
- Check pairs of non-unique columns
- Check triples of non-unique pairs ...
- Terminate if no new combinations can be enumerated

Apriori visualized



Characteristics of Apriori

- Works well for small uniques
 - Bottom-up checks columns first
- Best case: all columns are unique
 - n checks
- Worst case: no uniques = one duplicate row
 - 2^n checks
- Apriori is exponential in n

Extensions

- Top-down
 - Start from top and go down
 - Performs better if solution set is high up
 - Candidate pruning becomes more tricky
- Hybrid [\[Giannella, Wyss: Finding minimal keys in a relation instance. \(1999\)\]](#)
 - Combine bottom-up and top-down
 - Interleave checks
 - Works well if solution set has many small and large combinations
 - Worst case: solution set in the middle
- Statistics-based extensions [\[Abedjan, Naumann: Advancing the discovery of unique column combinations, CIKM'11\]](#)
 - More sophisticated candidate generation
 - Uses histograms for pruning
 - Finds and uses functional dependencies on-the-fly

DUCC

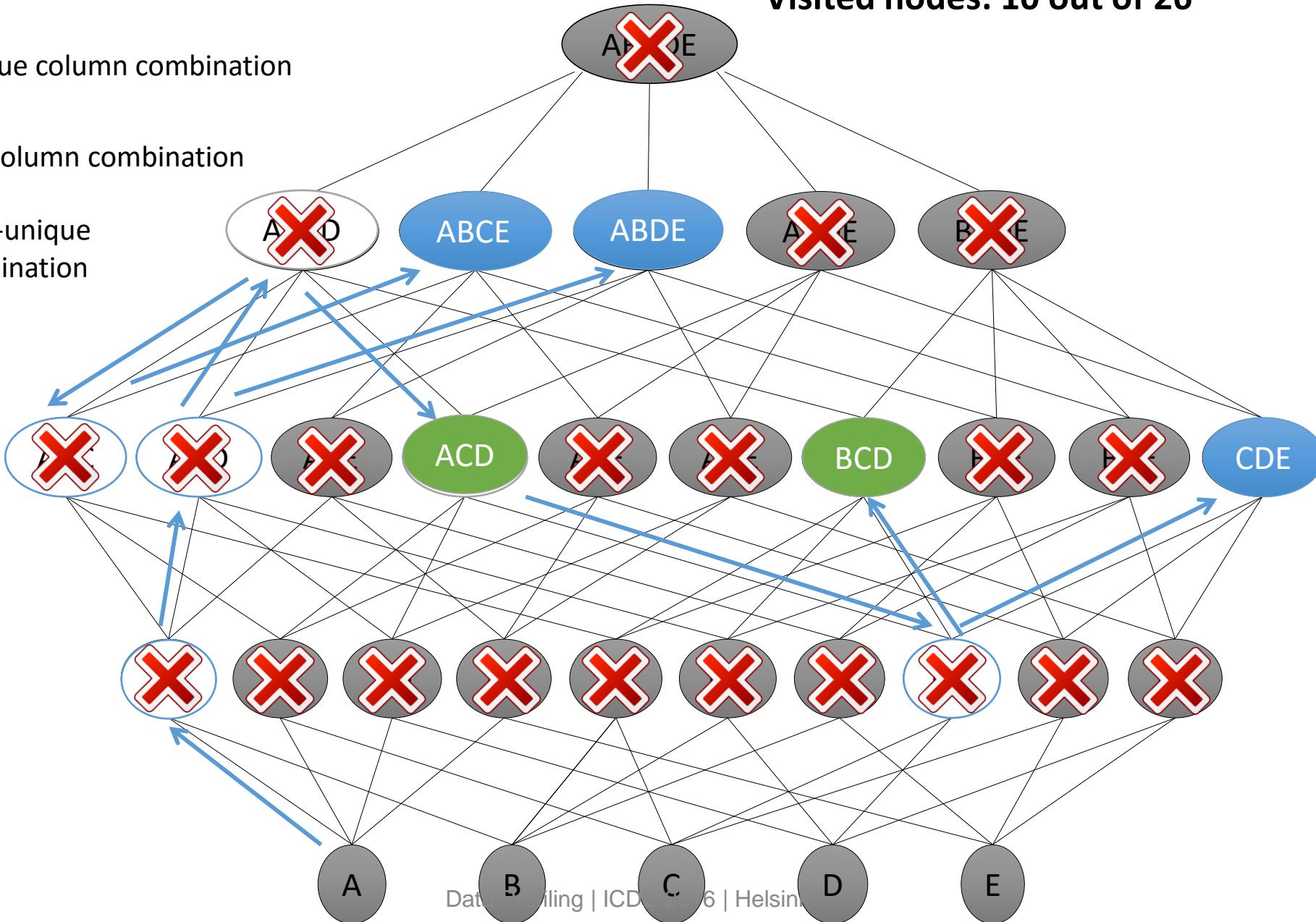
[Heise, Quiané-Ruiz, Abedjan, Jentzsch, Naumann: Scalable Discovery of Unique Column Combinations, PVLDB'14]

- Scalability is major design goal of DUCC
 - Random walk well suited for parallelization
 - Few coordination overhead
 - Threads/worker share findings through event bus
 - Uniques/non-uniques
 - Holes in graph
 - Lock-free to avoid bottlenecks
 - Only memory barrier in local event bus
- Basic idea: random walk through lattice
 - Pick random superset if current combination is non-unique
 - Pick random subset otherwise

- Unique column combination
- Minimal unique column combination
- Non-unique column combination
- Maximal non-unique column combination
- Pruned**

ACD and BCD are minimal uniques

Visited nodes: 10 out of 26



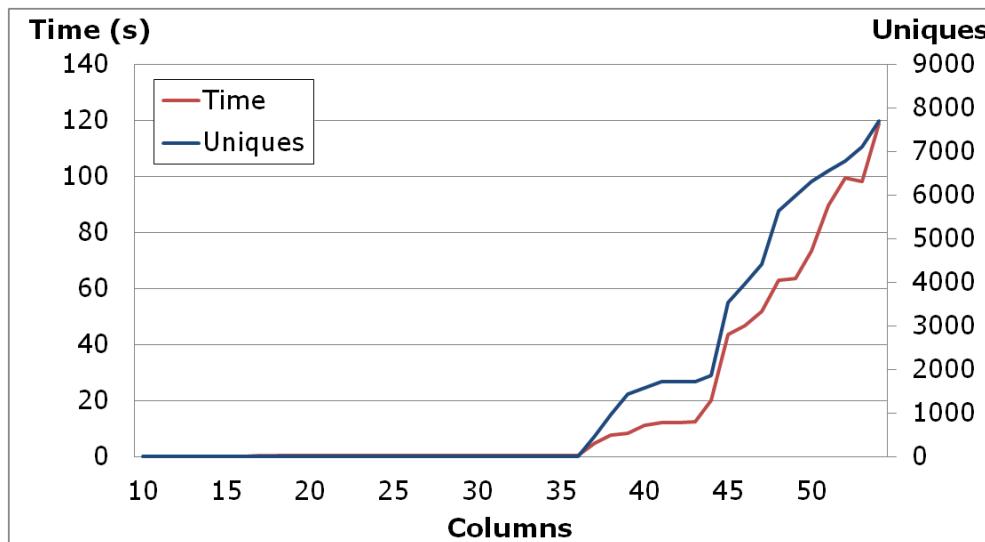
Position List Index

- Aka “partitions”
- Incorporates row-based pruning
- Intuition: number of duplicates decrease when going up the lattice
 - Many unnecessary rows are checked again and again
- Keep track of duplicates with inverted index
 - X: a->{r₁, r₂, r₃}, b->{r₄, r₅}
 - Y: 1->{r₁, r₃}, 2->{r₂, r₅}, 3->{r₄}
- Stripped partitions:
 - Remove clusters of size 1:
 - X: {{r₁, r₂, r₃}, {r₄, r₅}}
 - Y: {{r₁, r₃}, {r₂, r₅}}

X	Y
a	1
a	2
a	1
b	3
b	2

Analysis of DUCC

- Runtime mainly depends on size of solution set



- Worst case: Solution set is in the middle: $\binom{n}{n/2}$
- Aggressive pruning may lead to loss of minimal uniques!
 - Gordian's final step can be used to plug these holes

Gordian

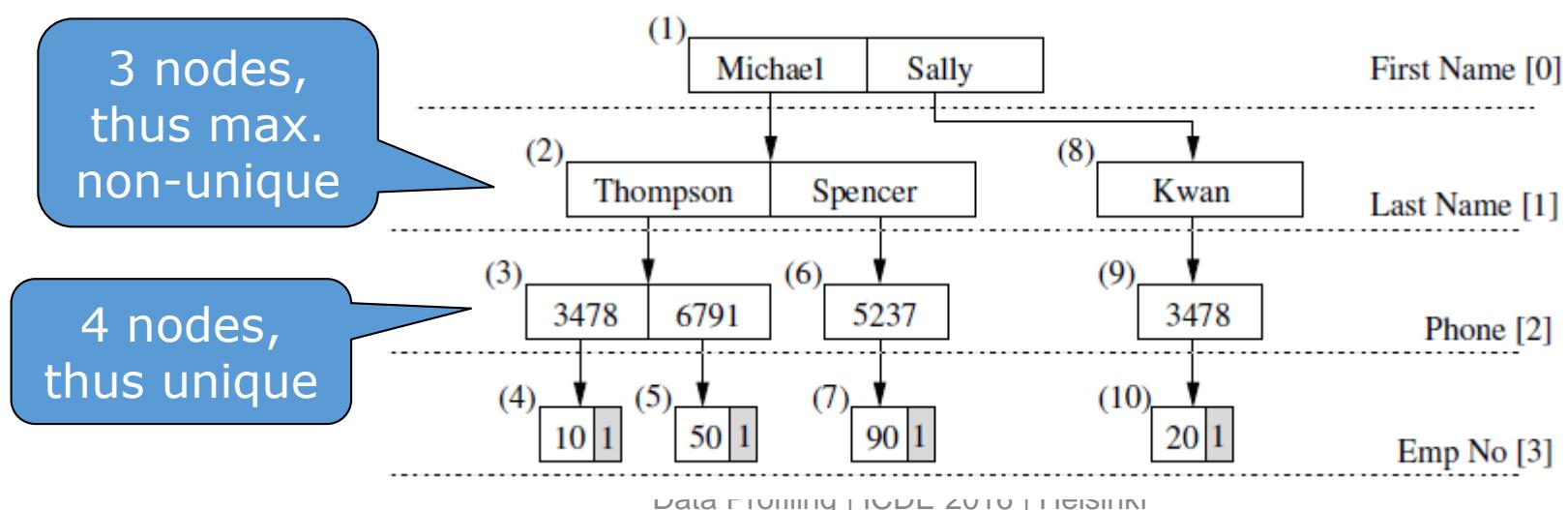
[Sismanis, Brown, Haas, Reinwald: GORDIAN: efficient and scalable discovery of composite keys, VLDB'06]

- Row-based algorithm
- Builds prefix tree while reading data
 - Discover maximal non-uniques on prefix tree
- Compute minimal uniques from maximal non-uniques
 - Complementation

Prefix tree

<i>FirstName</i>	<i>LastName</i>	<i>Phone</i>	<i>EmpNo</i>	<i>COUNT</i>
Michael	Thompson	3478	10	1
Sally	Kwan	3478	20	1
Michael	Spencer	5237	90	1
Michael	Thompson	6791	50	1

One tree
per
attribute
order



Analysis Gordian

- According to paper, polynomial in the number of tuples for data with a Zipfian distribution of values
 - Can abort scan as soon as duplicate has been found
- Worst case
 - Exponential in the number of columns
 - All data needs to be stored in memory
- Computing minimal uniques from maximal non-uniques
 - $O(\text{non-uniques}^3 \times \text{columns})$
 - Can be sped up with presorted list

Uniques on Dynamic Data: SWAN

[Abedjan, Quanie-Ruiz, Naumann: Detecting Unique Column Combinations on Dynamic Data, ICDE'14]

- **Inserts** may create new duplicate combinations
 - Minimal uniques might become non-unique
 - Maximal non-uniques might lose maximality
- **Deletes** remove duplicate value combinations
 - Non-uniques might get unique
 - Minimal uniques might lose minimality
- **SWAN**
 - Leverage the knowledge of previously discovered minimal uniques and maximal non-uniques
 - Create appropriate indices

Functional Dependencies



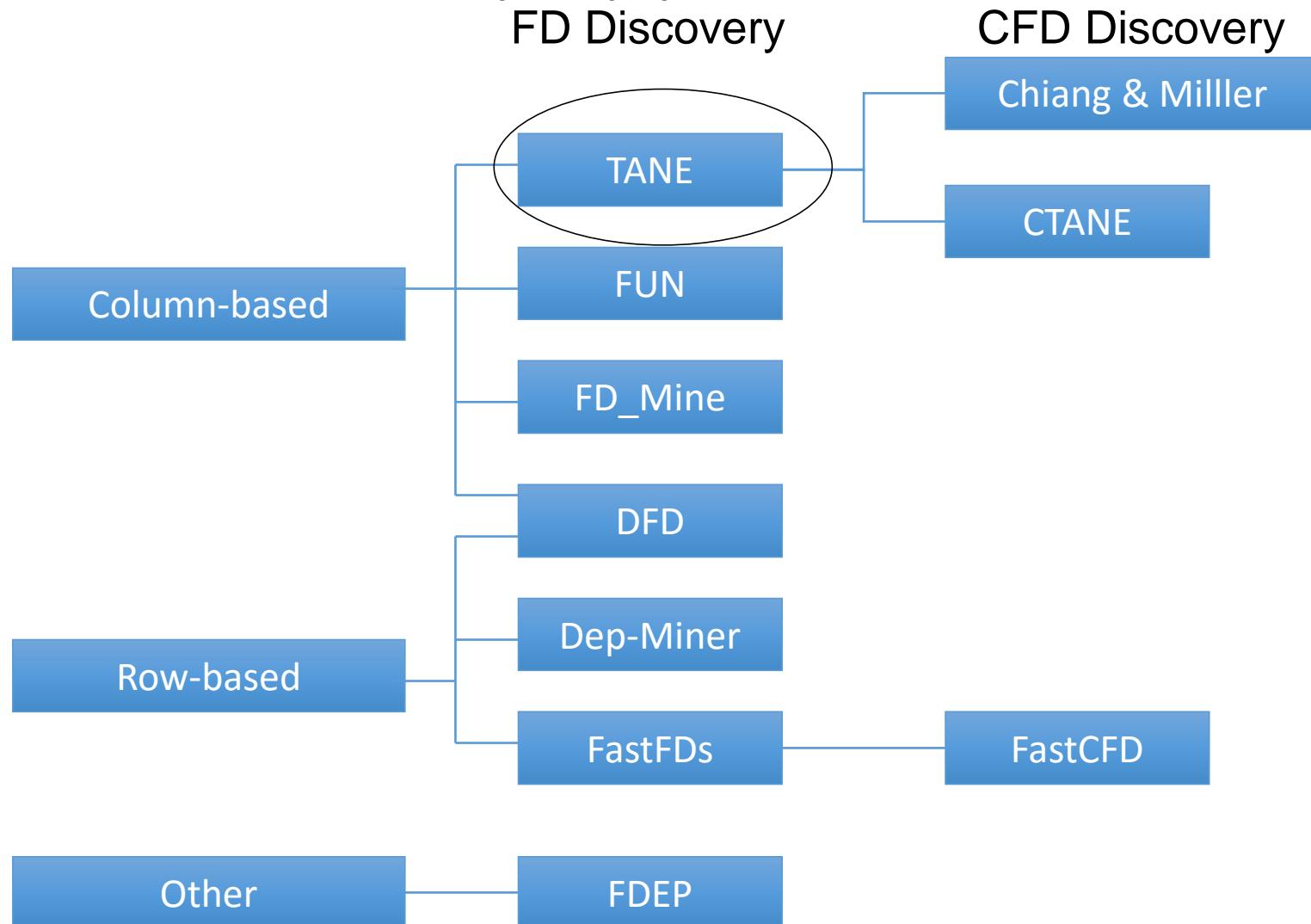
Trivial and minimal FDs

- „ $X \rightarrow A$ “ is a statement about a relation R: When two tuples have same value in attribute set X, they must have same values in attribute A.
- Non-trivial: At least one attribute on RHS does not appear on LHS
 - Street, City \rightarrow Zip, City
- Completely non-trivial: Attributes on LHS and RHS are disjoint.
 - Street, City \rightarrow Zip
- Minimal FD: RHS does not depend on any subset of LHS
- Typical goal: Given a relation R, find all minimal completely non-trivial functional dependencies.

Naive Discovery Approach

- Task: Given relation R , detect all minimal, non-trivial FDs $X \rightarrow A$.
- For each $A \in R$
 - For each column combination $X \setminus A$
 - For each pair of tuples (t_1, t_2)
 - If $t_1[X] = t_2[X]$ and $t_1[A] \neq t_2[A]$: Break
 - Return $X \rightarrow A$
- Complexity
 - For each of the $|R|$ possibilities for RHS
 - check $2^{|R|-1}$ combinations for LHS
 - And scan each record pair ($n^2/2$) for each check

Current FD Discovery approaches



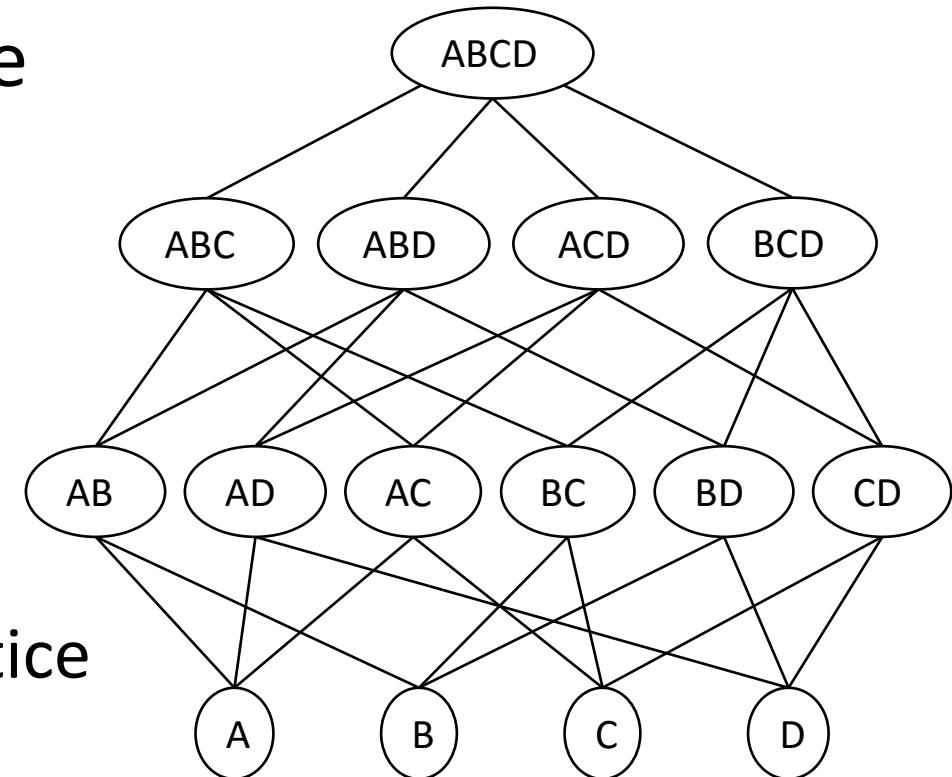
Tane – General Idea

[Huhtala, Kärkkäinen, Porkka, Toivonen:TANE: An Efficient Algorithm for Discovering Functional and Approximate Dependencies, Computer Journal'99]

- Two key ideas
 1. Reduce column combinations through pruning
 - Reasoning over FDs
 2. Reduce tuple sets through partitioning
 - Partition data according to attribute values
 - Level-wise increase of size of attribute set
 - Consider sets of tuples whose values agree on that set

TANE: Discovery strategy

- Bottom up traversal through lattice
 - \Rightarrow only minimal dependencies
 - Pruning
 - Re-use results from previous level
- For a set X , test all $X \setminus A \rightarrow A$, $A \in X$
 - \Rightarrow only non-trivial dependencies
 - Interpretation: Test each edge in lattice
 - Test on efficient data structure

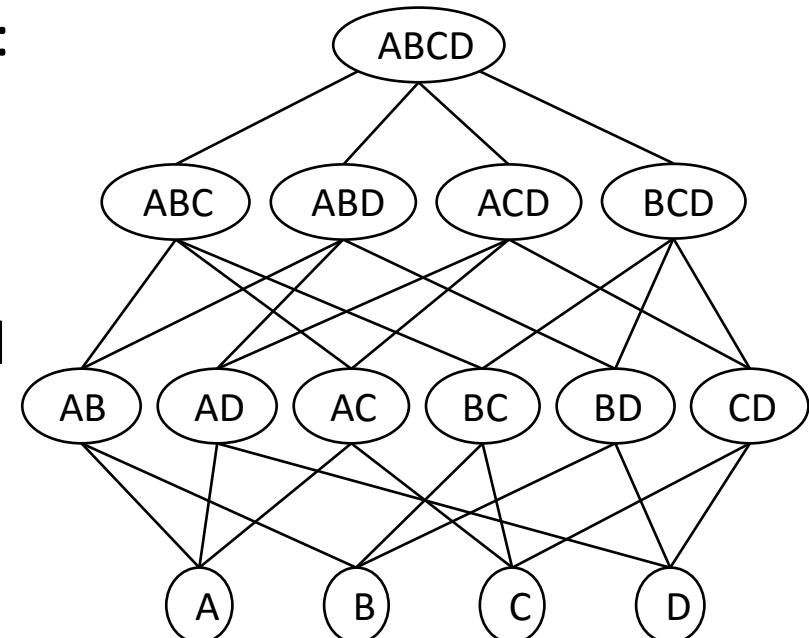


Candidate Sets

- RHS candidate set $C(X)$
- Stores only those attributes that might depend on **all** other attributes in X .
 - I.e., those that still need to be checked
 - If $A \in C(X)$ then A does not depend on any proper subset of X .
- $C(X) = R \setminus \{A \in X \mid X \setminus A \rightarrow A \text{ holds}\}$
- Examples: $R = \{ABCD\}$, and $A \rightarrow C$ and $CD \rightarrow B$ hold
 - $C(A) = \{ABCD\} \setminus \{\} = C(B) = C(C) = C(D)$
 - $C(AB) = \{ABCD\} \setminus \{\}$
 - $C(AC) = \{ABCD\} \setminus \{C\} = \{ABD\}$
 - $C(CD) = \{ABCD\} \setminus \{\}$
 - $C(BCD) = \{ABCD\} \setminus \{B\} = \{ACD\}$

RHS Candidate Pruning

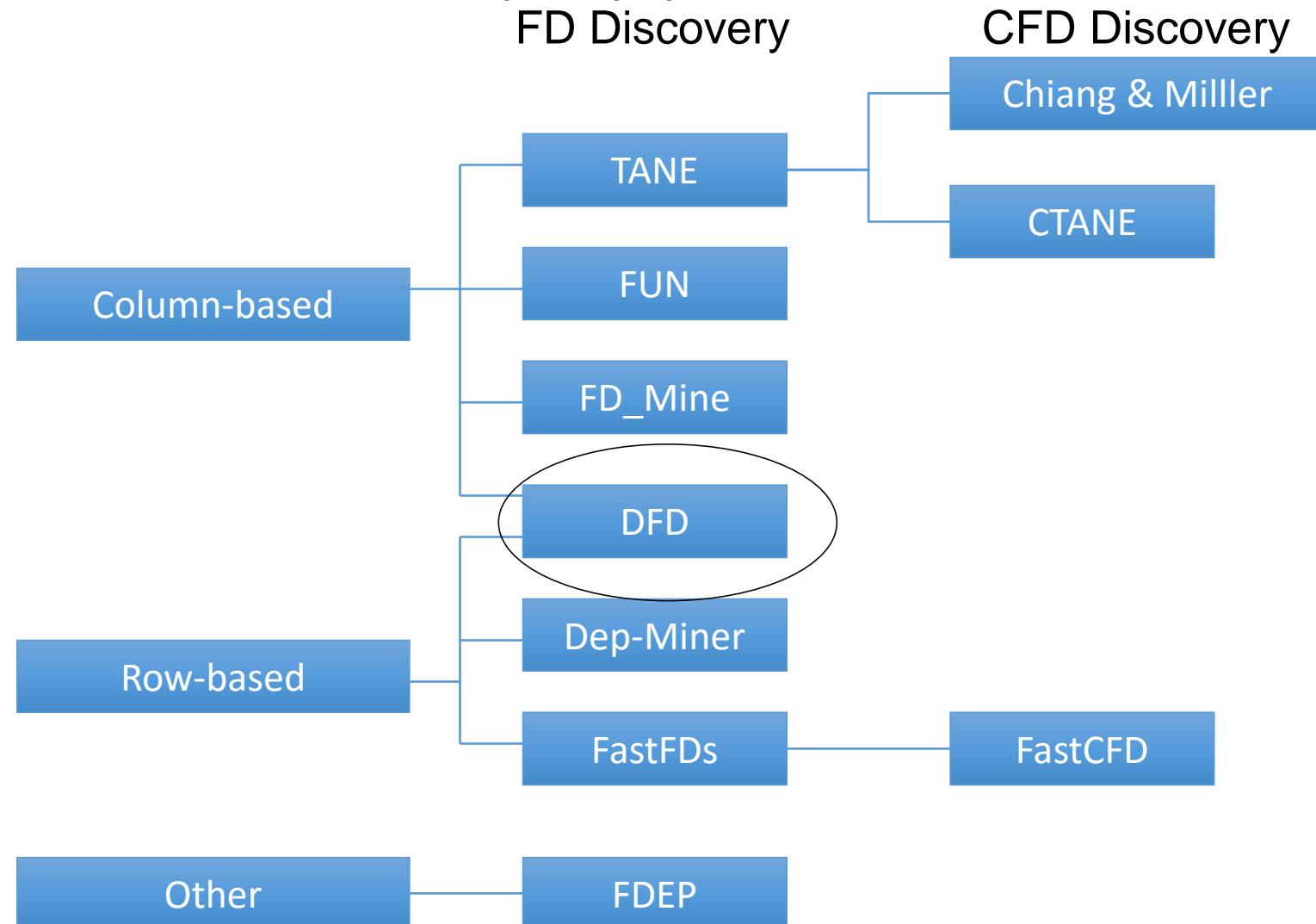
- RHS candidates: $C^+(X) = \{A \in R \mid \forall B \in X: X \setminus \{A, B\} \rightarrow B \text{ does not hold}\}$
 - Special case: $A = B$ corresponds to $C(X)$
 - Reminder: $C(X) = R \setminus \{A \in X \mid X \setminus A \rightarrow A \text{ holds}\}$
- This definition removes three types of candidates:
 - Minimality
 - Pseudotransitivity
 - Superkey
- Examples: $R = \{ABCD\}$, and $A \rightarrow C$ and $CD \rightarrow B$ hold
 - $C(ABC) = \{A\}$
 - $C(BCD) = \{ACD\}$



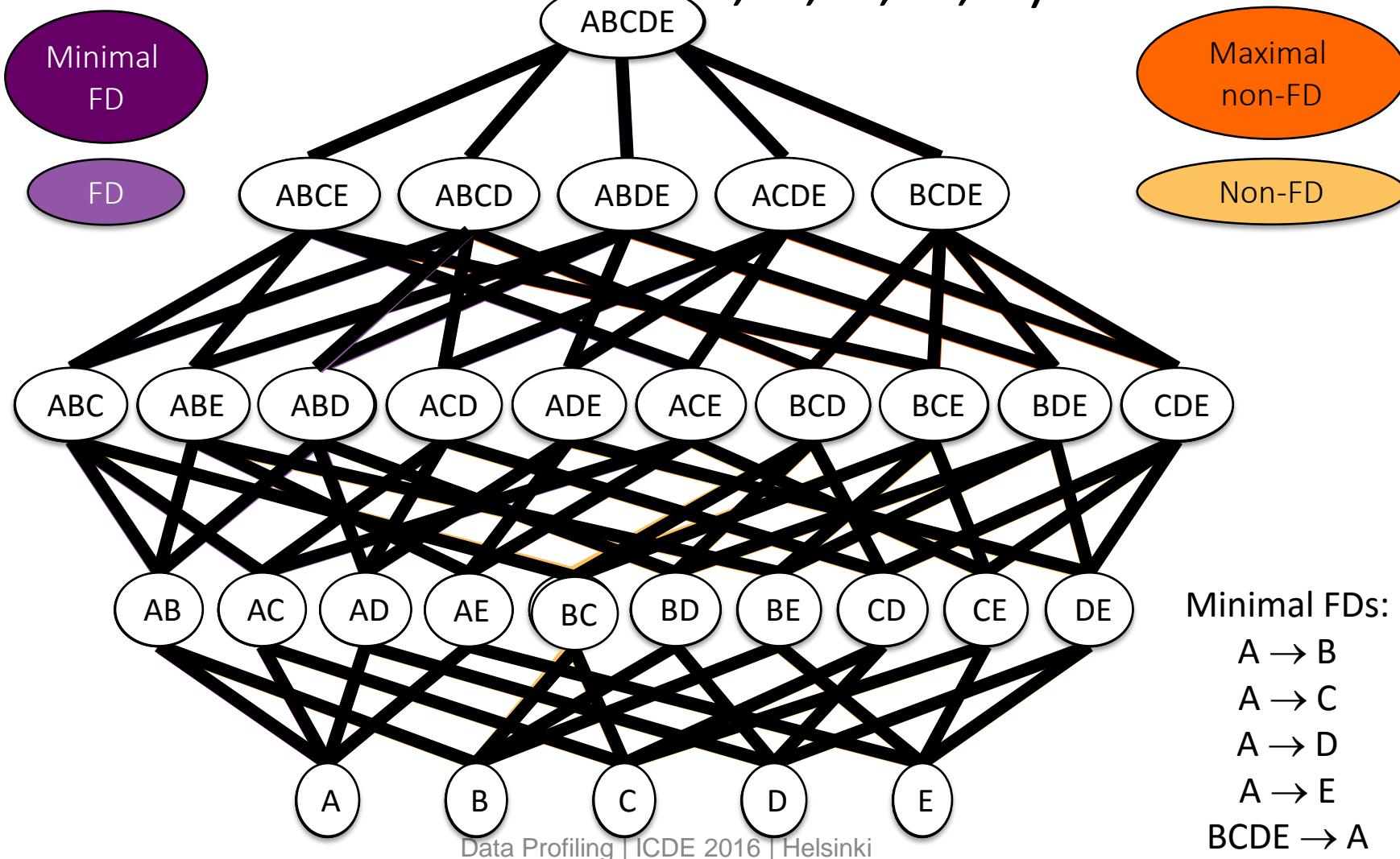
Partial FDs with TANE

- Definition based on minimum number of tuples to be removed from R for $X \rightarrow A$ to hold in R.
 - Discovery problem:
 - Given relation R and threshold ε , find all minimal non-trivial FDs $X \rightarrow A$ such that $e(X \rightarrow A) \leq \varepsilon$
 - Called “approximate” FDs in paper
1. Define error: Fraction of tuples causing FD violation
 - Error $e(X \rightarrow A) = \min\{|S| \mid S \subseteq R, R \setminus S \models X \rightarrow A\} / |R|$
 2. Specify error threshold ε
 3. Modify dependency checking algorithm
 - Efficient algorithm to compute error
 - Bounds to avoid error calculation

Current FD Discovery approaches



DFD Explanation: Tane visualized for $R = (A, B, C, D, E)$

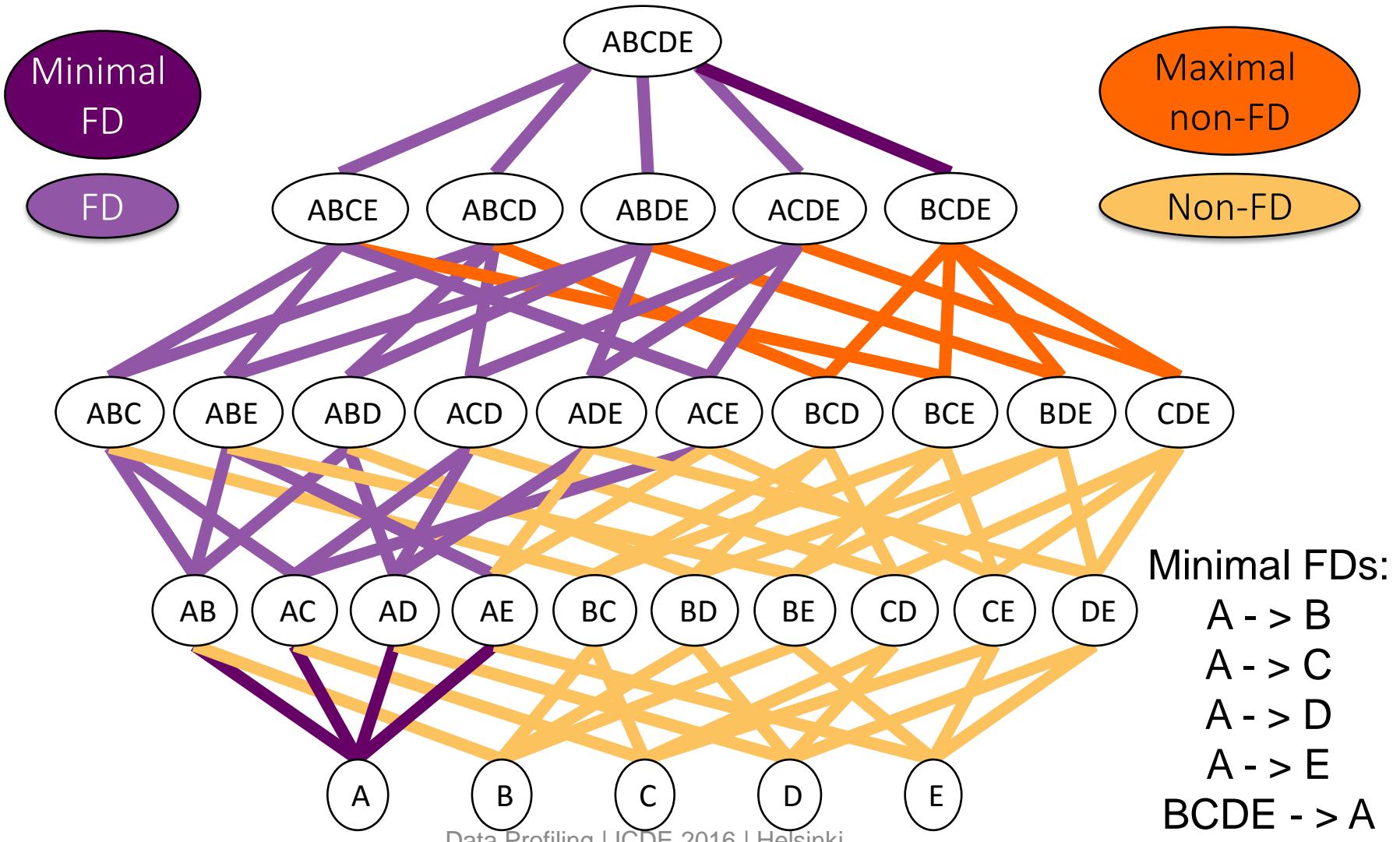


DFD: Depth-first approach for functional dependency discovery

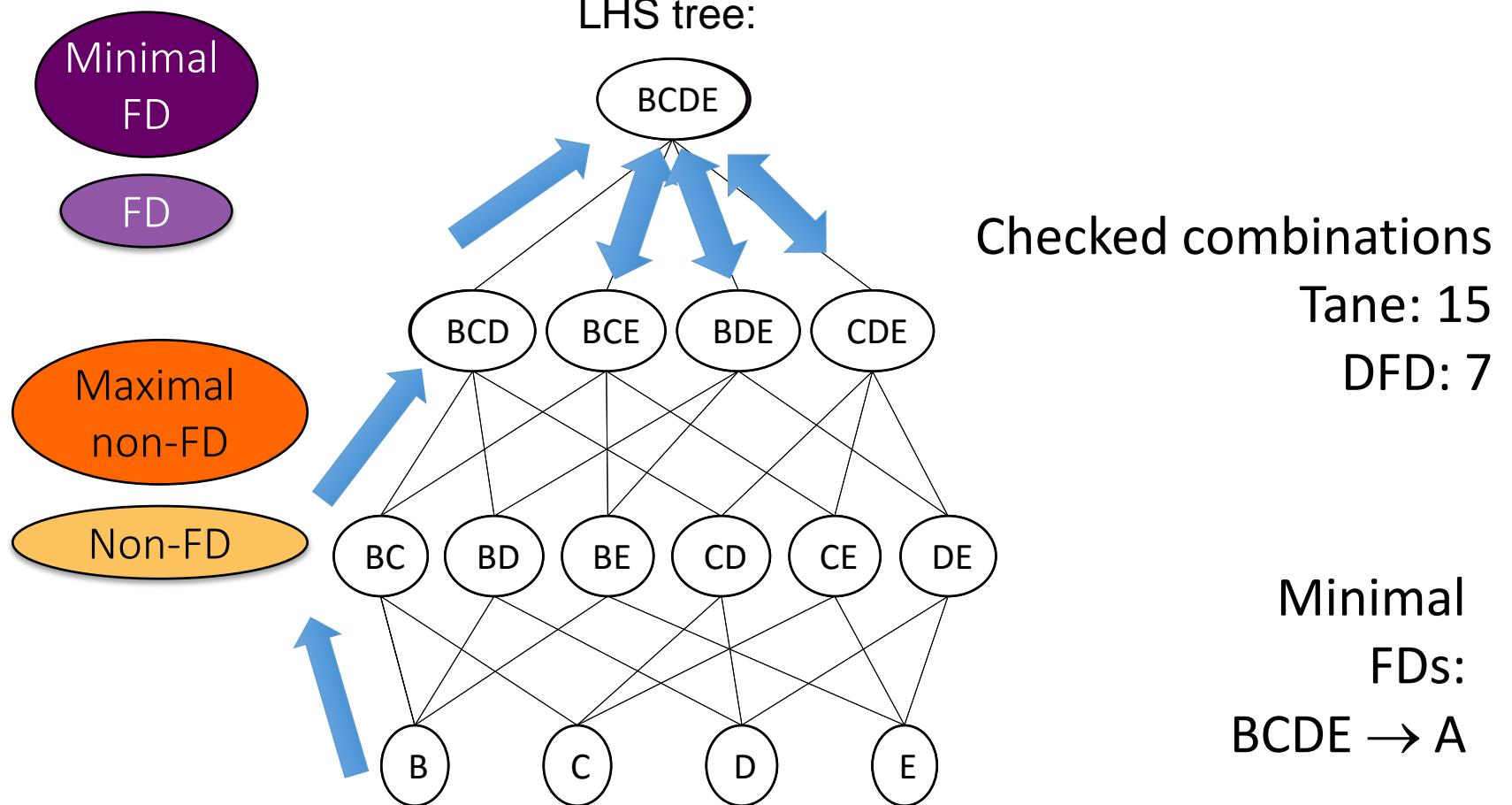
[Abedjan,Schulze,Naumann: DFD:Efficient Functional Dependency Discovery, CIKM'14]

- Traverse depth-first and prune upwards and downwards
- Applied for key/unique discovery: DUCC
 - Key discovery is a subproblem of FD discovery
 - Adapt the concept of minimality in keys to LHS of FDs:
 - An FD $X \rightarrow C$ is minimal if $" X' \subset X : X' \xrightarrow{NOT} C "$
 - A non-dependency $X \xrightarrow{NOT} C$ is maximal if $" X' \supset X : X' \rightarrow C "$

Decompose Relation for each RHS



Decomposition for RHS=A



Traversal Holes

- Aggressive traversal and pruning
 - As for DUCC: Some nodes might never be reached.
- GORDIAN [VLDB'06]:
 - Complement the set of **maximal non-keys**
 - = set of **minimal keys**
- Key observation from DUCC: the **difference** of one set and the complement of its counterpart delivers the **unvisited nodes!**
- Hole discovery works for FDs too:
 - Consider **minimal FD LHS** and **maximal non-FD LHS**

Execution time - uniprot

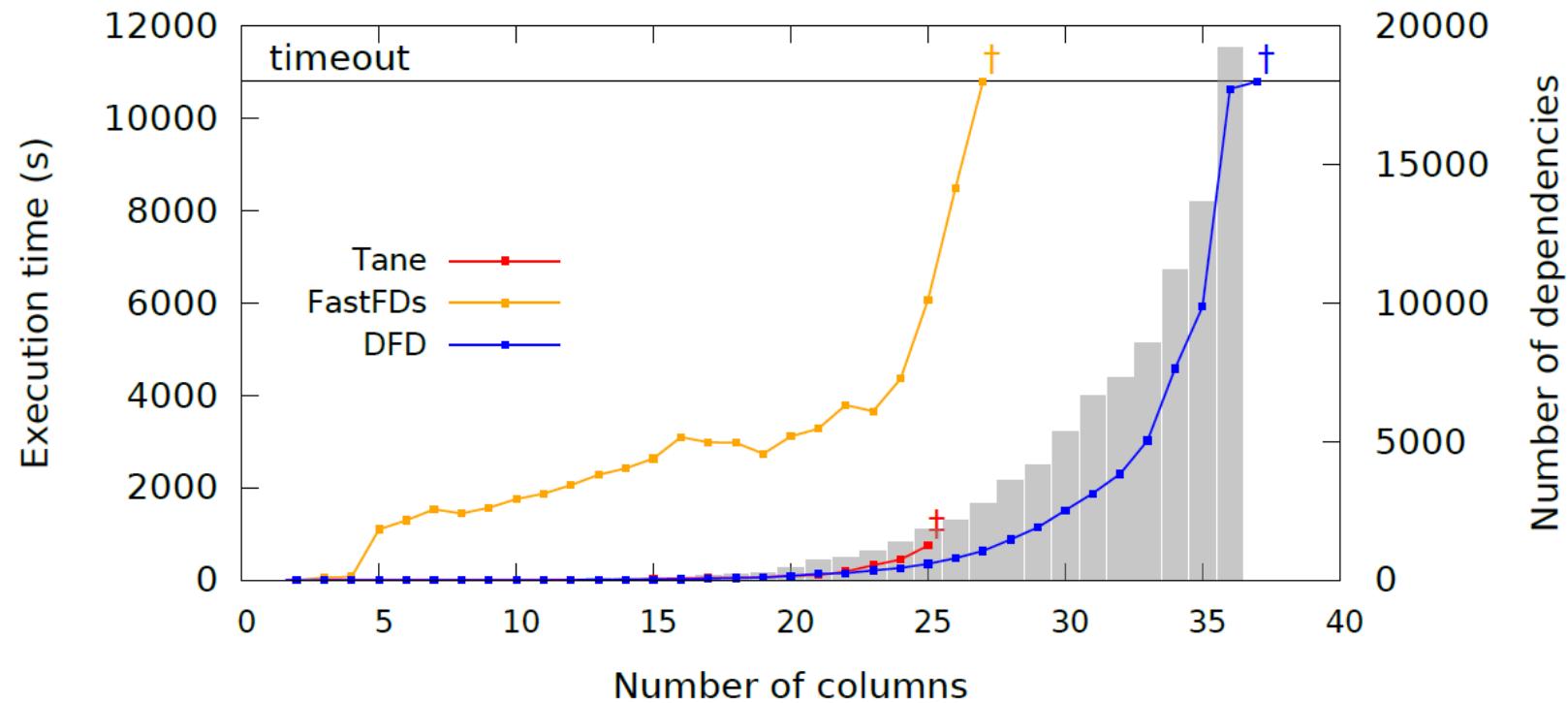


Figure 4.2: Execution time for Tane, FastFDs, and DFD on the first 100,000 rows of the uniprot dataset. († - Time Limit ‡ - Memory Limit)

Functional Dependency Evaluation

dataSet	Columns	Rows	FDs	Tane	FUN	FD_Mine	Dep-Miner	FastFDs	FDep	DFD
iris	5	150	4	0.6s	0.1s	0.1s	0.1s	0.1s	0.1s	0.1s
balance-scale	5	625	1	0.9s	0.4s	0.3s	0.2s	0.5s	0.3s	0.2s
chess	7	28,056	1	2.0s	1.0s	3.0s	200.8s	200.1s	202.5s	0.9s
abalone	9	4,177	137	1.0s	0.3s	1.0s	2.9s	3.0s	4.1s	0.9s
nursery	9	12,960	1	3.1s	1.5s	6.0s	132.0s	131.9s	56.6s	1.1s
breast-cancer	11	699	46	1.4s	0.4s	1.5s	0.9s	1.0s	0.4s	0.9s
bridges	13	108	142	1.3s	0.5s	2.9s	0.2s	0.2s	0.2s	0.9s
echocardiogram	13	132	538	0.8s	0.1s	69.9s	0.1s	0.1s	0.1s	1.6s
adult	14	48,842	78	81.2s	150.2s	485.3s	5982s	5946s	760.7s	6.8s
letter	17	20,000	61	326s	553.9s	ML	865.4s	853.9s	292.3s	9.1s
hepatitis	20	155	8,250	10.9s	321.6s	TL	5363.1s	9.3s	0.5s	317.8s
horse	27	368	128,726	5451.s	TL	TL	TL	386.8s	15.7s	TL
fd-reduced-30	30	250,000	89,571	41.1s	78.4s	TL	391.9s	391.3s	TL	TL
flight	109	1,000	982,631	ML	TL	ML	TL	TL	213.5s	TL
plista	125	1,000	178,152	ML	TL	TL	TL	TL	26.4s	TL

IND Discovery

1. DeMarchi's Algorithm

2. Spider

3. BINDER & MIND

- High performance IND detection
- Work by Thorsten Papenbrock



BINDER – divide & conquer based IND detection
 Linking web tables – an example

Name	Type	Equatorial diameter	Mass	Orbital radius	Orbital period	Rotation period	Confirmed moons	Rings	Atmosphere
Mercury	Terrestrial	0.382	0.06	0.47	0.24	58.64	0	no	minimal
Venus	Terrestrial	0.949	0.82	0.72	0.62	-243.02	0	no	CO ₂ , N ₂
Earth	Terrestrial	1.000	1.00	1.00	1.00	1.00	1	no	N ₂ , O ₂ , Ar
Mars	Terrestrial	0.532	0.11	1.52	1.88	1.03	2	no	CO ₂ , N ₂ , Ar
Jupiter	Giant	11.209	317.8	5.20	11.86	0.41	67	yes	H ₂ , He
Saturn	Giant	9.449	95.2	9.54	29.46	0.43	62	yes	H ₂ , He
Uranus	Giant	4.007	14.6	19.22	84.01	-0.72	27	yes	
Neptune	Giant	3.883	17.2	30.06	164.8	0.67	14	yes	
Mars		780		25.6			72		
Jupiter		399		13.1			121		
Saturn		378		12.4			138		
Uranus		370		12.15			151		
Neptune		367		12.07			158		

Sign	House	Domicile	Detriment	Exaltation	Fall	Planetary Joy
Aries	1st House	Mars	Venus	Sun	Saturn	Mercury
Taurus	2nd House	Venus	Pluto	Moon	Uranus	Jupiter
Gemini	3rd House	Mercury	Jupiter	N/A	N/A	Saturn
Cancer	4th House	Moon	Saturn	Jupiter	Mars	Venus
Leo	5th House	Sun	Uranus	Neptune	Mercury	Mars
Virgo	6th House	Mercury	Neptune	Pluto, Mercury	Venus	Saturn
Libra	7th House	Venus	Mars	Saturn	Sun	Moon
Scorpio	8th House	Pluto	Venus	Uranus	Moon	Saturn
Sagittarius	9th House	Jupiter	Mercury	N/A	N/A	Sun
	10th House	Saturn	Moon	Mars	Jupiter	Mercury
Capricorn	11th House	Uranus	Sun	Mercury	Neptune	Venus

Planet	Rotation Period	Revolution Period	Symbol	Unicode	Glyph
Mercury	58.6 days	87.97 days	Sun	U+2609	○
Venus	243 days	224.7 days	Moon	U+263D	☽
Earth	0.99 days	365.26 days	Moon	U+263E	☾
Mars	1.03 days	1.88 years	Mercury	U+263F	☿
Jupiter	0.41 days	11.86 years	Venus	U+2640	♀
Saturn	0.45 days	29.46 years	Earth	U+1F728	♁
Uranus	0.72 days	84.01 years	Mars	U+2642	♂
Neptune	0.67 days	164.79 years	Jupiter	U+2643	♃
Pluto	6.39 days	248.59 years	Saturn	U+2644	♄
Mercury	57.91	1	U+2645	♁	
Venus	108.21	1.86859	U+26E2	♂	
Earth	149.6	1.3825	Neptune	U+2646	♃
Mars	227.92	1.52353	Eris	≈ U+2641	♂
Ceres	413.79	1.81552	Eris	≈ U+29EC	♀
Jupiter	778.57	1.88154	Pluto	U+2647	♅
Saturn	1,433.53	1.84123	Pluto	not present	--
Uranus	2,872.46	2.00377	Aries	U+2648	♈
Neptune	4,495.06	1.56488	Taurus	U+2649	♉
Pluto	5,869.66	1.3058	Gemini	U+264A	♊
Mercury	0.4	0.387	Cancer	U+264B	♋
Venus	0.7	0.723	Leo	U+264C	♌
Earth	1	1	Virgo	U+264D	♍
Mars	1.6	1.524	Libra	U+264E	♎
Asteroid belt	2.8	2.767	Scorpio	U+264F	♏
Jupiter	5.2	5.203	Sagittarius	U+2650	♐
Saturn	10	9.539	Capricorn	U+2651	♑
Uranus	19.6	19.191	Capricorn	U+2651	♑
Neptune	38.8	30.061	Aquarius	U+2652	♒
Pluto	77.2	39.529	Pisces	U+2653	♓
	Conjunction	U+260C	☌

Unary IND detection complexity

Name	Type	Equatorial diameter	Mass	Orbital radius	Orbital period	Rotation period	Confirmed moons	Rings	Atmosphere
Mercury	Terrestrial	0.382	0.06	0.47	0.24	58.64	0	no	minimal
Venus	Terrestrial	0.949	0.82	0.72	0.62	-243.02	0	no	CO ₂ , N ₂
Earth	Terrestrial	1.000	1.00	1.00	1.00	1.00	1	no	N ₂ , O ₂ , Ar
Mars	Terrestrial	0.532	0.11	1.52	1.88	1.03	2	no	CO ₂ , N ₂ , Ar
Jupiter	Giant	11.209	317.8	5.20	11.86	0.41	67	yes	H ₂ , He
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Uranus	Giant	4.007	14.6	19.22	84.01	-0.72	27	yes	H ₂ , He
Neptune	Giant	3.883	17.2	30.06	164.8	0.67	14	yes	H ₂ , He

- Name ⊑ Type ?
- Name ⊑ Equatorial_diameter ?
- Name ⊑ Mass ?
- Name ⊑ Orbital_radius ?
- Name ⊑ Orbital_period ?
- Name ⊑ Rotation_period ?
- Name ⊑ Confirmed_moons ?
- Name ⊑ Rings ?
- Name ⊑ Atmosphere ?
- Type ⊑ Name ?
- Type ⊑ Equatorial_diameter ?
- Type ⊑ Mass ?
- Type ⊑ Orbital_radius ?
- Type ⊑ Orbital_period ?
- Type ⊑ Rotation_period ?
- Type ⊑ Confirmed_moons ?
- Type ⊑ Rings ?
- Type ⊑ Atmosphere ?

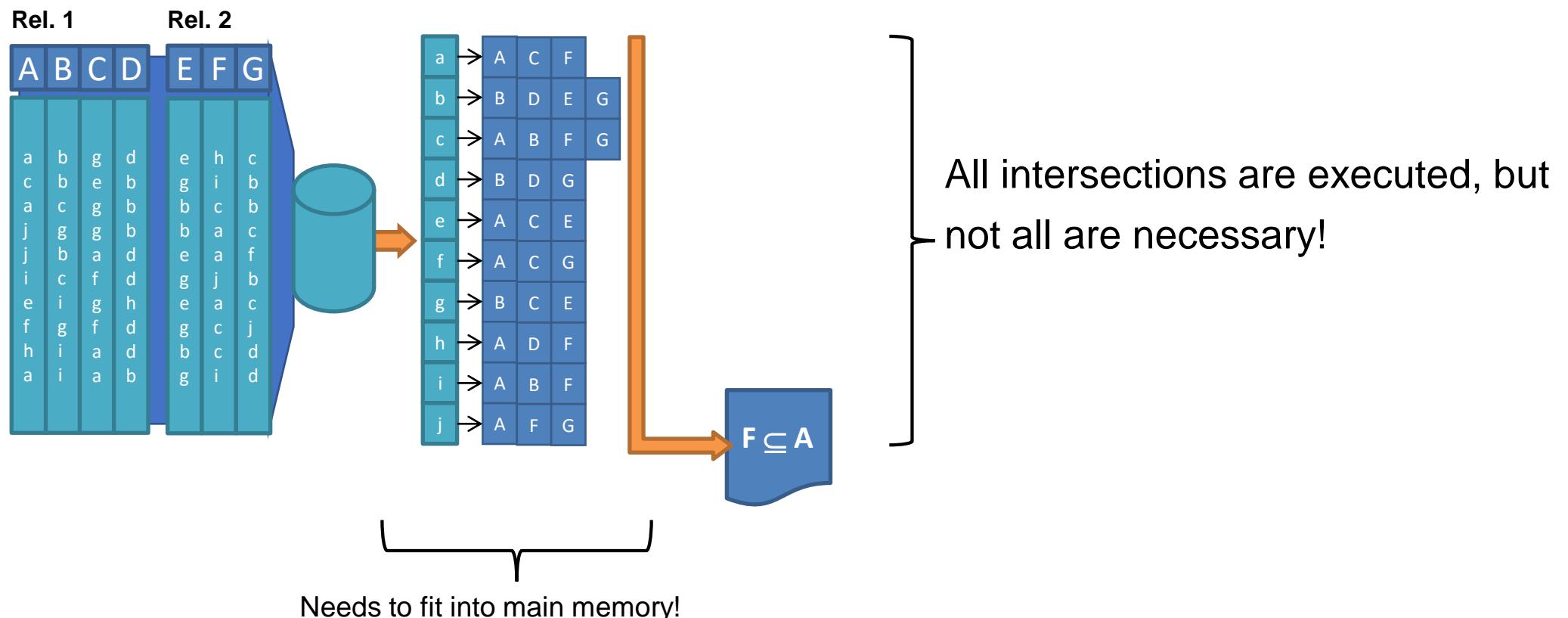
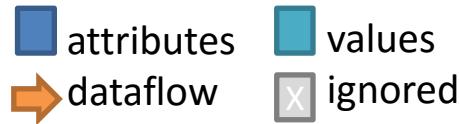
Complexity: $O(n^2 \cdot n)$
for n attributes

Example:
10 attr ~ 90 checks
1,000 attr ~ 999,000 checks

- Mass ⊑ Name ?
- Mass ⊑ Type ?
- Mass ⊑ Equatorial_diameter ?
- ...

MIND

[Marchi, Lopes, Petit: Unary and n-ary inclusion dependency discovery in relational databases, JIIS'09]



BINDER algorithm – workflow

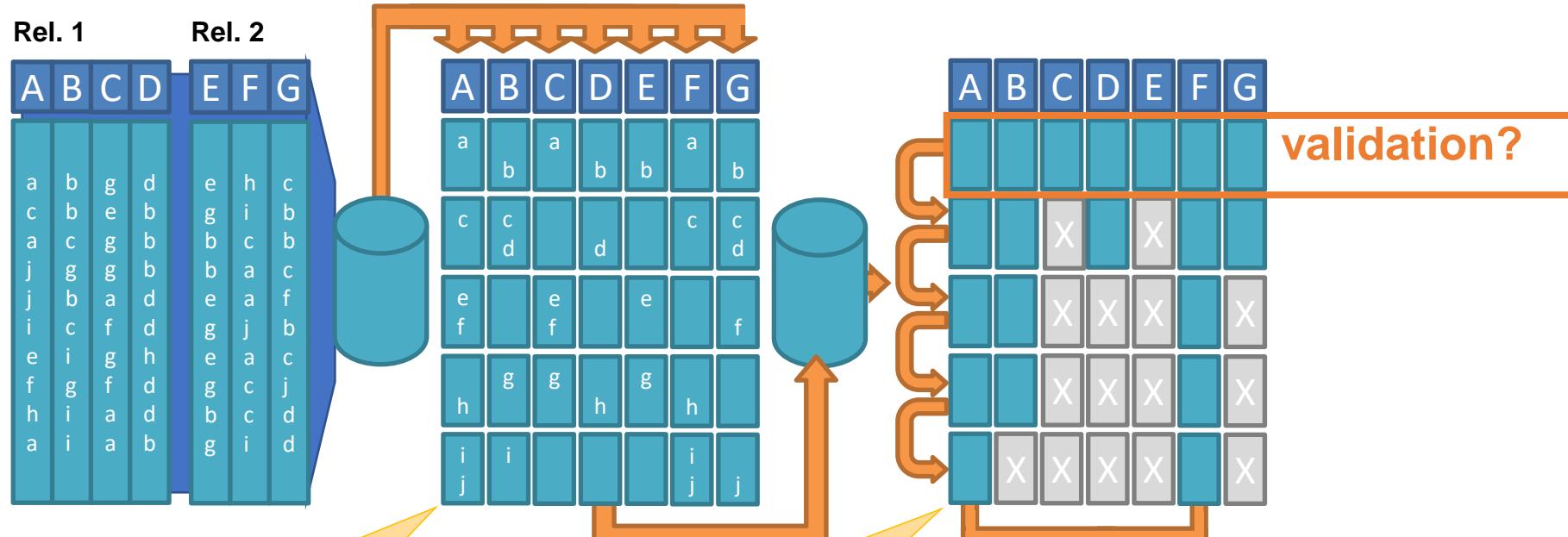
[Papenbrock, Quiane, Naumann: Divide & Conquer-based Inclusion Dependency Discovery, PVL]

■ attributes ■ values
➡ dataflow ✕ ignored

Divide

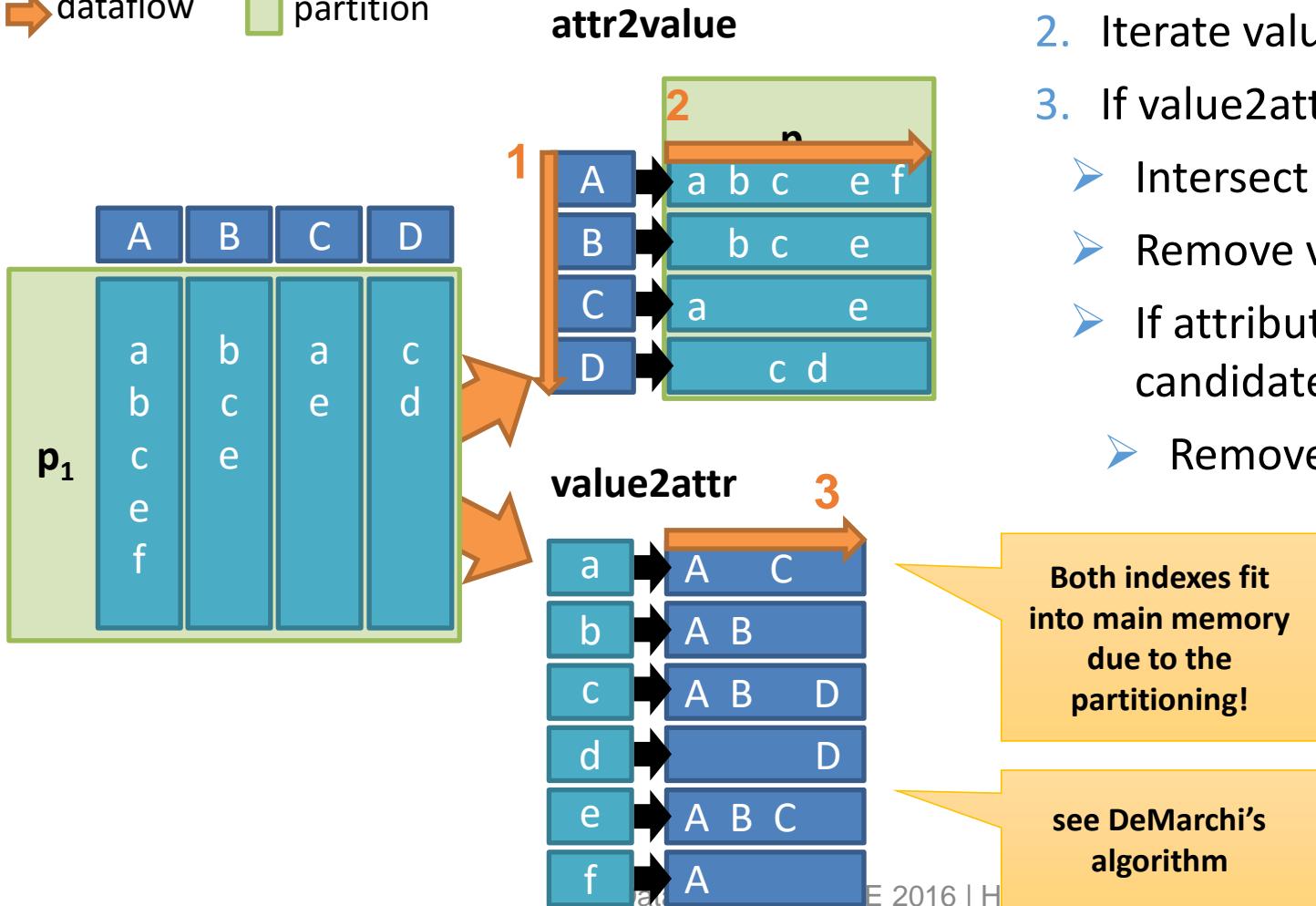
No sortation
needed, just
hashing

Conquer



BINDER algorithm – validation

■ attributes ■ values
➡ dataflow ■ partition



1. Iterate attributes
2. Iterate values
3. If $value2attr$ entry exists
 - Intersect candidates with this list
 - Remove $value2attr$ entry
 - If attribute removed from all candidates
 - Remove entry from $attr2value$

Both indexes fit
into main memory
due to the
partitioning!

see DeMarchi's
algorithm

BINDER algorithm – validation example

attr2value

a b c e f
b c e
a e
c d

Never tested! →

value2attr

A C
A B
A B D
D
A B C
A

	A	B	C	D
look up	B,C,D	A,C,D	A,B,D	A,B,C

1. Iterate attributes

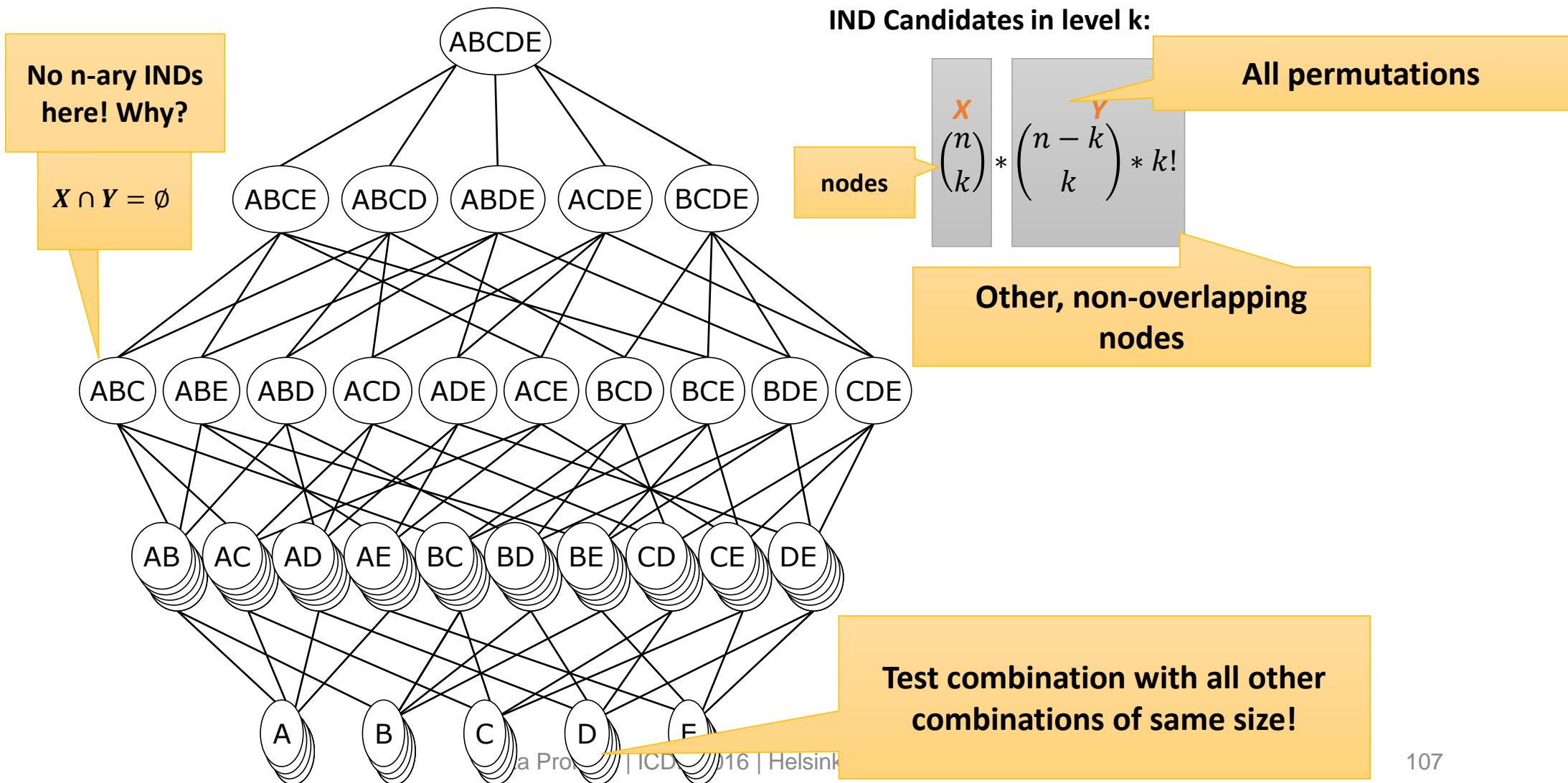
2. Iterate values

3. If value2attr entry exists

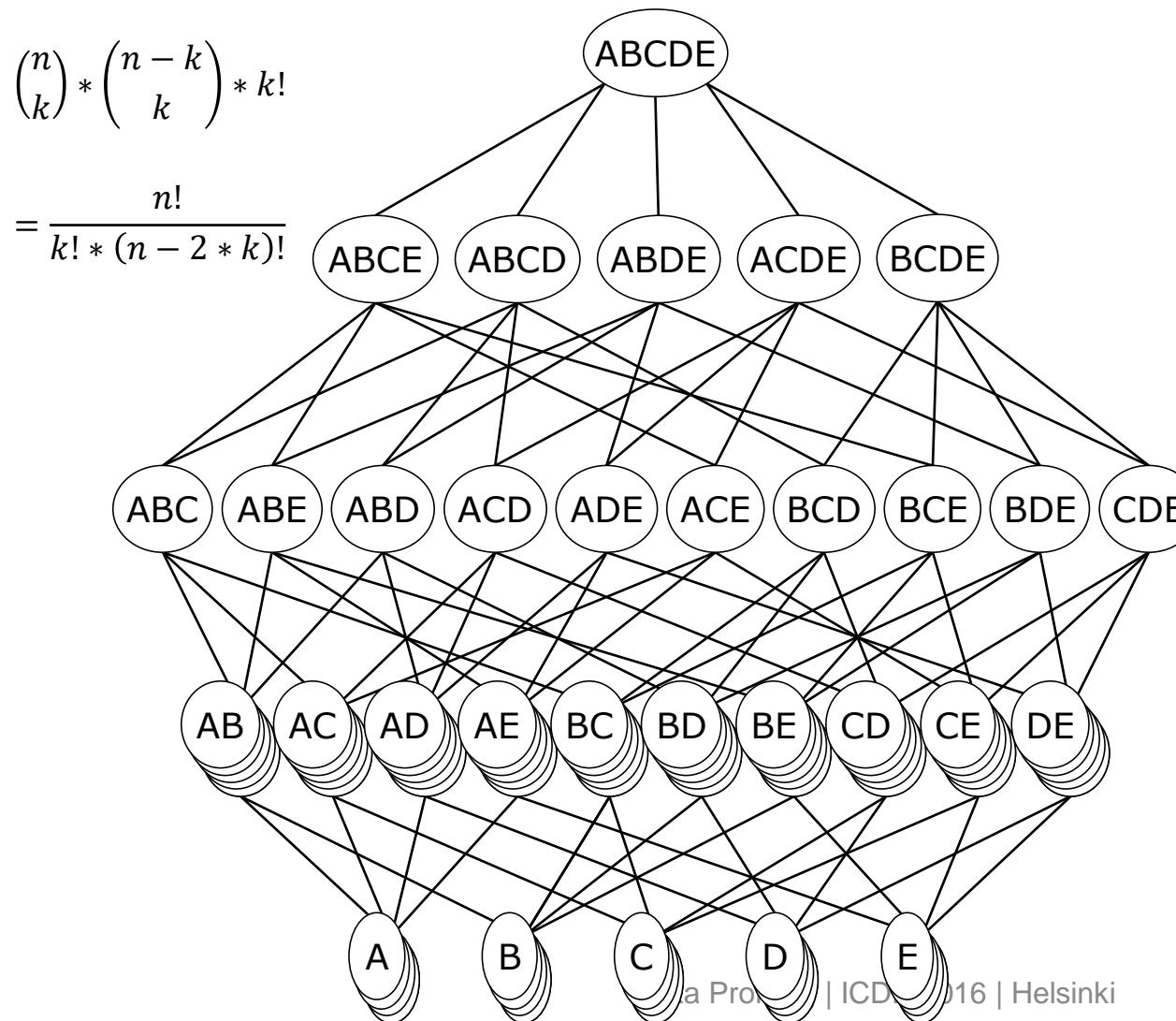
- Intersect candidates with this list
- Remove value2attr entry
- If attribute removed from all candidates
- Remove entry from attr2value

$B \subseteq A$
 $C \subseteq A$

N-ary IND detection complexity



N-ary IND detection complexity



$$\binom{5}{5} * \binom{5-5}{5} * 5! \sim 0$$

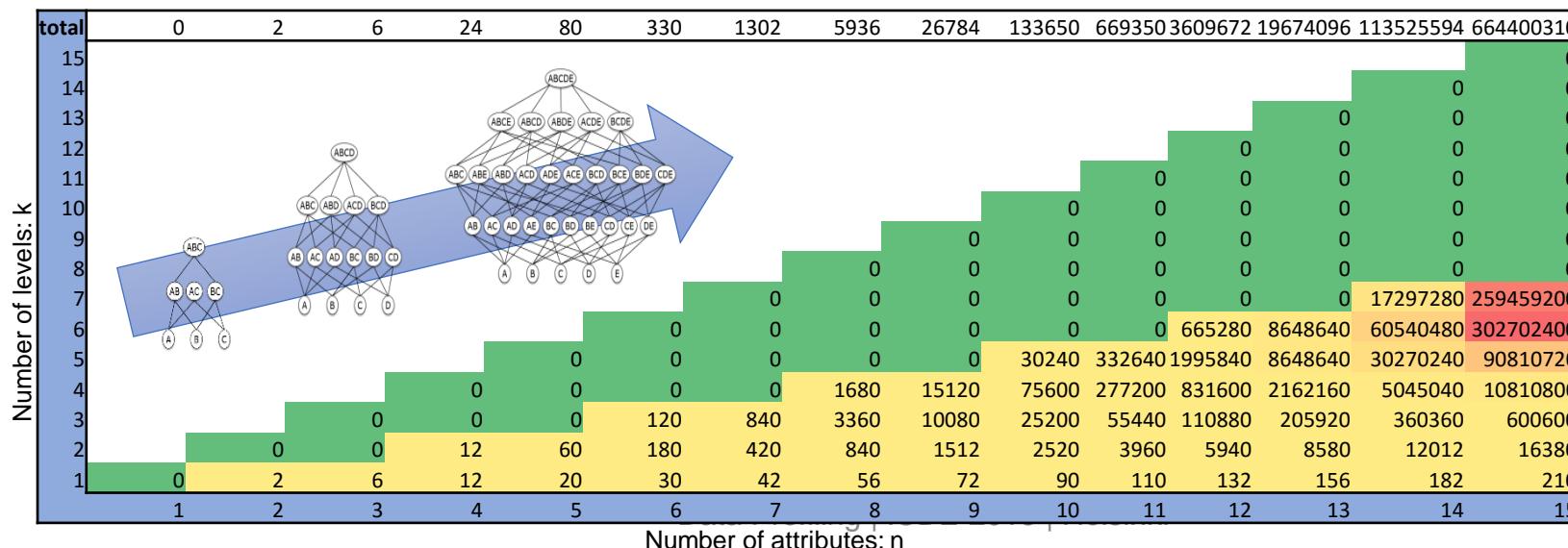
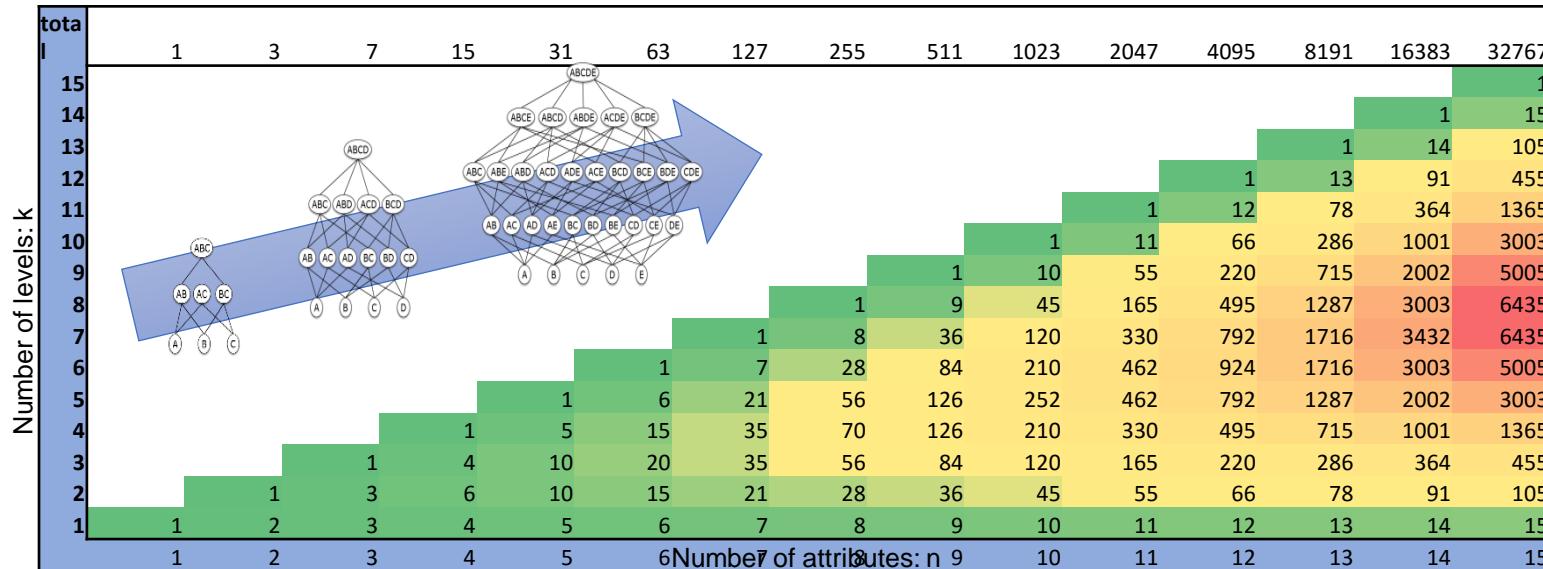
$$\binom{5}{4} * \binom{5-4}{4} * 4! \sim 0$$

$$\binom{5}{3} * \binom{5-3}{3} * 3! \sim 0$$

$$\binom{5}{2} * \binom{5-2}{2} * 2! = 60$$

$$\binom{5}{1} * \binom{5-1}{1} * 1! = 20 = n^2 - n$$

N-ary IND detection complexity



MIND & BINDER – candidate generation

- **Apriori algorithm:**
 - Bottom-up lattice traversal strategy
 - Input: all valid attribute combinations of size n
 - Output: all candidate attribute combinations of size n+1
- **Adaption for n-ary IND detection:**
 - Let R_i be the i-th relation in the relational schemata R. For each valid IND $R_j[X] \subseteq R_k[Y]$ with $|X|=|Y|=n$ generate all IND candidates $R_j[XA] \subseteq R_k[YA]$ so that:
 1. $R_j[X] \subseteq R_k[Y]$ and $R_j[A] \subseteq R_k[B]$ (both are valid INDs)
 2. $\forall X_i \in X: X_i < A$ (INDs are permutable; do not generate them twice)
 3. $A \notin X, B \notin Y$ (do not generate trivial candidates)

Intrinsic limitations of IND algorithms

- Observations: all IND algorithms follow a common pattern

Algorithm	Phase 1 Data Reorganization	Phase 2 Comparison
De Marchi	Create Inverted Index	Intersect Attribute Groups
SPIDER	Sort Columns	Value-based Iteration
BINDER	Partition Columns	In-Memory Partition Comparison

- e.g., $\text{IND } A \subseteq B$
 - to prove, need to read A completely
 - to disprove, need to read B completely
- Data reorganization is the most expensive phase
 - I/O-heavy workload, but other phase brings considerable I/O as well

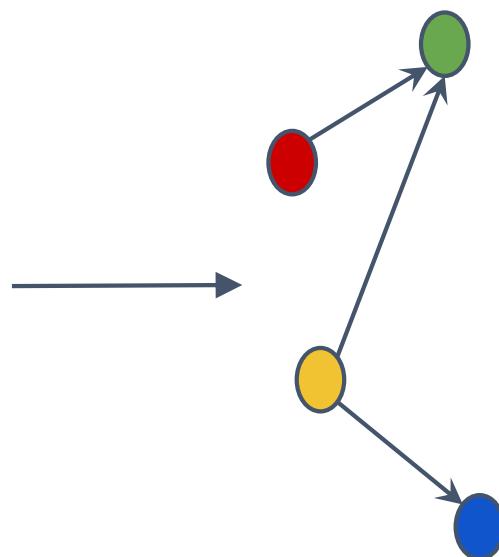
Visualisation

[1011066.Name] = [1011057.Name]
[129284.Reference] = [1223862.null] [586920.Ref.] [1030730.RCDB page] [108435.No.] [1248790.Source] [983315.References] [207338.Home railway
(external link)] [975850.Ref] [1375996.Source] [1129539.References] [1168707.References] [744488.Ref] [1169311.Ref] [1068498.Ref]
[163214.Reference] [604676.References] [1002900.Ref] [749972.Reference] [951640.References] [939700.Page] [900853.Ref] [788203.Ref]
[788409.References] [978758.Ref] [652885.Link] [652377.Ref] [1320358.Reference] [1287392.Ref] [1012269.Report] [1180077.References]
[1274408.Ref] [856227.NFL Recap] [1286480.Ref] [1354142.null] [525501.References] [630016.Notes] [762537.Refs] [902406.Report]
[1005369.Link] [1255682.Source] [1157534.Source] [1065320.Ref] [956840.Ref] [775466.References] [988811.Ref] [1005838.Link] [1005593.Link]
[576411.References] [1134428.Ref] [1170953.Reference(s)] [699144.Note] [268733.References] [931606.Notes] [1284557.Ref.] [1357973.Source]
[1238931.Report] [867400.Reference] [794774.Ref] [716064.Refs] [377521.References] [995370.Ref] [1282132.References] [1358158.Ref.]
[1120007.Ref] [1342522.Ref] [1319381.null] [889114.Ref] [1004839.Link] [697527.Website] [980509.Ref(s)] [1078901.Ref]
[1390416.Rank] = [1169921.Rank] [1183098.Rank] [1011765.Rank] [1225076.Rank] [454782.Rank] [1186535.Rank] [1209635.Rank] [1161665.Rank]
[708465.Rank] [708648.Rank]
[637307.Date] = [1311505.Date] [1337020.Date]
[1083420.Event] = [976659.Event] [976901.Event] [975917.Event] [1060037.Event] [1068182.Event] [1067251.Event] [1067097.Event] [1000067.Event]
[972968.Event] [1058267.Event] [988323.Event] [1003312.Event] [1063506.Event] [1027145.Event] [1078507.Event] [1062268.Event]
[302006.Role:] = [391330.Role:] [703281.Role:] [387497.Role:] [735612.Role:] [151885.Role:] [150598.Role:]
[1083410.Event] = [983546.Event] [975773.Event] [1071989.Event] [1068219.Event] [1002900.Event] [1074984.Event] [967160.Event] [1052352.Event]
[1066949.Event] [1082562.Event] [1151162.Event] [1042660.Event] [1056643.Event] [950860.Event] [958921.Event] [1063309.Event]
[973967.Event] [1027145.Event] [1062263.Event]
[73362.State] = [1185141.State]
[1083402.Event] = [1083339.Event] [1068498.Event] [1060027.Event] [1002823.Event] [1046135.Event] [1249836.Event] [1000145.Event]
[994576.Event] [990543.Event]
[854590.Venue] = [883202.Venue] [890993.Venue] [1104659.Venue]
[648260.TEAM] = [1286540.Club] [1308745.Club]
[627822.Division Record] = [466958.Sets W - L]
[1236345.Match] = [1231569.Match]
...

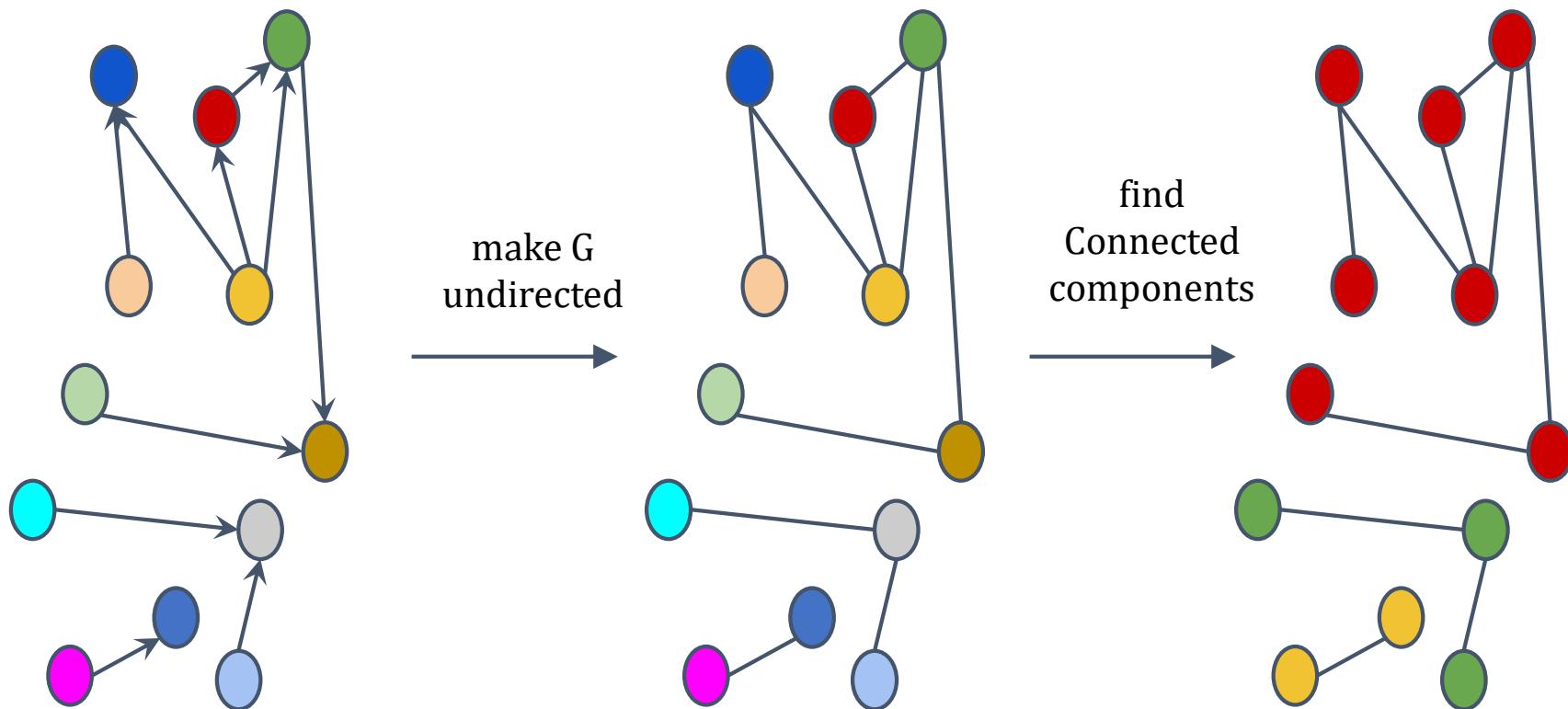
Visualisation

```
INDS = {  
R1.A ⊆ R2.B,  
R3.A ⊆ R1.D,  
R3.C ⊆ R2.A,  
R3.B ⊆ R4.A  
}
```

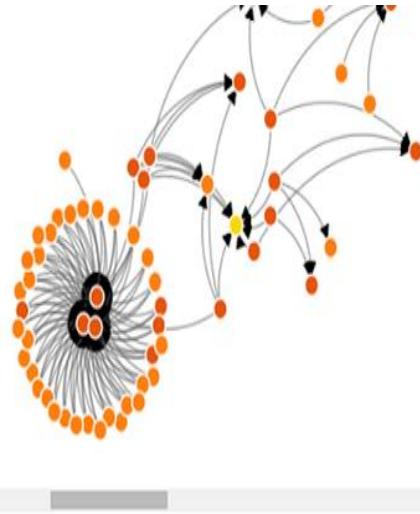
```
G= (   
V = {  
    R1, R2, R3, R4  
},  
E = { (R1, R2), (R3, R1),  
        (R3, R2), (R3, R4)  
}  
)
```



Visualisation



Interactive Application



96242-1	'Astrology_and_the_classical_elements'.csv
43666-3	43666-3.'BBC_Radio_Stoke'.Programming.csv
53064-1	53064-1.'Rotation_period'.Rotation period of selected objects.csv
562884-4	562884-4.'Planets_in_astrology'.Ruling planets of the astrological signs and houses.csv
175797-1	175797-1.'Sun_sign_astrology'.Sun signs.csv
177750-2	177750-2.'BBC_Radio_Manchester'.Programming.csv
89462-4	89462-4.'Astrology_and_the_classical_elements'.Triplicities by season.csv
213213-1	213213-1.'Dalton_Park'.Opening times.csv
	470402-

Celestial Objects	Rotation period	Rotation period
Sun	25.379995 days (equatorial) 35 days (high latitude)	25 d 9 h 7 m 11.6 s 35 d
Mercury	58.6462 days	58 d 15 h 30 m 30 s
Venus	?243.0187 days	?243 d 0 h 26 m
Earth	0.99726968 days	0 d 23 h 56 m 4.100 s
Moon	27.321661 days (synchronous toward Earth)	27 d 7 h 43 m 11.5 s
Mars	1.02595675 days	1 d 0 h 37 m 22.663 s
Ceres	0.37809 days	0 d 9 h 4 m 27.0 s
Jupiter	0.4135344 days (deep interior) 0.41007 days (equatorial) 0.41369942 days (high latitude)	0 d 9 h 55 m 29.37 s 0 d 9 h 50 m 30 s 0 d 9 h 55 m 43.63 s
Saturn	0.44403 days (deep interior) 0.426 days (equatorial) 0.443 days (high)	0 d 10 h 39 m 24 s 0 d 10 h 14 m 0 d 10 h 38 m

Zoom (1-5)

Range (logarithmic)

Dataset

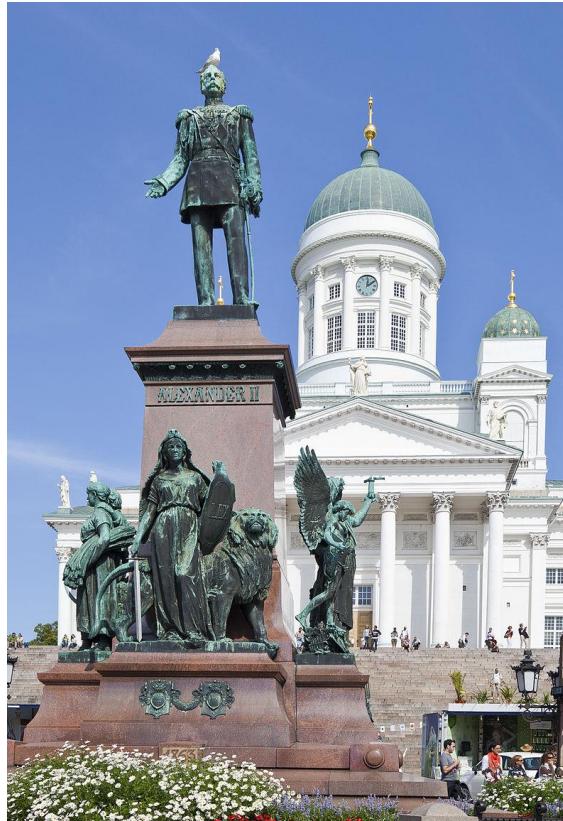
allFilters

More Dependencies

- Conditional ...
 - Uniques
 - FDs
 - INDs
- Approximate ..
 - ..
- Order dependencies [Langer, Naumann: Discovering Order Dependencies, VLDBJ'15]
- Matching dependencies [Fan et al.:Reasoning about record matching rules, VLDB'09]

Tutorial Overview

- Motivation
 - Task classification
 - Use cases
- Tools
 - Research and industry
 - Shortcomings
- Single and Multiple Column Analysis
 - Cardinalities and datatypes
 - Co-occurrences and summaries
- Dependencies
 - UCCs, INDs, FDs
 - and their discover algorithms
- Outlook
 - **Functionality**
 - **Semantics**



Part Overview

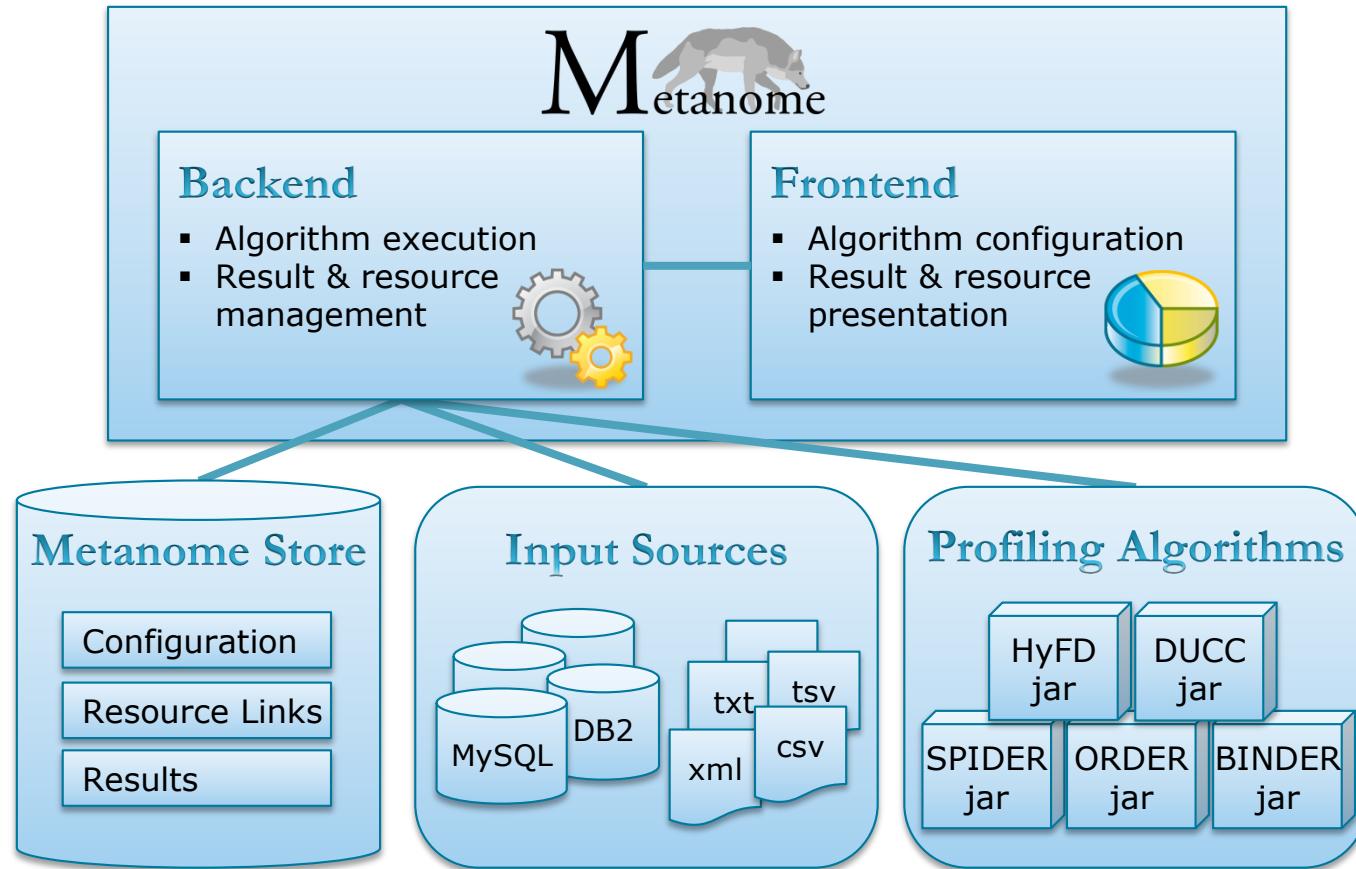
- The Metanome Data Profiling Framework
- Functional challenges
- Non-functional challenges
- Semantics of Dependencies





The Metanome Data Profiling Framework

Metanome Data Profiling Tool



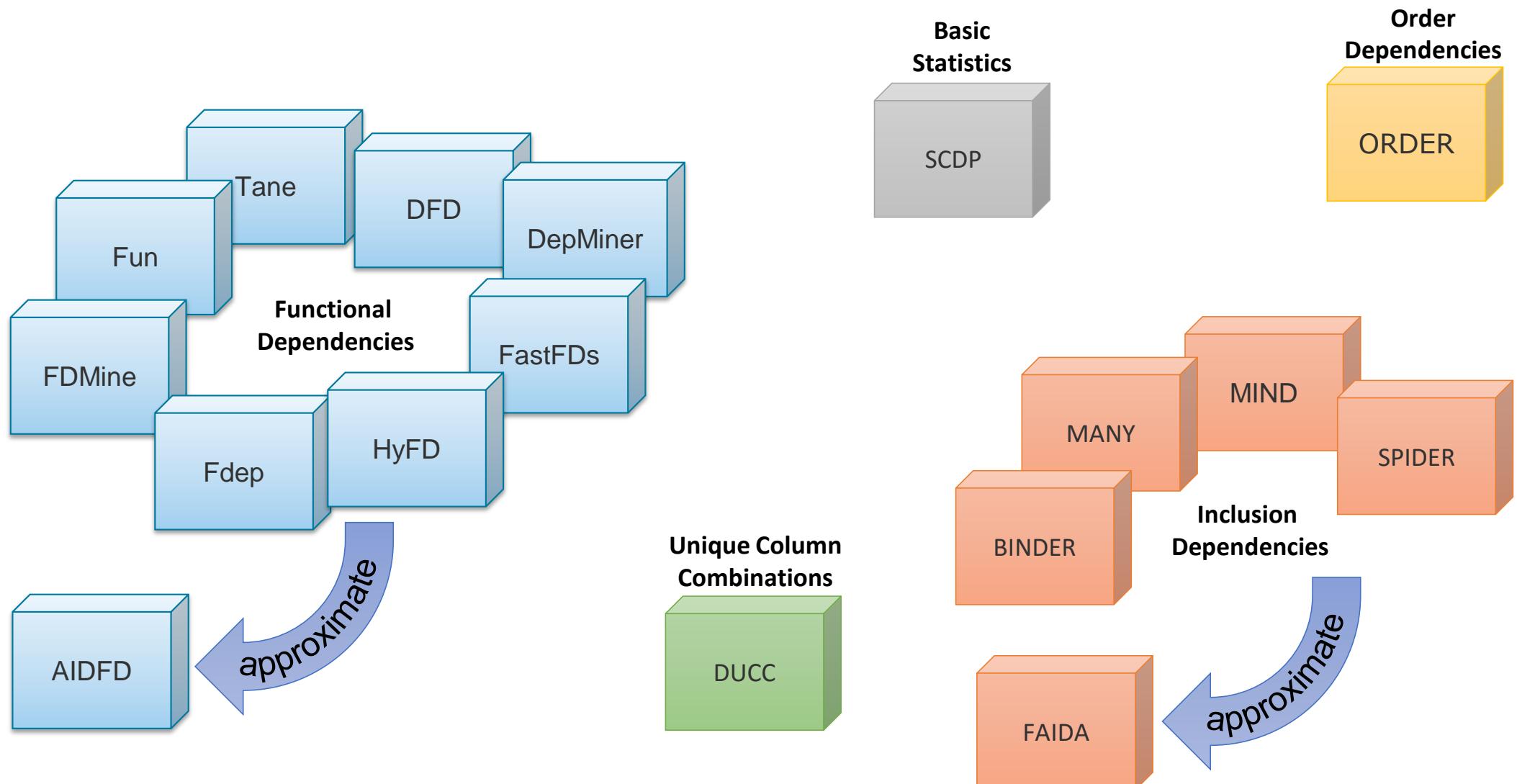
Open source framework, tool plus many algorithms

www.metanome.de

Data Profiling | ICDE 2016 | Helsinki

120

Profiling Algorithms



Metanome User Experience

The screenshot shows the Metanome web application interface. At the top, there is a navigation bar with tabs for NEW, HISTORY, and ABOUT, and a logo on the right.

Choose algorithm (Left Panel):

- Functional Dependency Algorithms
 - AIDFD-1.1-SNAPSHOT
 - Approximate FD detection
 - dfdMetanome-1.1-SNAPSHOT
 - Random Walk-based FD discovery
 - fastfds_algorithm-1.1-SNAPSHOT
 - Difference- and Agree-Set-based FD discovery
 - fdep_algorithm-1.1-SNAPSHOT
 - Dependency Induction-based FD discovery
 - fun_for_metanome-1.1-SNAPSHOT
 - Lattice Traversal-based FD discovery
 - HyFD-1.1-SNAPSHOT
 - Hybrid Sampling- and Lattice-Traversals-based FD discovery

Select datasource (Right Panel):

- File Input (choose 1)
 - MLR_abalone.csv
 - No description
 - MLR_adult.csv
 - No description
 - MLR_breastcancer.csv
 - No description
 - MLR_bridges.csv
 - No description
 - MLR_chess.csv
 - No description
 - MLR_ecgocardiogram.csv

Additional configuration (Bottom Panel):

- MAX_DETERMINANT_SIZE:
-1
- NULL_EQUALS_NULL
- VALIDATE_PARALLEL
- ENABLE_MEMORY GUARDIAN
- Result handling**
 - Cache result and write it to disk when the algorithm is finished.
 - Write result immediately to disk.
 - Just count the results.
- Memory (in MB):
- EXECUTE** button

Metanome User Experience

Metanome - Chromium

localhost:8888/#/result/1?cached=true&ind=false&fd=true&ucc=false&cucc=false&od=false&basicStat=false

NEW HISTORY RESULT ABOUT

Results for algorithm 'HyFD-1.1-SNAPSHOT.jar' executed in 115 ms

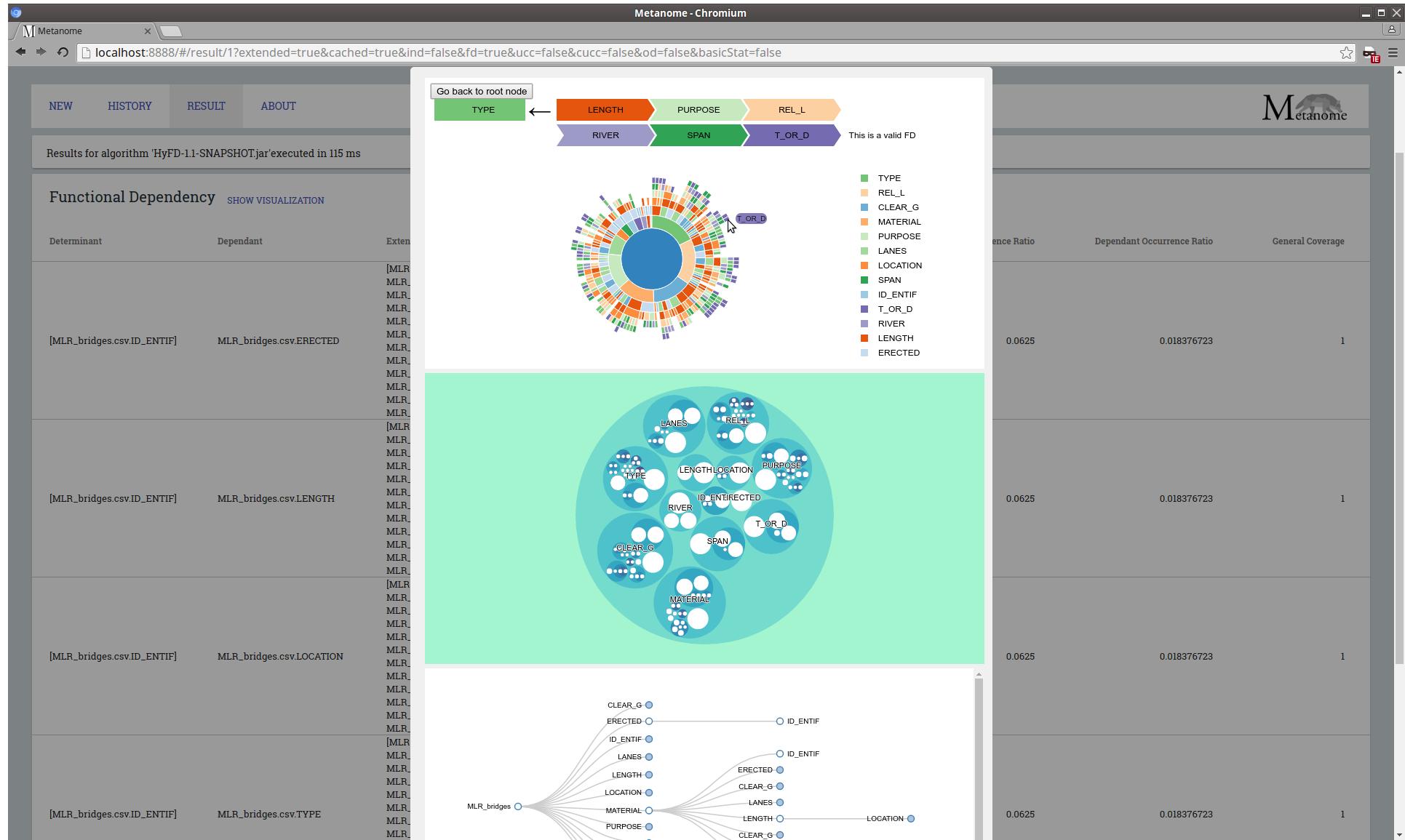
LOAD EXTENDED RESULT

Functional Dependency

Determinant	Dependant
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.ERECTED
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.LENGTH
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.LOCATION
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.TYPE
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.LANES
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.RIVER
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.PURPOSE
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.MATERIAL
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.SPAN
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.REL_L
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.CLEAR_G
[MLR_bridges.csv.ID_ENTIF]	MLR_bridges.csv.T_OR_D
[MLR_bridges.csv.ERECTED, MLR_bridges.csv.LENGTH]	MLR_bridges.csv.LANES
[MLR_bridges.csv.ERECTED, MLR_bridges.csv.LENGTH]	MLR_bridges.csv.RIVER
[MLR_bridges.csv.ERECTED, MLR_bridges.csv.LENGTH]	MLR_bridges.csv.MATERIAL

15 ▾ 1 - 15 of 142 < >

Metanome User Experience



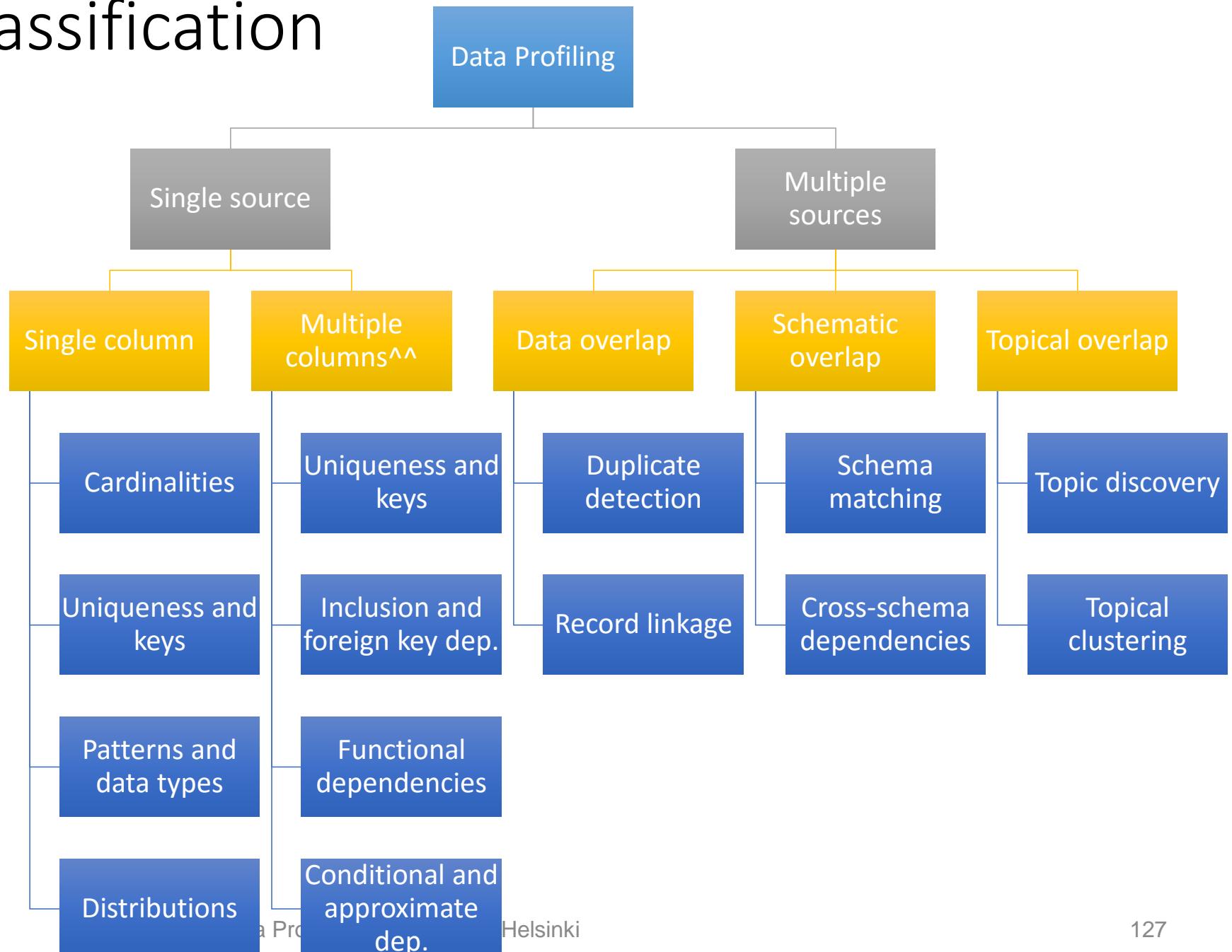
Extending the Functionality of Data Profiling



Many Other Kinds of Dependencies

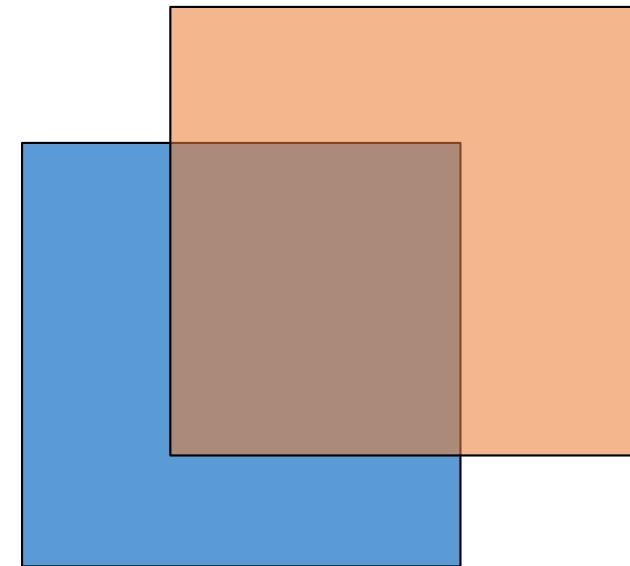
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Extended Classification of Profiling Tasks

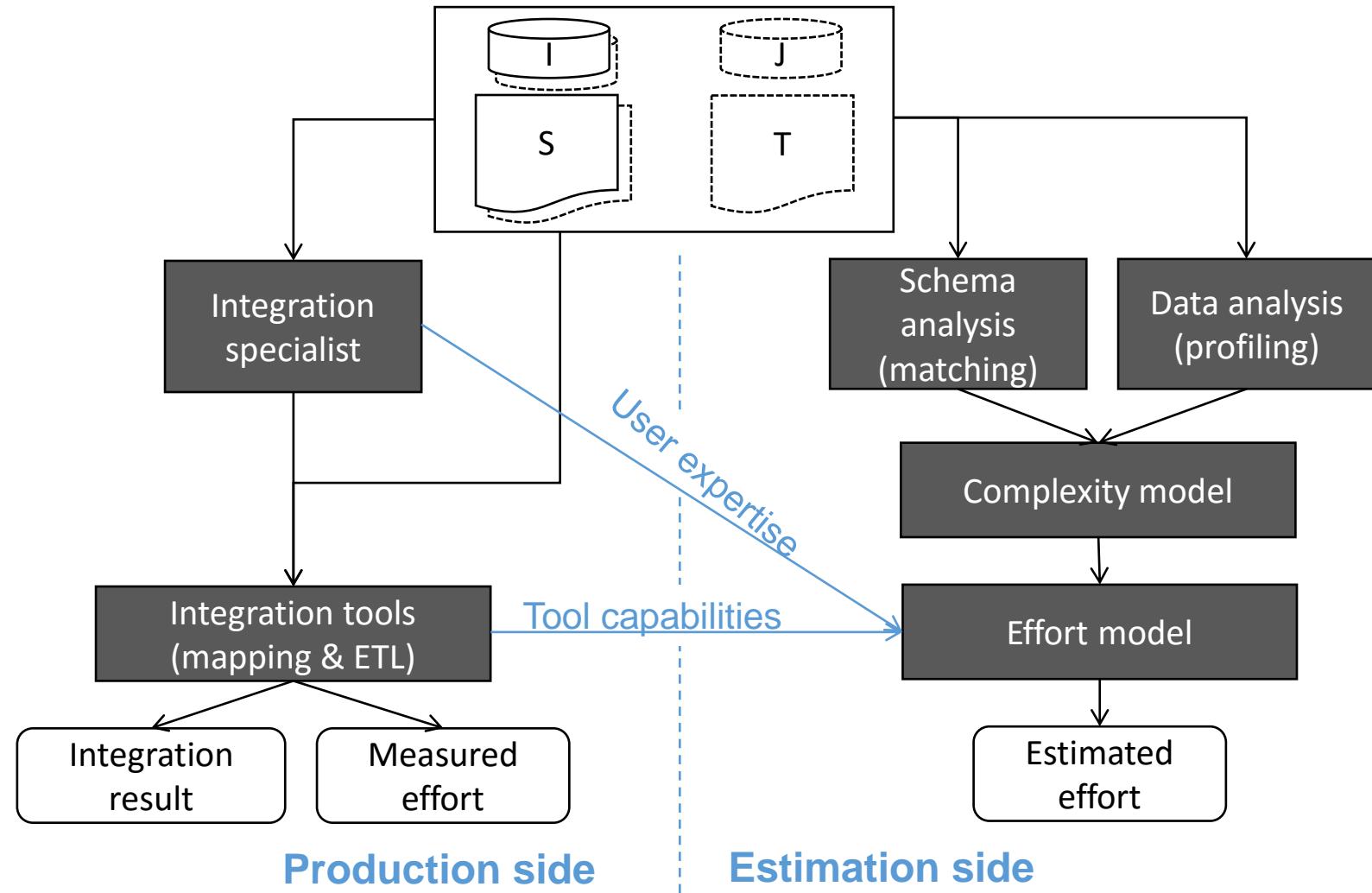


Profiling for Integration

- Create measures to estimate integration (and cleansing) effort
 - Schema and data overlap
 - Severity of heterogeneity
- Schema matching/mapping
 - What constitutes the “difficulty” of matching/mapping?
- Duplicate detection
 - Estimate data overlap
 - Estimate fusion effort
- Overall: Determine integration complexity and integration effort
 - Intrinsic complexity: Schema and data
 - Extrinsic complexity: Tools and expertise



Integration Effort Estimation



[Kruse, Papotti, Naumann: Estimating Data Integration and Cleaning Effort. EDBT 2015]

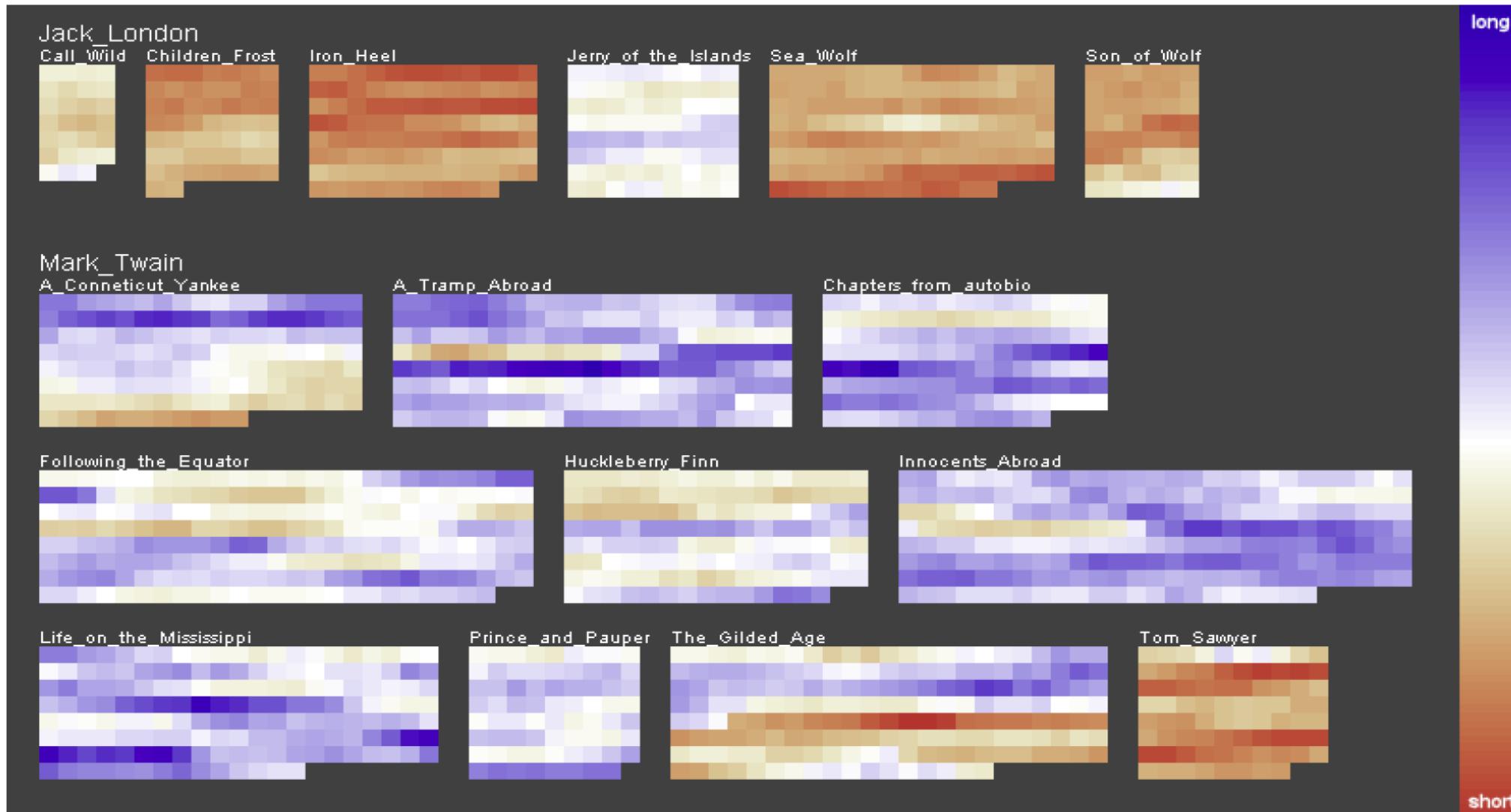
Profiling new Types of Data

- Traditional data profiling: Single table or multiple tables
- More and more data in other models
 - XML / nested relational / JSON
 - RDF triples
 - Textual data: Blogs, Tweets, News
 - Multimedia data
- Different models offer new dimensions to profile
 - XML: Nestedness, measures at different nesting levels
 - RDF: Graph structure, in- and outdegrees
 - Multimedia: Color, video-length, volume, etc.
 - Text: Sentiment, sentence structure, complexity, and other linguistic measures

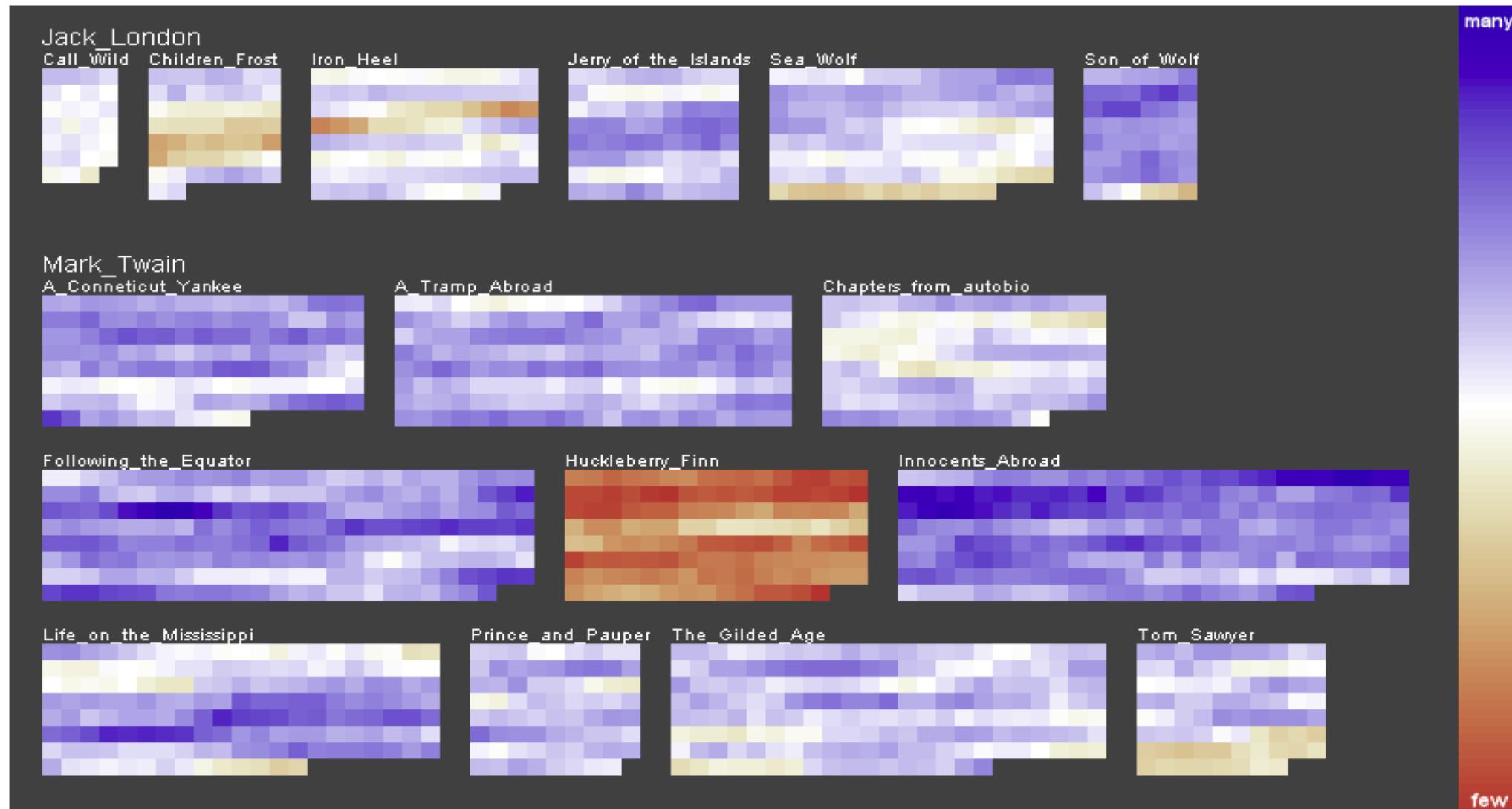
Example: Text Profiling

- Statistical measures
 - Syllables per word
 - Sentence length
 - Proportions of parts of speech
- Vocabulary measures
 - Frequencies of specific words
 - Type-token ratio
 - Simpson's index (vocabulary richness)
 - Number of hapax (dis)legomena
 - Token that occurs exactly once (twice) in the corpus
 - Characterize style of an author

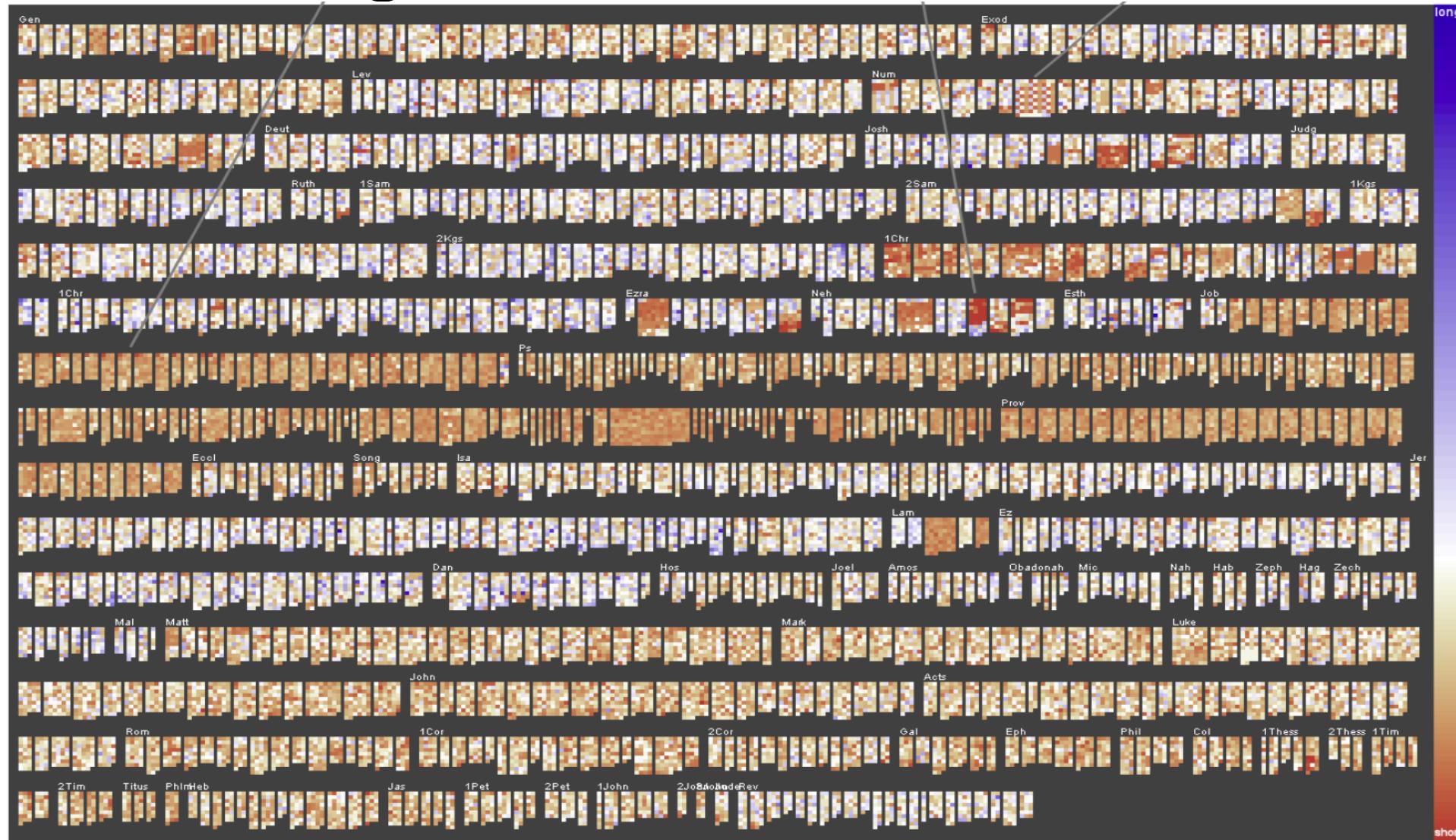
Average Sentence Length



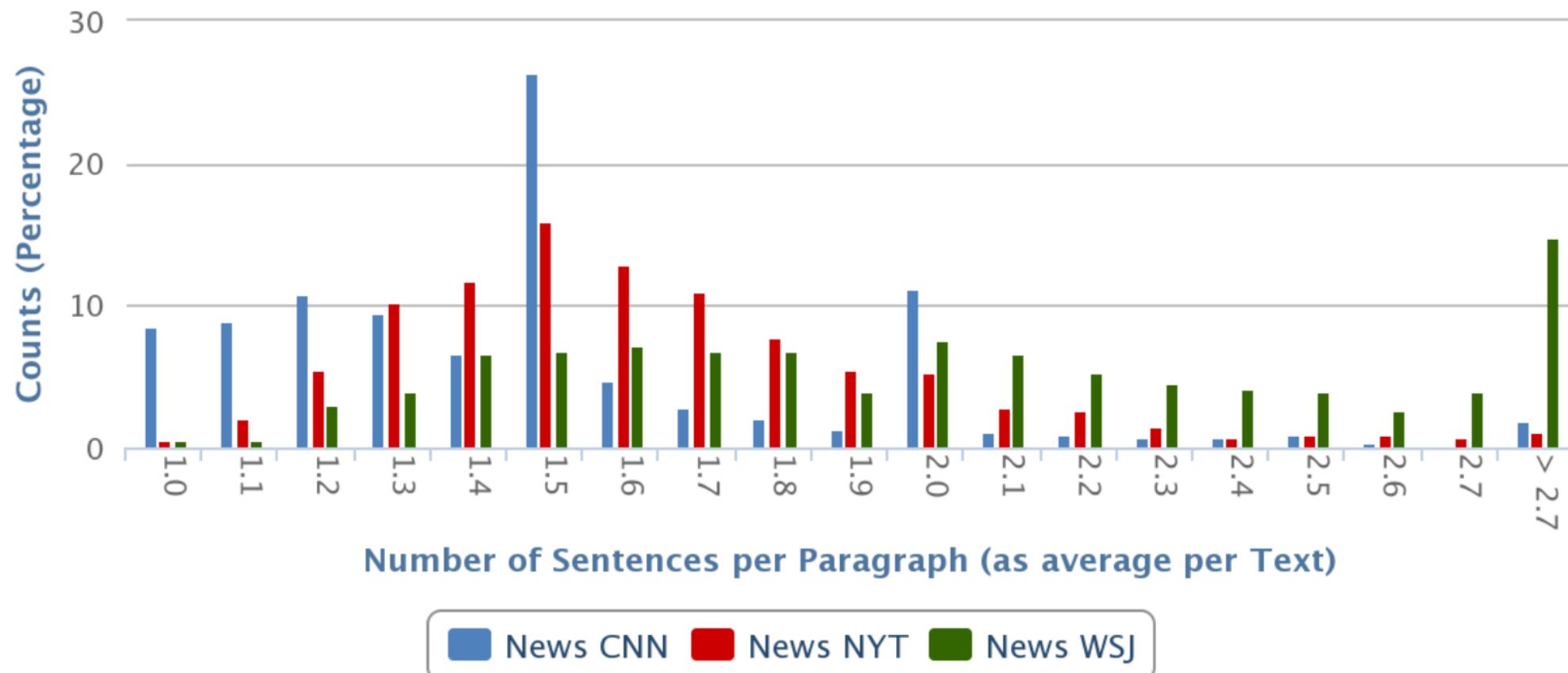
Hapax Legomena



Verse Length



Example: News Article Statistics



Improving Non-Functional Properties of Data Profiling



Profiling Challenges

- Efficient profiling
- Scalable profiling
- Holistic profiling
- Incremental profiling
- Online profiling
- Temporal profiling
- Profiling query results
- Profiling new types of data
- Data generation and testing
- Data profiling benchmark

Holistic Profiling

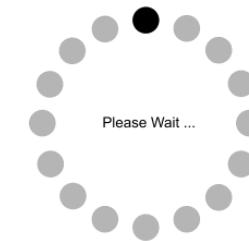
- Various profiling methods for various profiling tasks
- Commonalities/similarities
 - Search space: All column combinations (or pairs thereof)
 - I/O: Read all data at least once
 - Data structure: Some index or hash table
 - Pruning and candidate generation: based on subset/superset relationships
 - Sortation: Benefit from sorted sets
- Challenge: Develop single method to output all/most profiling results

Incremental Profiling

- Data is dynamic
 - Insert (batch or tuple-based)
 - Updates
 - Deletes
- Problem: Keep profiling results up-to-date without reprofiling the entire data set
 - Easy examples: SUM, MIN, MAX, COUNT, AVG
 - Difficult examples: MEDIAN, uniqueness, FDs, etc.

Online Profiling

- Profiling is long procedure
 - Boring for developers
 - Expensive for machines (I/O and CPU)
- Challenge: Display intermediate results
 - ... of improving/converging accuracy
 - Allows early abort of profiling run
- Gear algorithms toward that goal
 - Allow intermediate output
 - Enable early output: “progressive” profiling

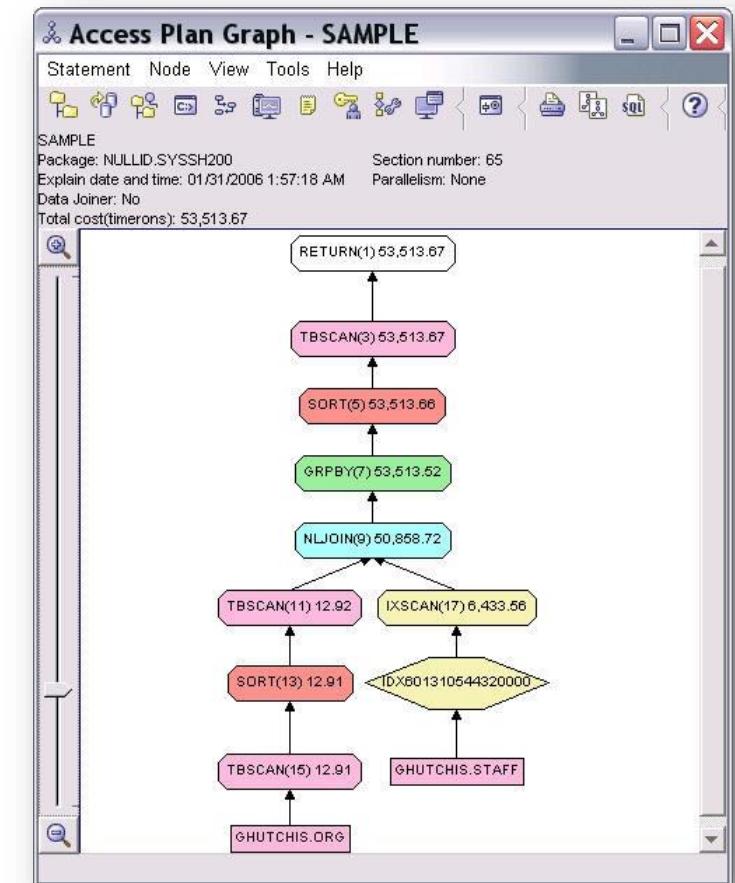


Temporal Profiling

- Observe behavior of dependencies over time
 - Do FDs appear and disappear?
 - Does a partial IND become less partial over time?
 - ...
- Metadata monitoring
 - Meta-Metadata

Profiling Query Results

- Query results are boring: Spruce them up with some metadata
 - Usually only: Row count
 - For each column, give some statistics
- Idea: Piggy-back profiling on query execution
 - Re-use sortations, hash tables, etc.



Data Generation and Testing

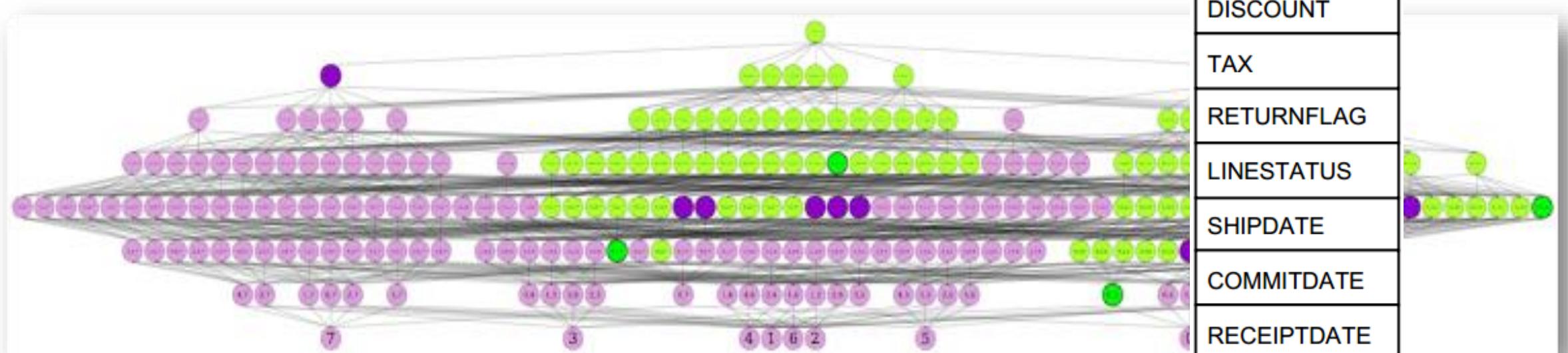
- Generate volumes of data with certain properties
 - Test extreme cases
 - Test scalability
- Problem: Interaction between properties
 - FDs vs. uniqueness
 - Patterns vs. conditional INDs
 - Distributions vs. all others...
- Problem: Create realistic data
 - Distributions, patterns
 - Placement of dependencies (tight or spread out)
 - Example: TPCH (next slide)

Recent work

[Arocena et al. : Messing Up with BART: Error Generation for Evaluating Data-Cleaning Algorithms. PVLDB 9(2), 2015]
[Arocena et al. : The iBench Integration Metadata Generator . PVLDB 9(3), 2015]

TPCH – Uniques and Non-Uniques

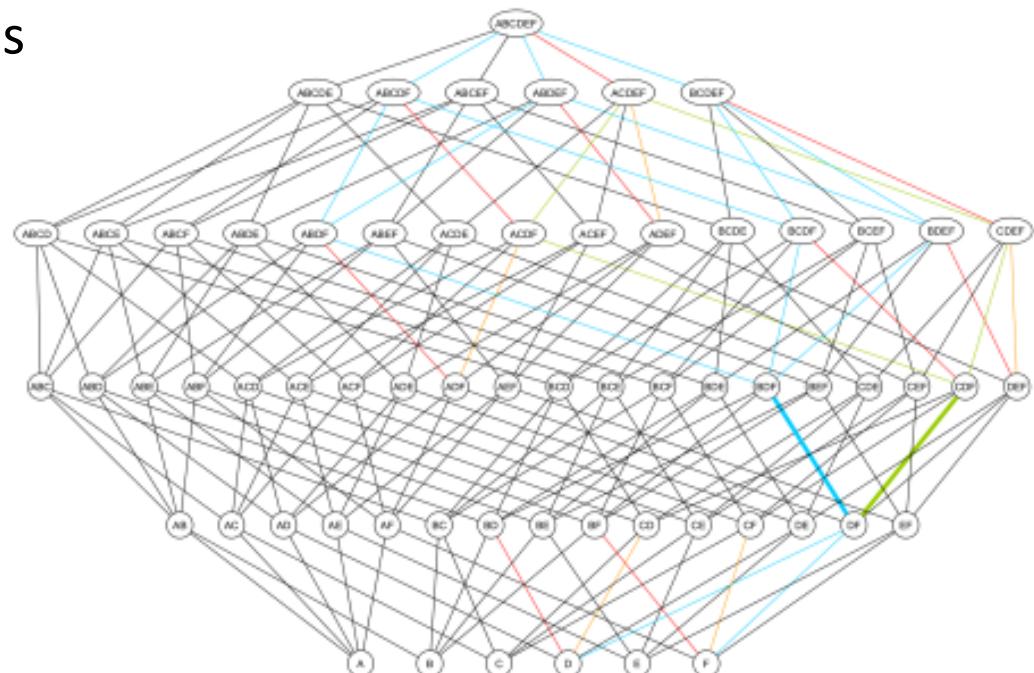
- Using the first 8 columns of the lineitems table
- Using a scale-factor of 0.1



LINEITEM (L_)
SF*6,000,000
ORDERKEY
PARTKEY
SUPPKEY
LINENUMBER
QUANTITY
EXTENDEDPRICE
DISCOUNT
TAX
RETURNFLAG
LINESTATUS
SHIPDATE
COMMITDATE
RECEIPTDATE
SHIPINSTRUCT
SHIPMODE
COMMENT

Data Profiling Benchmark

- Define data
 - Data generation
 - Real-world dataset(s)
 - Different scale-factors: Rows and columns
- Define tasks
 - Individual tasks
 - Sets of tasks
- Define measures
 - Speed
 - Speed/cost
 - Minimum hardware requirements
 - Accuracy for approximate approaches



Semantic Interpretation of Profiling Results



Turning Instance-based Observations to Schema-based Constraints

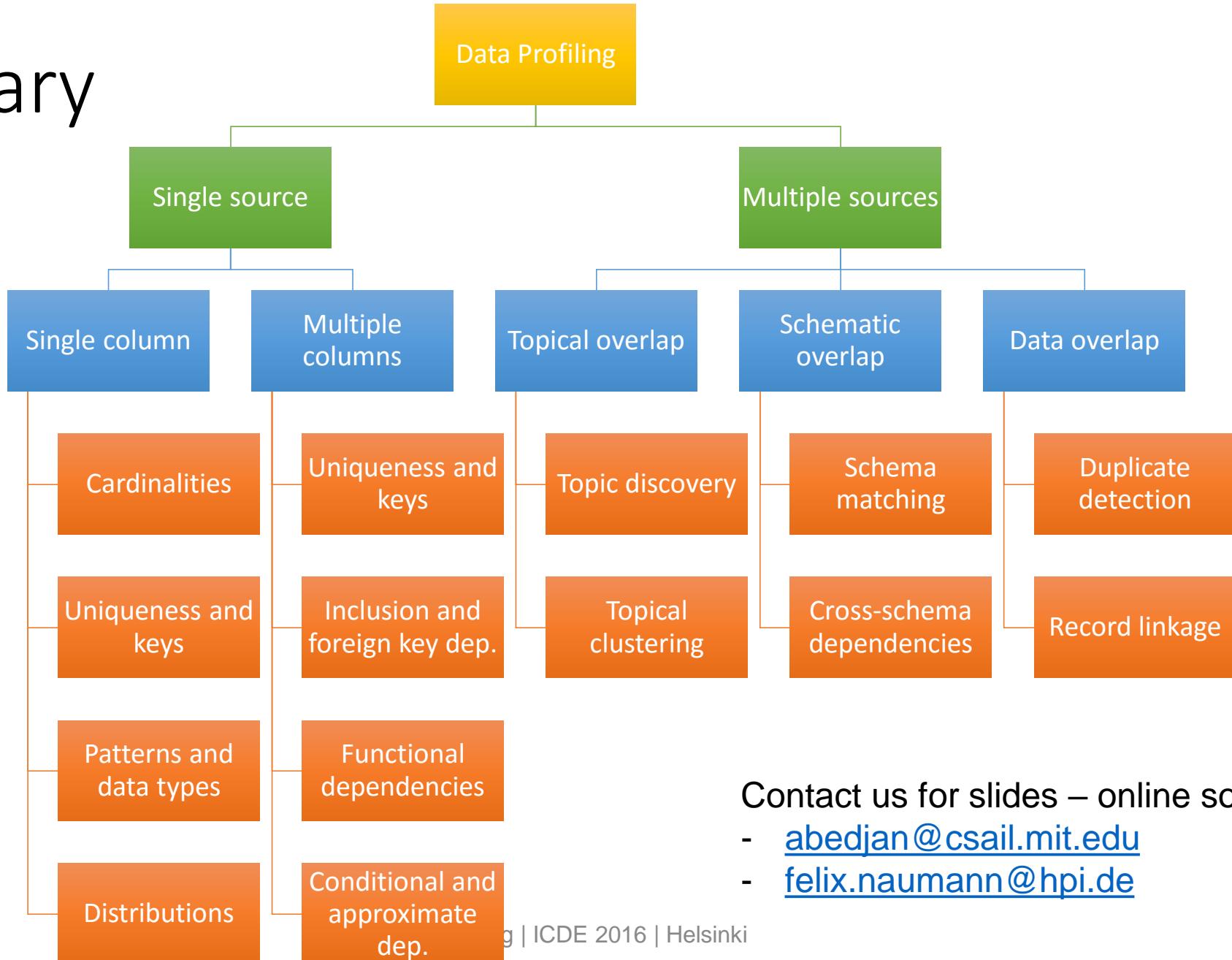
- Hundreds of UCCs – which ones are keys?
- Thousands of FDs – which ones are true?
- Millions of INDs – which ones are foreign keys?
- User-driven interpretation:
 - Rank and visualize metadata
- Machine-driven interpretation
 - Machine learning



Thanks to co-authors, colleagues and team!

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Summary



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