

Overview: Relational Abstractions

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Flavor network and the principles of food pairing

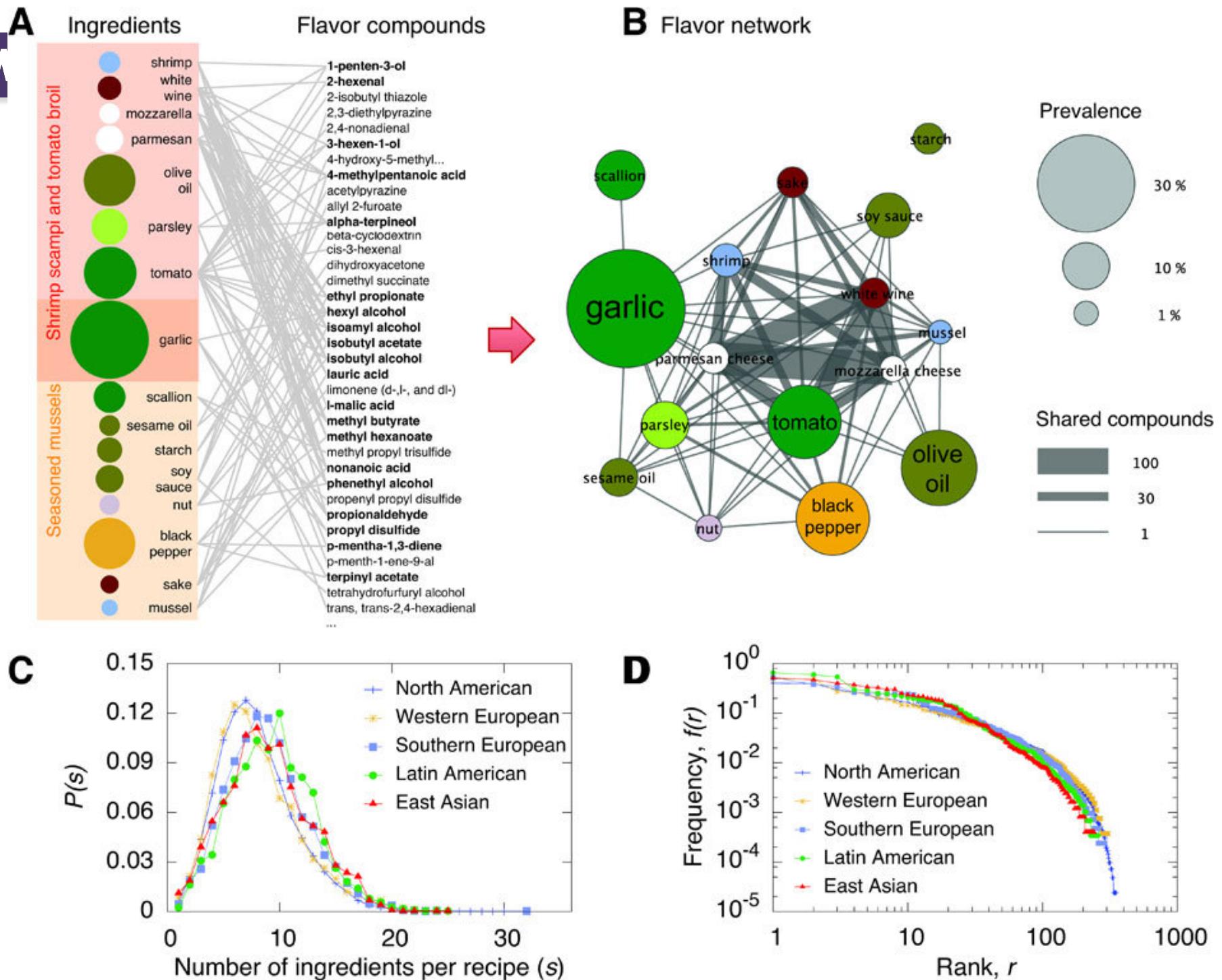
Yong-Yeol Ahn, Sebastian E. Ahnert, James P. Bagrow & Albert-László Barabási

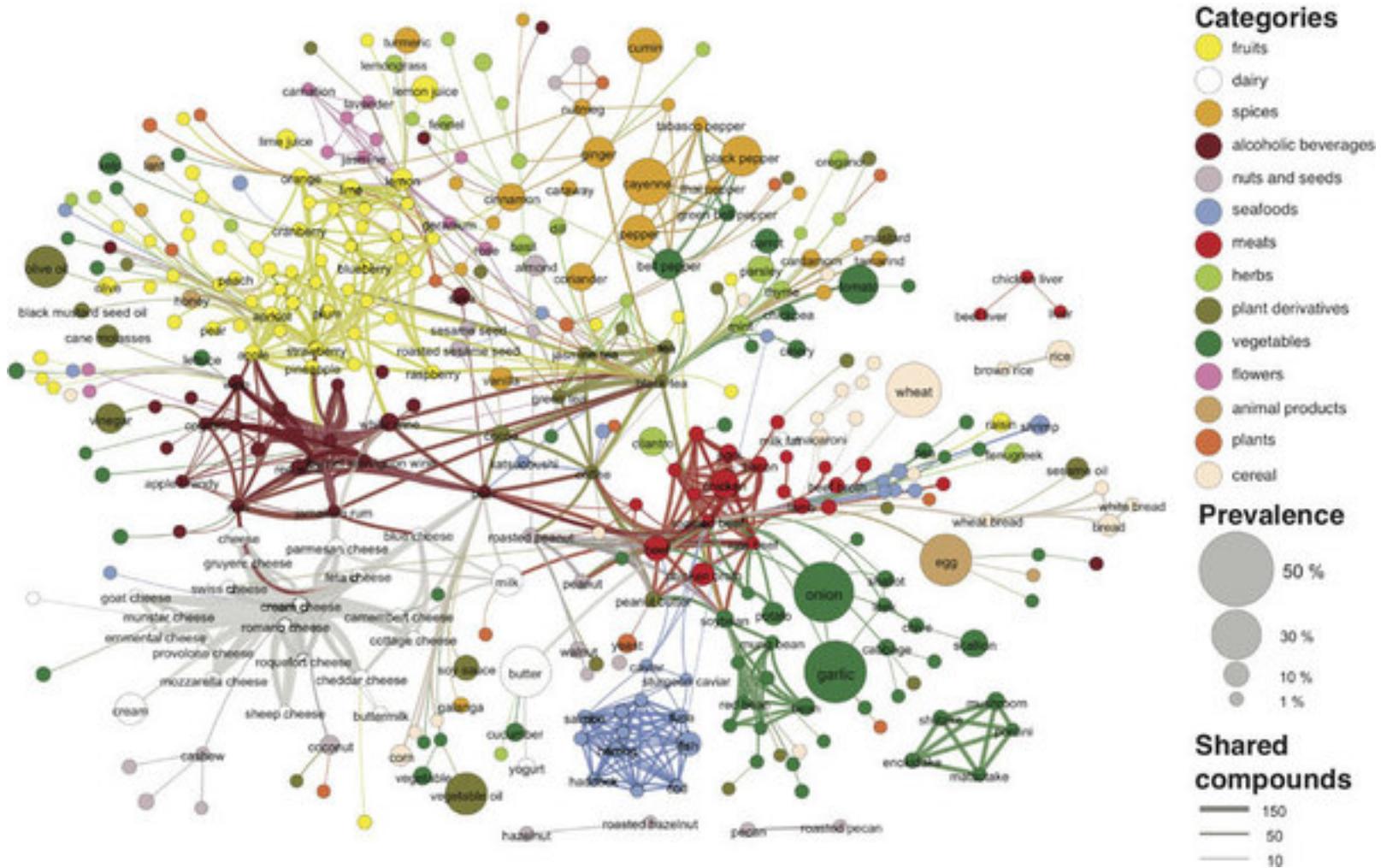
[Affiliations](#) | [Contributions](#) | [Corresponding authors](#)

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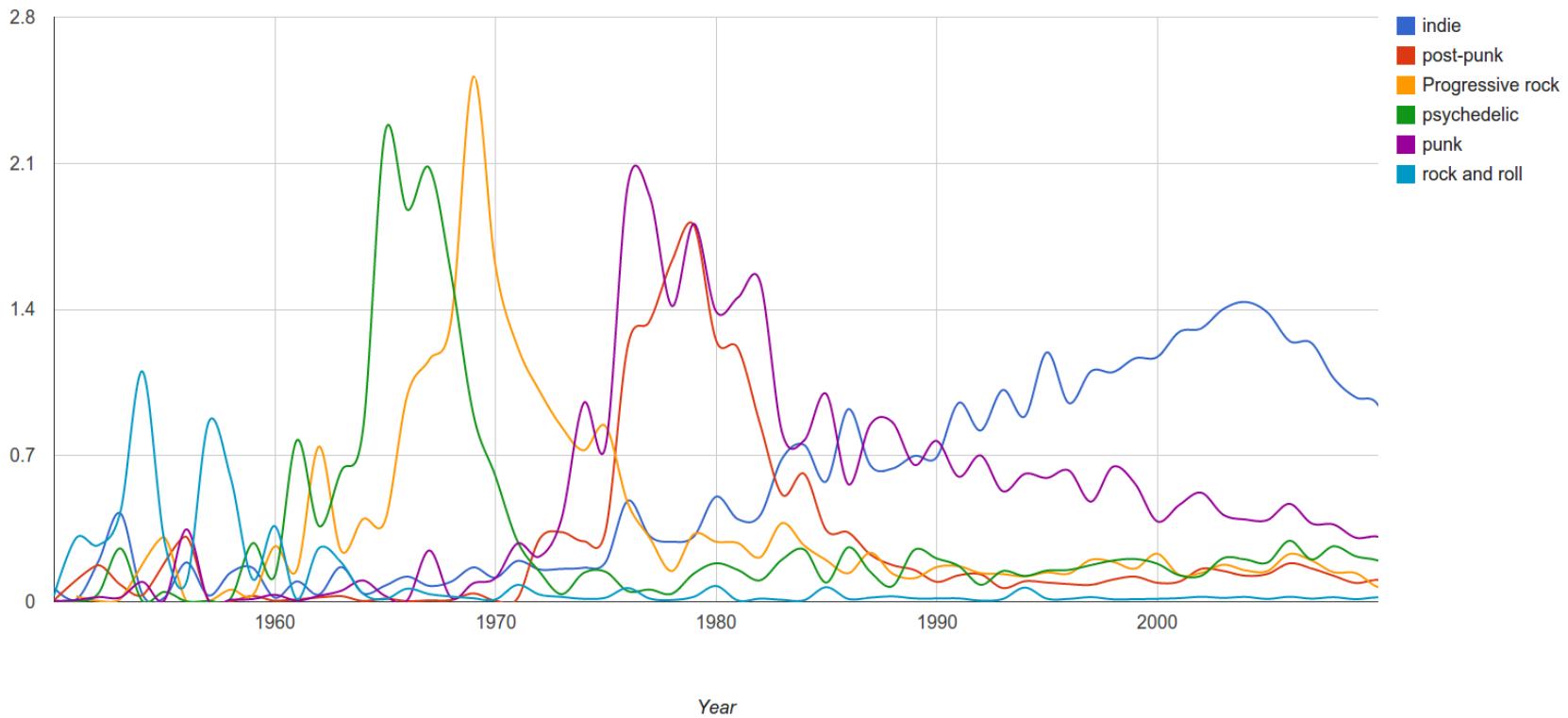
V





Last.FM

Popularity



Since we have a massive amount of user tag data available we can easily correlate tags and years and measure “popularity” of a genre by counting the number of artists formed in a specific year.

THE TWITTER POLITICAL INDEX

OCTOBER 23, 2012: OBAMA +18 [LEARN MORE](#)

 Tweet

 Follow @gov

D



BARACK OBAMA

44

↑ +2 SINCE
YESTERDAY

 Follow @BarackObama

10M+ followers

R



MITT ROMNEY

26

+0 SINCE
YESTERDAY

 Follow @MittRomney

1.5M followers

The Million Song Dataset Challenge: Part I

"people who listened to X also listened to Y"

How journalists can use data to improve the news



The Data Journalism Handbook

Edited by Jonathan Gray, Liliana Bounegru
and Lucy Chambers

L'Aquila quake: Italy scientists guilty of manslaughter

 [COMMENTS \(1022\)](#)



The BBC's Alan Johnston in Rome says the prosecution argued that the scientists were "just too reassuring"

Six Italian scientists and an ex-government official have been sentenced to six years in prison over the 2009 deadly earthquake in L'Aquila.

10/25/12

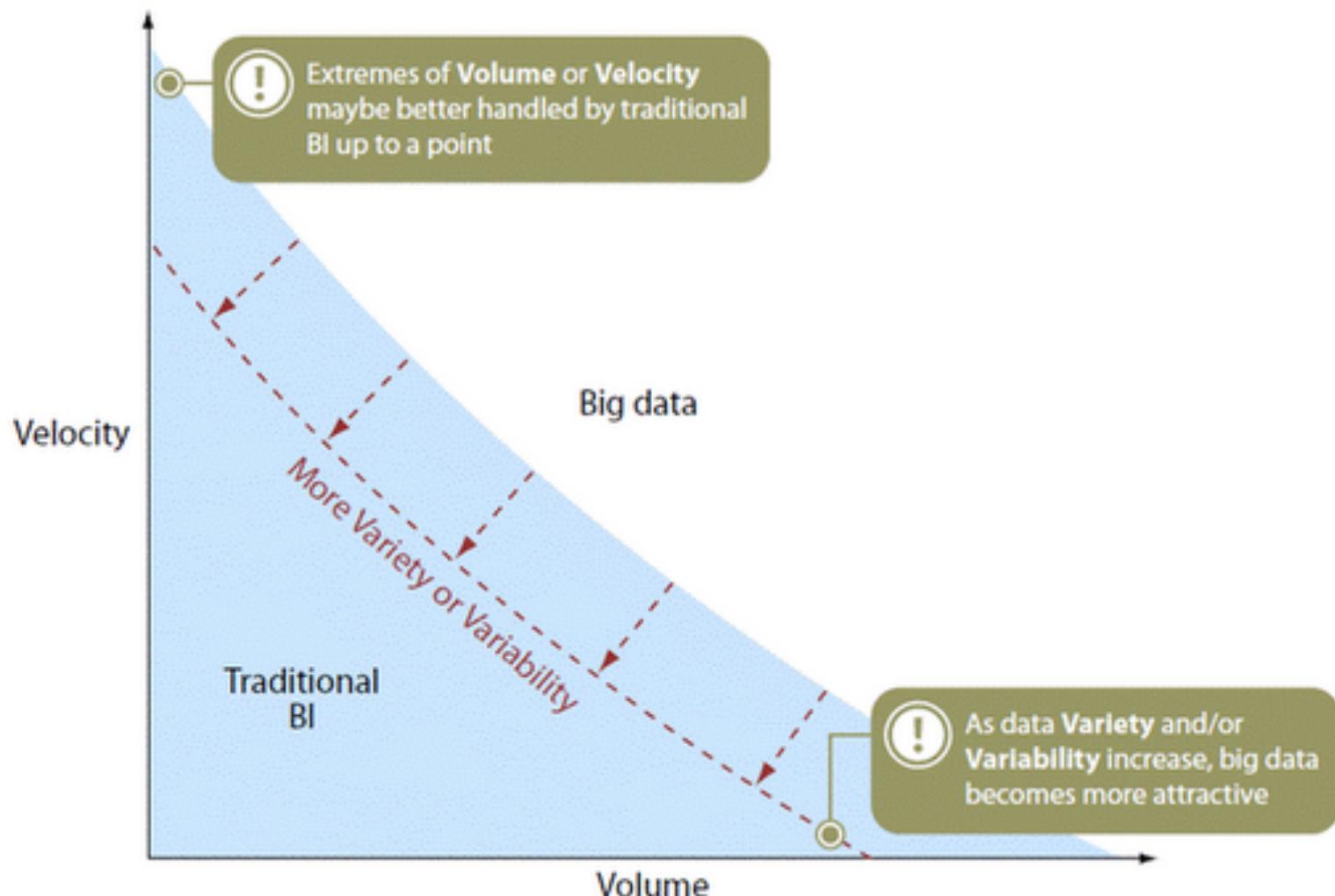
Bill Howe, UW

Related Stories

Roadmap

- Bonus slides from last lecture
- Relational Algebra and query evaluation
- In-Database Analytics
- Demo

Volume — exceeds physical limits of vertical scalability	Velocity — decision window small compared to data change rate
Variety — many different formats makes integration expensive	Variability — many options or variable interpretations confound analysis



Relational Database History

Pre-Relational: if your data changed, your application broke.

Early RDBMS were buggy and slow (and often reviled), but required only 5% of the application code.

“Activities of users at terminals and most application programs should remain unaffected when the internal representation of data is changed and even when some aspects of the external representation are changed.” -- Codd 1979

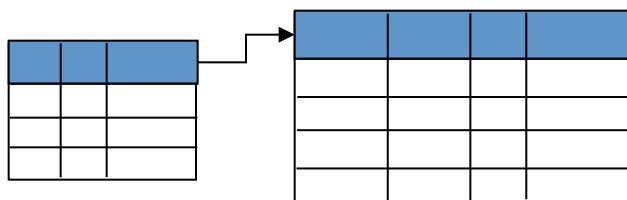
Key Ideas: Programs that manipulate tabular data exhibit an algebraic structure allowing reasoning and manipulation independently of physical data representation

Key Idea: Data Independence

views

```
SELECT *
  FROM my_sequences
```

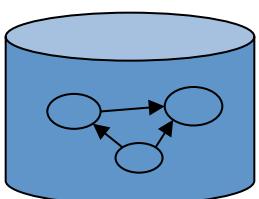
logical data independence



relations

```
SELECT seq
  FROM ncbi_sequences
 WHERE seq = 'GATTACGATATTA' ;
```

physical data independence



files and
pointers

```
f = fopen('table_file');
fseek(10030440);
while (True) {
    fread(&buf, 1, 8192, f);
    if (buf == GATTACGATATTA) {
```

Key Idea: Indexes

- Databases are especially, but not exclusively, effective at “Needle in Haystack” problems:
 - Extracting small results from big datasets
 - Transparently provide “old style” scalability
 - Your query will **always*** finish, regardless of dataset size.
 - Indexes are easily built and automatically used when appropriate

```
CREATE INDEX seq_idx ON sequence(seq) ;
```

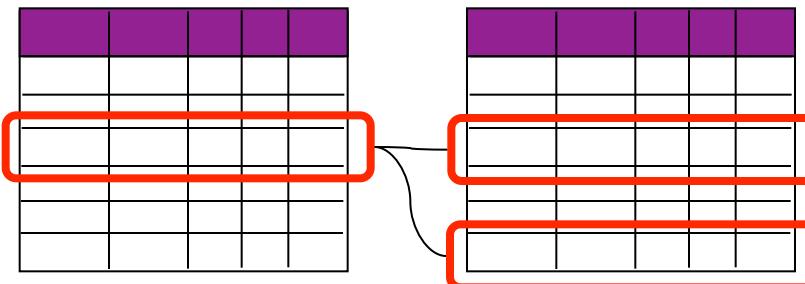
```
SELECT seq  
      FROM sequence  
     WHERE seq = 'GATTACGATATTA' ;
```

*almost

Key Idea: An *Algebra of Tables*

select

project



join

Other operators: aggregate, union, difference, cross product

Key Idea: Algebraic Optimization

$$N = ((z^*2) + ((z^*3) + 0)) / 1$$

Algebraic Laws:

1. (+) identity: $x+0 = x$
2. (/) identity: $x/1 = x$
3. (*) distributes: $(n*x + n*y) = n*(x+y)$
4. (*) commutes: $x*y = y*x$

Apply rules 1, 3, 4, 2:

$$N = (2+3)*z$$

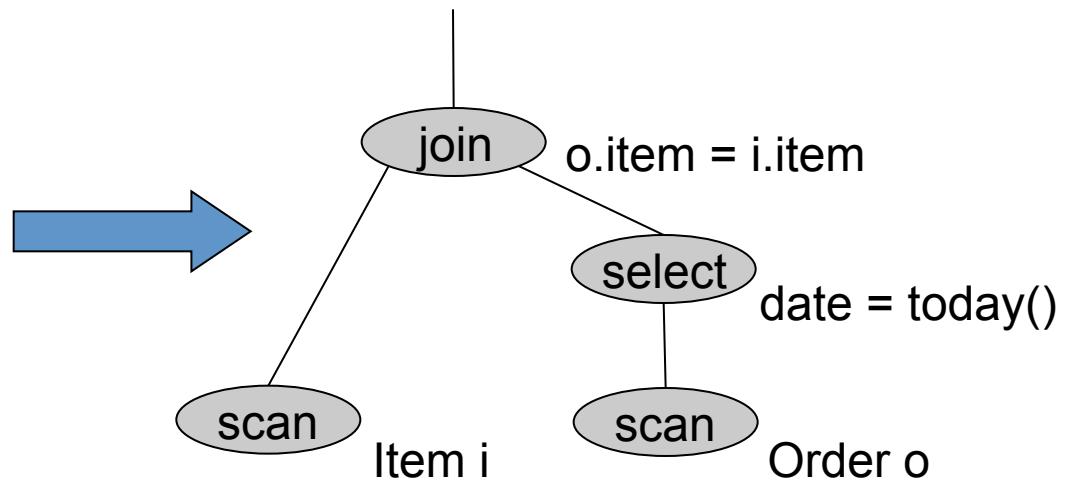
two operations instead of five, no division operator

Same idea works with the Relational Algebra!

Key Idea: Declarative Languages

Find all orders from today, along with the items ordered

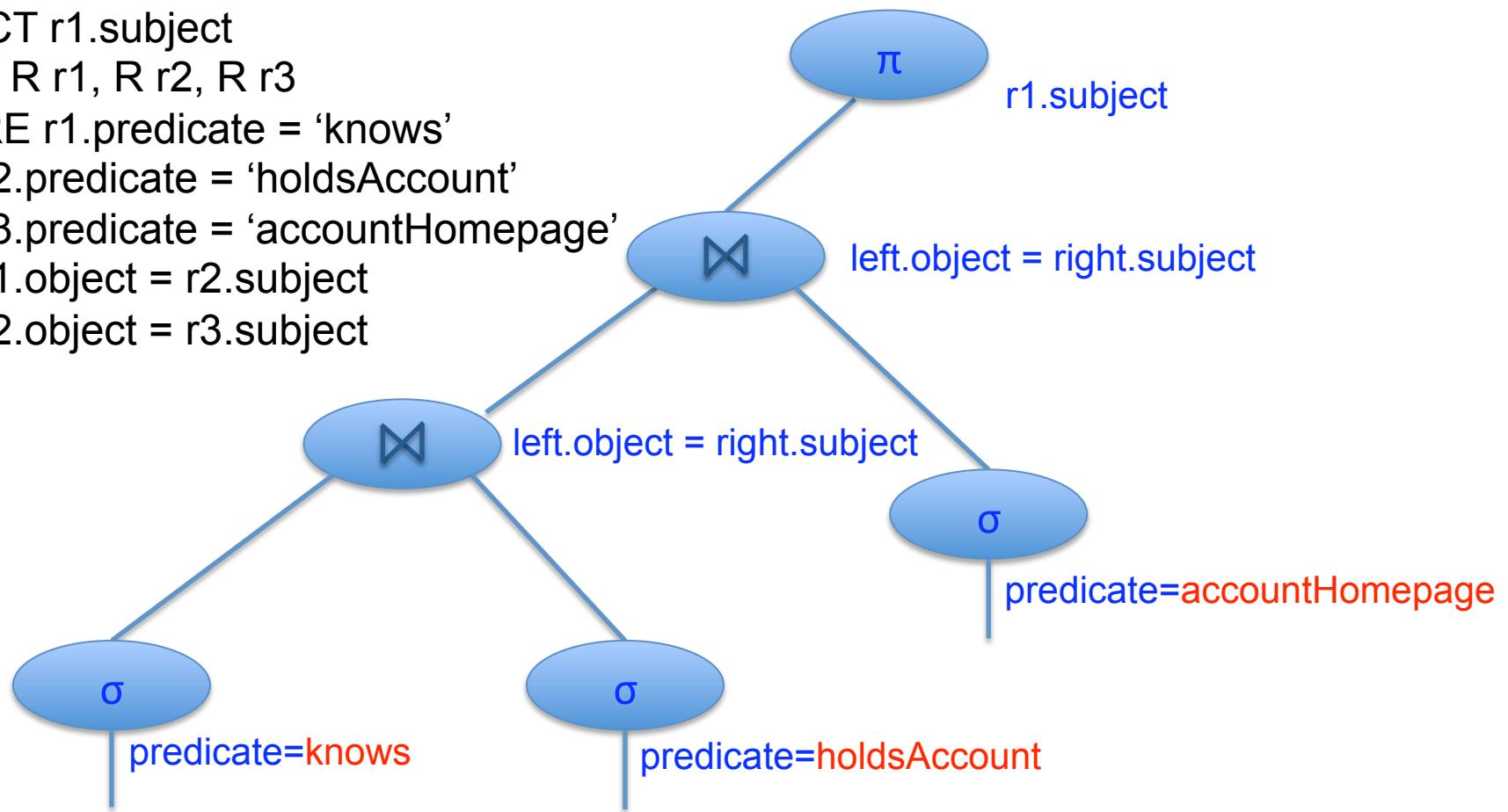
```
SELECT *
  FROM Order o, Item i
 WHERE o.item = i.item
   AND o.date = today()
```



Example Query in Relational Algebra

R(subject, predicate, object)

```
SELECT r1.subject  
FROM R r1, R r2, R r3  
WHERE r1.predicate = 'knows'  
AND r2.predicate = 'holdsAccount'  
AND r3.predicate = 'accountHomepage'  
AND r1.object = r2.subject  
AND r2.object = r3.subject
```



Equivalent logical expressions; different costs

$$\sigma_{p=\text{knows}}(R) \bowtie_{o=s} (\sigma_{p=\text{holdsAccount}}(R) \bowtie_{o=s} \sigma_{p=\text{accountHomepage}}(R))$$

right associative

$$(\sigma_{p=\text{knows}}(R) \bowtie_{o=s} \sigma_{p=\text{holdsAccount}}(R)) \bowtie_{o=s} \sigma_{p=\text{accountHomepage}}(R)$$

left associative

$$\sigma_{p1=\text{knows} \ \& \ p2=\text{holdsAccount} \ \& \ p3=\text{accountHomepage} \ \& \ r1.o=r2.s \ \& \ r2.o=r3.s} \quad (R \times R \times R)$$

cross product

Same logical expression, different physical algorithms

A = select($p=knows$)

B = select($p=holdsAccount$)

C = select($p=accountWebpage$)

hA = hash(A)

AB = probe hA with B

hC = hash(C)

ABC = probe hC with AB

A = select($p=knows$)

B = select($p=holdsAccount$)

C = select($p=accountWebpage$)

hB = hash(B)

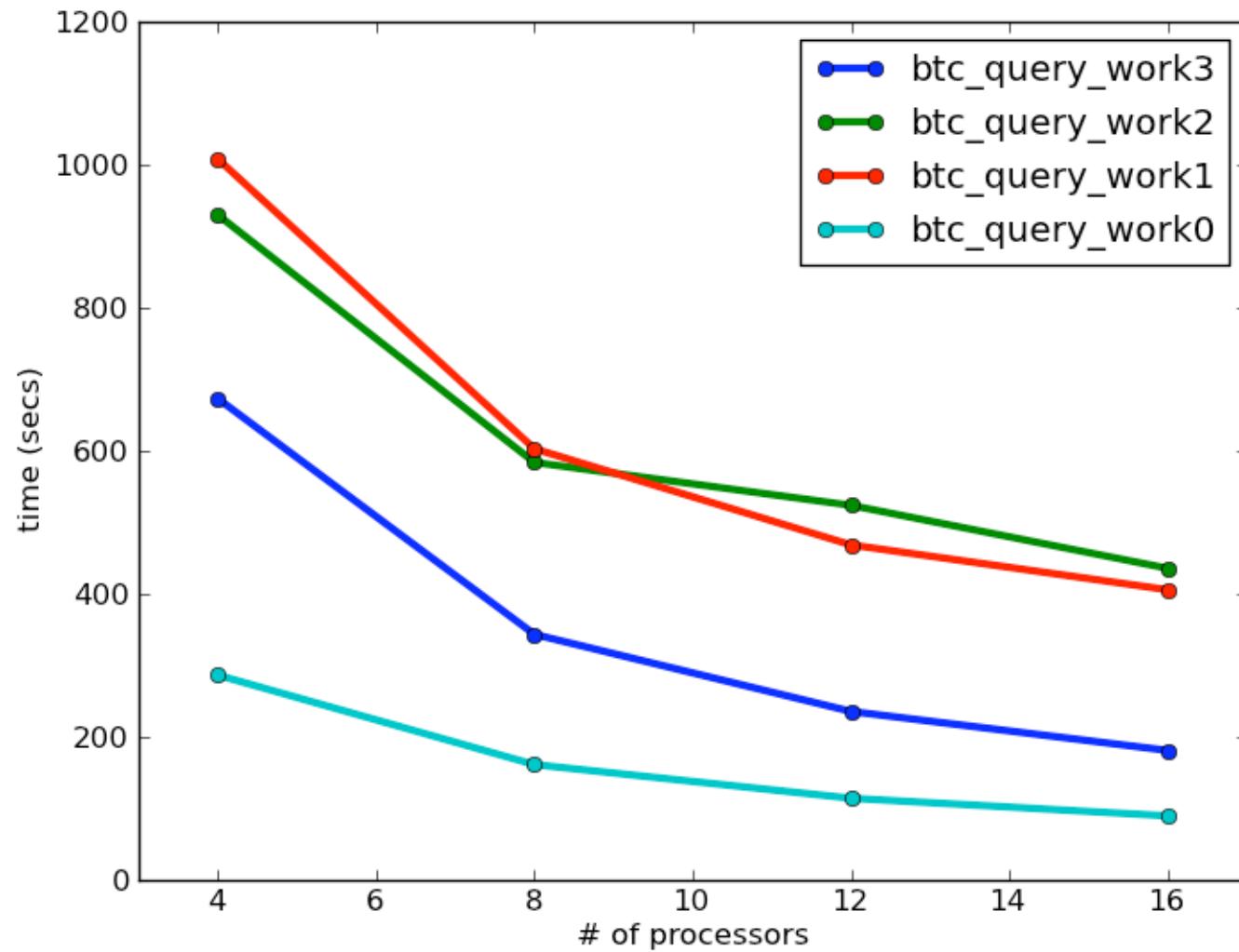
AB = probe hB with A

hC = hash(C)

ABC = probe hC with AB

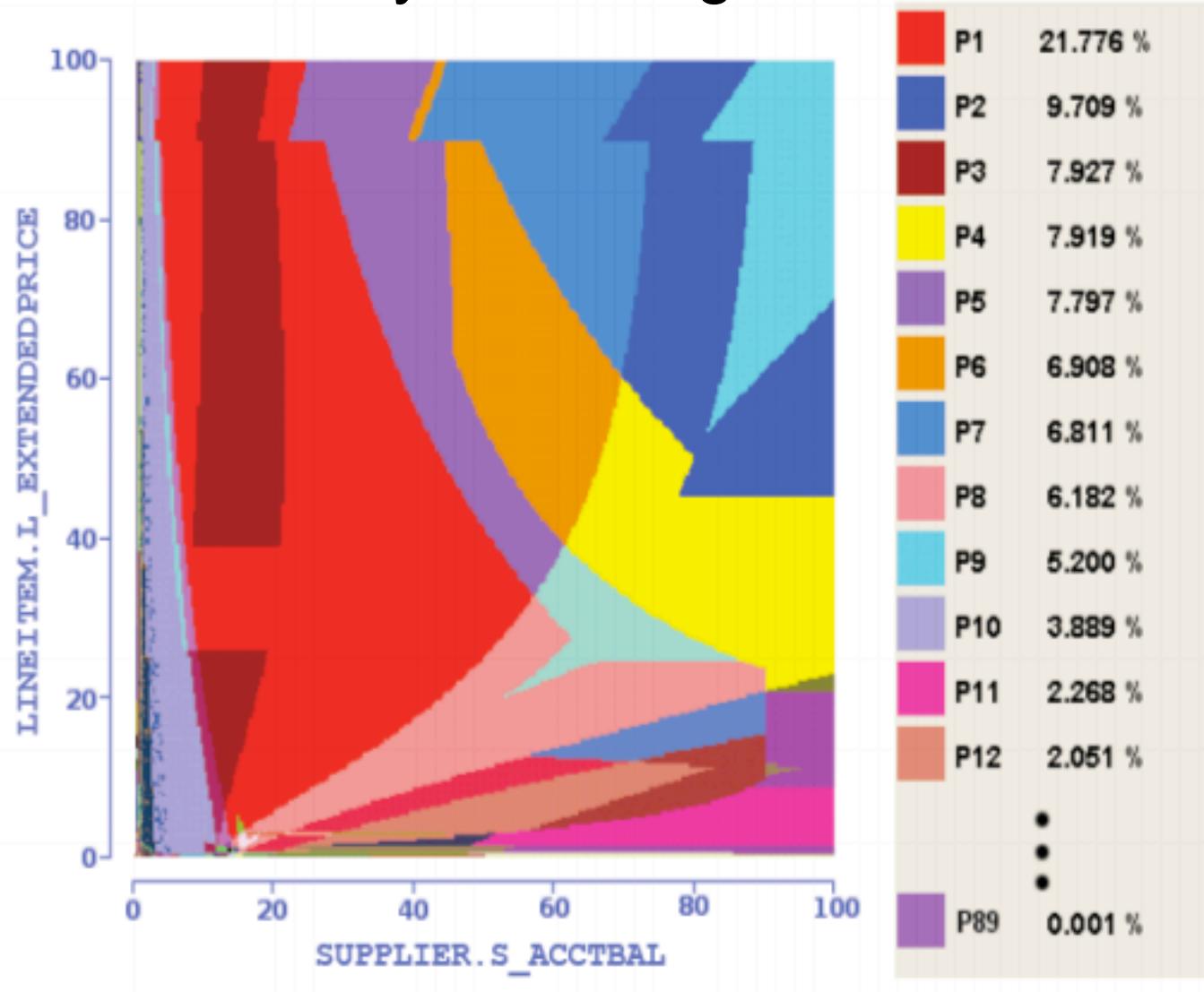
Which is faster?

Algebraic Optimization Matters



*BTC 2010 Dataset
3B quads
623 GB processed*

Picasso Query Plan Diagrams



IN-DATABASE ANALYTICS

“There is no point in bringing data ... into the data warehouse environment without integrating it. If the data arrives at the data warehouse in an unintegrated state, it cannot be used to support a corporate view of data. And a corporate view of data is one of the essences of the architected environment.”

Inmon, 2005, “Building the Data Warehouse”

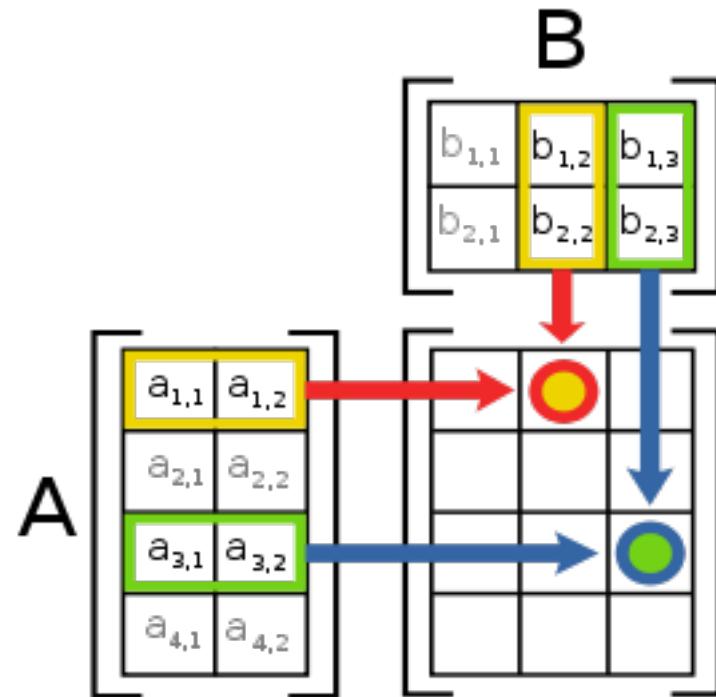
Not a good fit for analytics!

Matrix Addition, vector representation

A(row_number, row float[])
B(row_number, row float[])

```
SELECT A.row_number, A.vector + B.vector  
FROM A, B  
WHERE A.row_number = B.row_number;
```

Matrix Multiplication



Matrix multiplication, vector representation

A(row_number, row float[])

SELECT 1, array_accum(row_number, vector*v) FROM A;

$$\vec{x} \cdot \vec{y} = \sum_i x_i y_i$$

Matrix Transpose, vector representation

A(row_number, row float[])

```
SELECT S.col_number, array_accum(A.row_number, A.vector[S.col_number])
FROM A, generate_series(1,3) AS S(col_number)
GROUP BY S.col_number;
```

generate_series is a *table function*

Matrix Multiplication, sparse representation

```
A(row_number, column_number, value)  
B(row_number, column_number, value)
```

```
SELECT A.row_number, B.column_number, SUM(A.value * B.value)  
FROM A, B  
WHERE A.column_number = B.row_number  
GROUP BY A.row_number, B.column_number
```

Aside: User-defined functions and types

- Scalar functions

```
SELECT myfunc(r.a, r.b) ...
```

- Aggregate functions

```
SELECT concat(r.s) ...
```

- Table functions

```
SELECT ... FROM tablefunc(a,b)
```

Experiment design

```
CREATE VIEW design AS
SELECT a.trial_id,
       floor (100 * random()) AS row_id
  FROM generate_series(1,10000) AS a (trial_id),
       generate_series(1,3) AS b (subsample_id)
```

```
CREATE VIEW trials AS
SELECT d.trial_id, AVG(a.values) AS avg_value
  FROM design d, T
 WHERE d.row_id = T.row_id
 GROUP BY d.trial_id
```

src: Cohen et al., VLDB 2009

Prior to the implementation of this functionality within the DBMS, one Greenplum customer was accustomed to calculating the OLS by exporting data and importing the data into R for calculation, a process that took several hours to complete. They reported significant performance improvement when they moved to running the regression within the DBMS. Most of the benefit derived from running the analysis in parallel close to the data with minimal data movement.

src: Cohen et al., VLDB 2009

Assignment

- Matrix Addition for Sparse Representation in SQL
 - Assume the following schema:
A(row_number, column_number, value)
B(row_number, column_number, value)
- **Bonus:** tf-idf in a series of SQL statements
 - Assume the following schema:
frequency(term, docid, count)

Query Database-as-a-Service for the 99%



Approach: Strip down databases to bare essentials

- Upload -> Query -> Share
 - Try to eliminate **installation, configuration, schema design, data loading, tuning, app-building**
easy, thanks to the cloud
- harder

```

SELECT substring(d.name, 1, 5) as construct
,substring(d.name, 7, 5) as glycerol_id
,
FROM [billhowe].[dnasamples.csv] d
    
```

Your datasets

All datasets

Shared datasets

Recent activity... 2

Recently viewed »

Getting Data In

- Upload data through a browser to a cloud database (so no need to manage a local system)
- No need to “design a database” before you use your data – just upload and get to work
- Write SQL queries, with some automated help and some guided help from experts
- Build on your own results
 - Output of one query can be the input to another
 - Encourages sharing and reuse
 - Avoids everyone on the team running the same data processing steps over and over
 - Provides provenance – “how did you get this result?”



The screenshot shows the SQLShare web application interface. On the left, there's a sidebar with links like 'Your datasets', 'All datasets', 'Shared datasets', 'Recent activity...', 'Recently viewed', 'Upload dataset', and 'New query'. Below that is a section for 'YOUR TOP VIEWED' datasets. On the right, the main area is titled 'Your Datasets' and lists various datasets with their descriptions and last modified dates. A blue box highlights the first dataset, 'GO0005515_domains'. A red box highlights the second dataset, 'GO0005515_112gene_blast_nr.csv'. A red circle with a blue outline is drawn around the 'GO0005515_112gene_blast_nr.csv' entry.

Name	Description	Last Modified
GO0005515_domains	How many genes have which domain	Apr 18, 2011 6:58 PM
WD other domains	These are the unique domains that showed up with WD domain	Apr 21, 2011 2:34 PM
domains per gene		Apr 21, 2011 2:58 PM
GO0005515_112gene_blast_nr.csv	blast results for the 112 genes in the enriched G	Apr 21, 2011 3:54 PM
GO0005515_best_anno	These annotations have had all the unknowns and predicted removed	Apr 22, 2011 11:10 AM
GO0005515_lowest_eval_best_anno	With all the unknowns and predicted removed	Apr 22, 2011 11:11 AM
GO0005515_lowest_eval	With all the unknowns and predicted removed	Apr 22, 2011 11:17 AM
GO0005515_lowest_eval	With all the unknowns and predicted removed	Apr 22, 2011 11:17 AM
146_enriched_gene_allbest_annotation	no unknowns or predicted are included	Apr 22, 2011 4:14 PM
146_enriched_gene_lowest_evals	no unknowns or predicted are included ==92 g	Apr 22, 2011 4:22 PM
146_enriched_best_anno_lowest_eval	Best blasts excluding unknowns and predicted	Apr 22, 2011 4:31 PM
809_interpro2GO0005515_sort.csv	809_enrich_GO0005515	Apr 22, 2011 5:03 PM

Select from a list of English descriptions

Edit a Query

```
SELECT interpro_description, count(DISTINCT gene_id) as num_genes
FROM [koesterj@washington.edu].[809_interpro2GO0005515_sort.csv]
GROUP BY interpro_description
order by num_genes DESC
```

Your datasets
All datasets
Shared datasets
Recent activity... 0
Recently viewed »

Upload dataset
New query

YOUR TOP VIEWED
exon.feature.o... 36
Lengths_hotsp... 28
noncoding_po... 25
hotspot_desert... 17
HD_NC_overla... 16

POPULAR TAGS
protein 49
oceanography 23
fluorimeter 16
suna 16

GO0005515_domains

Last modified: Apr 21, 2011 2:34 PM koesterj@washington.edu

How many genes have which domain

[Click here to add a tag](#)

```
SELECT interpro_descripti
FROM [koesterj@washington
GROUP BY interpro_descrip
order by num_genes DESC
```

Save the results, share them with others

DATASET PREVIEW Rows 1 - 100 of 148 | Columns 2 of 2

<< first < prev 1 2 3 4 5 next > last >>

interpro_description	num_genes
NULL	112
WD40/YVTN repeat-like-containing domain	18
WD40 repeat-like-containing domain	14
Zinc finger RING/FYVE/PHD-type	13
WD40 repeat subgroup	13
Tetratricopeptide-like helical	13
WD40 repeat	12
Tetratricopeptide repeat	11
Tetratricopeptide repeat-containing	9
WD40-repeat-containing domain	9

Your datasets[All datasets](#)[Shared datasets](#)[Recent activity... 2](#)[Recently viewed »](#)[Upload dataset](#)[New query](#)**YOUR TOP VIEWED**

csv2.csv 115

csv2.csv 115

blackhole 16

Vizlet Scores 14

vizlets_23nov... 14

POPULAR TAGS

biomed 138

ht_screening_r... 81

seqvalidation 52

protein 47

oceanography 23

tsg 16

suna 16

Your DatasetsFilter dataset by keyword:

Name	Sharing / Owner	Modified	
topic.csv research topics simpleschema	billhowe@washington.edu	Feb 24, 2012 9:04 AM	
mhip_zip_eScience_022112a.csv additional outcome measures for mhip dataset mhip	billhowe@washington.edu	Feb 21, 2012 5:05 PM	
total students taking AMATH301 and CSE142 csse	billhowe@washington.edu	Feb 4, 2012 11:46 PM	
total students taking amath301 prior to cse142 csse	billhowe@washington.edu	Feb 4, 2012 11:46 PM	
amathe_analysis.csv anonymized course registrations for AMATH301 and CSE142 csse	billhowe@washington.edu	Feb 4, 2012 11:46 PM	
elements_with_atomic_numbers_92_and_below.csv test dataset for alicia	billhowe@washington.edu	Jan 20, 2012 1:45 PM	
SeaFlow Example Dataset Clean SeaFlow Example Dataset seawflow	billhowe@washington.edu	Jan 20, 2012 12:40 PM	
categorized_fat.xlsx.txt health	billhowe@washington.edu	Dec 7, 2011 1:06 PM	
Vizlet Scores and Features Score is the number of promote actions for each vizlet type for each column p vizdeck	billhowe@washington.edu	Dec 2, 2011 1:40 AM	
VizDeck User Study Timing and Success vizdeck	billhowe@washington.edu	Dec 1, 2011 11:56 PM	
vizstudy_analysisv7.csv	billhowe@washington.edu	Dec 1, 2011 11:44 PM	
Vizlet Scores Score of each (session_x column_y column_z vizlet_type)			

<http://sqlshare.escience.washington.edu>

Upload Dataset

1. Choose File

2. Import Settings

3. Save

File:

2010.csv

7.47 MB



Analysing your file...

[Cancel](#)

<http://sqlshare.escience.washington.edu>

Upload Dataset

1. Choose File **2. Import Settings** 3. Save

Dataset was imported with the following settings:
You can change the parser options if your data was not properly imported.

Contains column header Values are separated by

DATASET PREVIEW (Imported table with **3 columns**)

activity	thrust	time in past 12 months
SQLShare Engineering	long-tail	1
SQLShare research	long-tail	1
Client+Cloud	long-tail	1
HaLoop	scalable analytics	1.5
Cloud Vis	scalable analytics	1

[Cancel](#) [Back](#) [Next](#)

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SQLSHARE

dnsamples_parsed_n... x topic.csv x

Logged in: billhowe@washington.edu

Your datasets All datasets Shared datasets Recent activity... 2 Recently viewed »

Upload dataset New query

topic.csv Only you can view this research topics simpleschema

Last modified: Feb 24, 2012 9:04 AM billhowe@washington.edu

SELECT * FROM [table_topic.csv]

Edit dataset Derive dataset Create snapshot More actions ▾

YOUR TOP VIEWED

csv2.csv	115
csv2.csv	115
blackhole	16
Vizlet Scores	14
vizlets_23nov...	14

POPULAR TAGS

biomed	138
ht_screening_r...	81
seqvalidation	52

DATASET PREVIEW Rows 1 - 20 of 20 | Columns 2 of 2

activity	topic
Astroinformatics	disc
Client+Cloud	cloud
Client+Cloud	vis
cloud certificate	cloud
Cloud Vis	cloud
Cloud Vis	disc
Cloud Vis	vis
cloud workshop	cloud
escience appliances	cloud
Graph query	db

<http://sqlshare.escience.washington.edu>

Your da

All data

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csv2.cs

csv2.csv

blackhole

Vizlet Scores

vizlets_23nov

Your datasets

All datasets

Shared datasets

Recent activity...

2

Recently viewed »

Sharing with others

Last modified: Nov 28, 2011 12:32 PM

billhowe@washington.edu

and glycerol_id parsed

```
string(d.name, 1, 5) as construct
string(d.name, 7, 5) as glycerol_id
[howe].[dnasamples.csv] d
```

[Edit dataset](#)[Derive dataset](#)[Create snapshot](#)[More actions ▾](#)[Edit dataset](#)[Derive dataset](#)[Create snapshot](#)[More actions ▾](#)

POPULAR TAGS

biomed

138

AnphA	00176	452	2	AnphA.00176.a.A1.GU26581.D1		2010-02-10 17:02:47.760147	2010-02-10 17:02:47.760147
AnphA	00202	453	1	AnphA.00202.a.B1.GE26906.D1		2010-02-10 17:02:47.760147	2010-02-10 17:02:47.760147
AnphA	00202	454	1	AnphA.00202.a.B1.GU26910.D1		2010-02-10 17:02:47.760147	2010-02-10 17:02:47.760147
AnphA	00205	455	3	AnphA.00205.a.A1.GE26620.D1		2010-02-10 17:02:47.760147	2010-02-10 17:02:47.760147

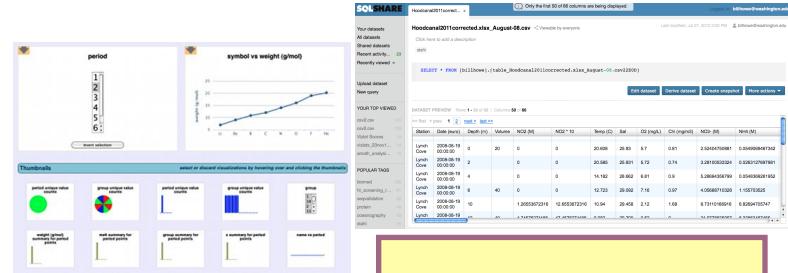
suna

16

astoda

10

<http://sqlshare.escience.washington.edu>



VizDeck

“Flagship”
SQLShare App
(Python) on EC2



Spreadsheet
Crawler

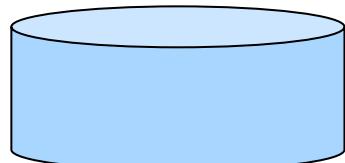
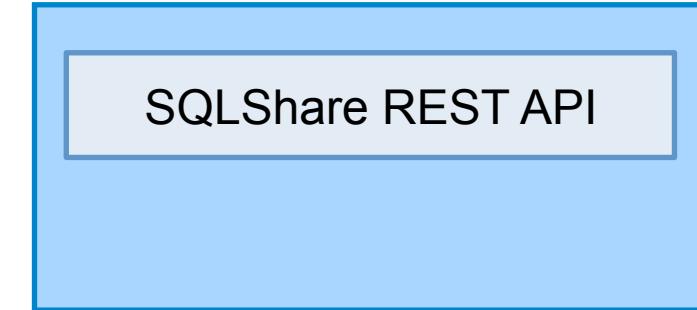
Excel
Addin

Python
Client

R
Client



Windows Azure™



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BACKUP SLIDES

Term frequency, inverse document frequency (tf-idf)

a(row_id, row_v)

row_v is a document vector

```
SELECT a1.row_id AS document_i  
, a2.row_id AS document_j,  
(a1.row_v * a2.row_v) /  
((a1.row_v * a1.row_v) * (a2.row_v * a2.row_v)) AS theta
```

```
FROM a AS a1, a AS a2  
WHERE a1.row_id > a2.row_id
```