



# Scientific Analysis by Queries in Extended SPARQL over a Scalable e-Science Data Store

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- Introduction
- SciSPARQL overview
- Evaluation
- RDF views over external storage systems
- Related approaches
- Summary

# Motivation

## Big data needs

- *scalable* data management
- good *documentation*
- *easy access*
- *reuse* of existing software packages

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- easy access
  - **standard** W3C query language for searching RDF databases: SPARQL
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  - specialized e-Science data stores
- good *documentation*
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- easy access
  - **standard** W3C query language for searching RDF databases: SPARQL
- reuse of existing software packages
  - calling **standard** and custom libraries from queries

# Problem with RDF databases

RDF (Resource Description Framework) – a W3C standard  
“metadata data model”

- RDF is very suitable for describing properties about scientific experiments (metadata) **but:**
  - Scientific data usually involves numerical arrays
  - Arrays are represented in a very inefficient way in RDF
- **Our approach:** Extend RDF with compact numerical array representation called *Numeric Multidimensional Arrays* (*NMA*)

# Problem with SPARQL queries

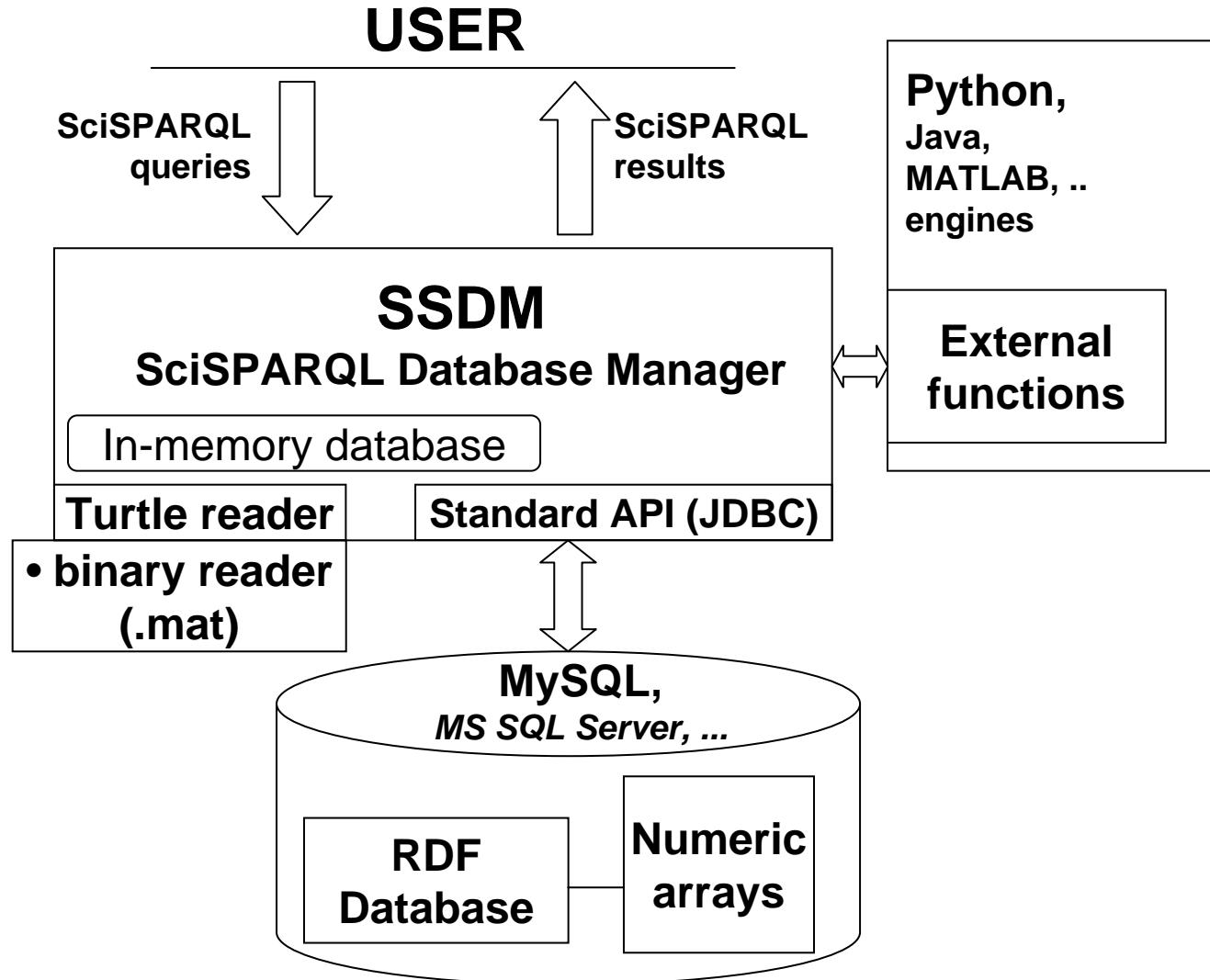
SPARQL (**S**PARQL **P**rotocol **a**nd **R**D**F** **Q**uery **L**anguage) – a W3C standard language for querying RDF

- SPARQL is very suitable for searching scientific RDF-based metadata, **but**:
  - SPARQL has no support for queries involving array operations
- **Our approach:** Extend SPARQL with common array operators => SciSPARQL

# Reusing program libraries

- Often need for using existing program libraries when processing experiments data, **but**:
  - SPARQL has no standard way of plugging in external program libraries and algorithms
- **Our approach:** SciSPARQL provides a general mechanism to call functions in C, Java, Python, or MATLAB

# Our System Architecture





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# Basic RDF experimental metadata

## EXAMPLE



# Basic RDF experimental metadata

## EXAMPLE



# Basic SPARQL metadata query

## EXAMPLE



- *RDF database of triples:*

```
prefix : <http://udbl.it.uu.se/bistab#>
```

```
:Task1 :inExperiment :Experiment1 .  
:Experiment1 :responsibleName "Andrej" .
```

# Basic SPARQL metadata query

## EXAMPLE



- *RDF database of triples:*

```
prefix : <http://udbl.it.uu.se/bistab#>
```

```
:Task1 :inExperiment :Experiment1 .  
:Experiment1 :responsibleName "Andrej" .
```

- *Select all the tasks that Andrej is responsible for*

```
SELECT ?task  
WHERE { ?task :inExperiment ?experiment .  
        ?experiment :responsibleName "Andrej" }
```

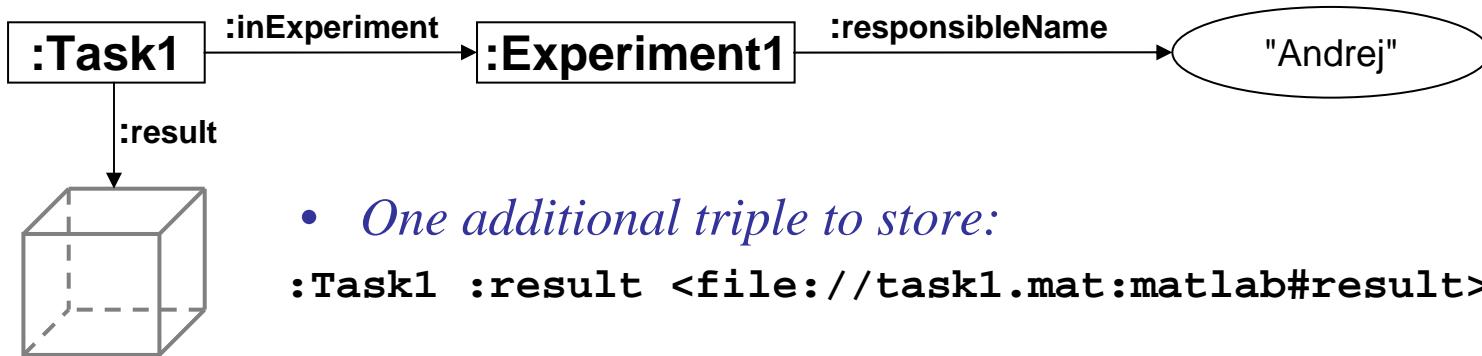
# SSDM extends RDF with arrays

## EXAMPLE



# SciSPARQL extends SPARQL with array access

## EXAMPLE

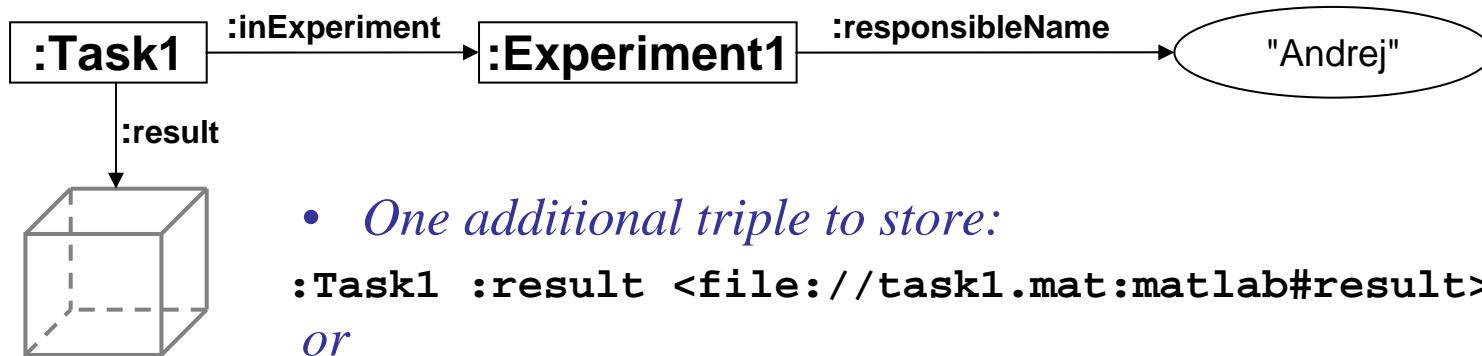


- *One additional triple to store:*

```
:Task1 :result <file://task1.mat:matlab#result> .
```

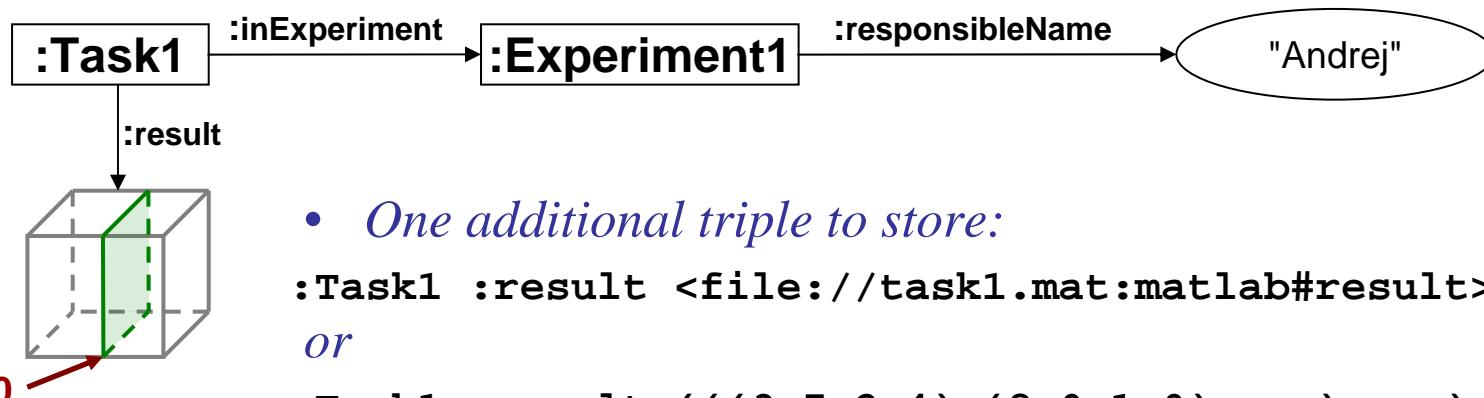
# SciSPARQL extends SPARQL with array access

## EXAMPLE



# SciSPARQL extends SPARQL with array access

## EXAMPLE



- Select 50-slice of "result" arrays  
of all tasks that Andrej is responsible for

```
SELECT (?result[50,:,:,:] AS ?slice50)  
WHERE { ?task :result ?result ;  
        :inExperiment ?experiment .  
        ?experiment :responsibleName "Andrej" }
```

# Relational Representation

## EXAMPLE

of a relational database describing a BISTAB scientific experiments

### Experiment

id	mesh	simulation algorithm	# cells	# species	specie ids				time points				
					A	B	E_A	E_B					
1	triangular #1	nsm	11107	4	0	1	2	3	<table border="1"> <tr><td>0.0</td><td>0.5</td><td>1.0</td><td>....</td></tr> </table> <p>(sequence of 201 real numbers)</p>	0.0	0.5	1.0	....
0.0	0.5	1.0	....										

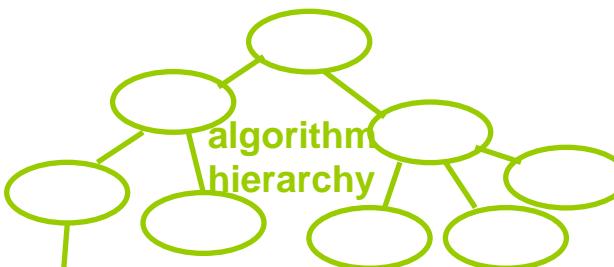
### Task

id	experiment id	parameters				realization	result
		k_1	k_a	k_d	k_4		
1	1	32.159	79.279	782750669.857	53.286	1	
2	1	19.151	39.044	300035857.676	73.445	1	

# Relational Representation

## Experiment

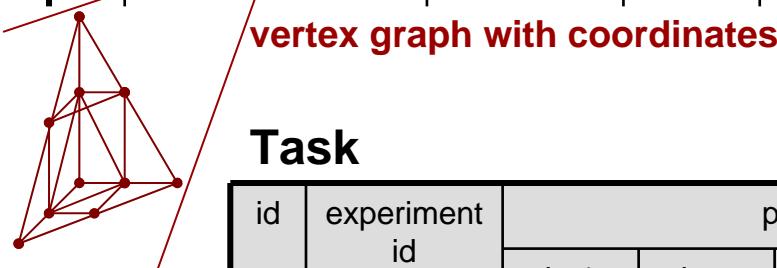
id	mesh	simulation algorithm	# cells	# species	specie ids				time points
					A	B	E_A	E_B	
1	triangular #1	nsm	11107	4	0	1	2	3	[0.0, 0.5, 1.0, ...]



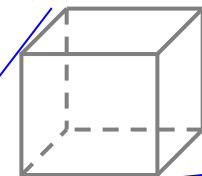
more types of species

## Task

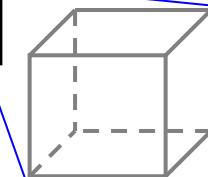
id	experiment id	parameters				realization	result
		k_1	k_a	k_d	k_4		
1	1	32.159	79.279	782750669.857	53.286	1	
2	1	19.151	39.044	300035857.676	73.445	1	



more parameters



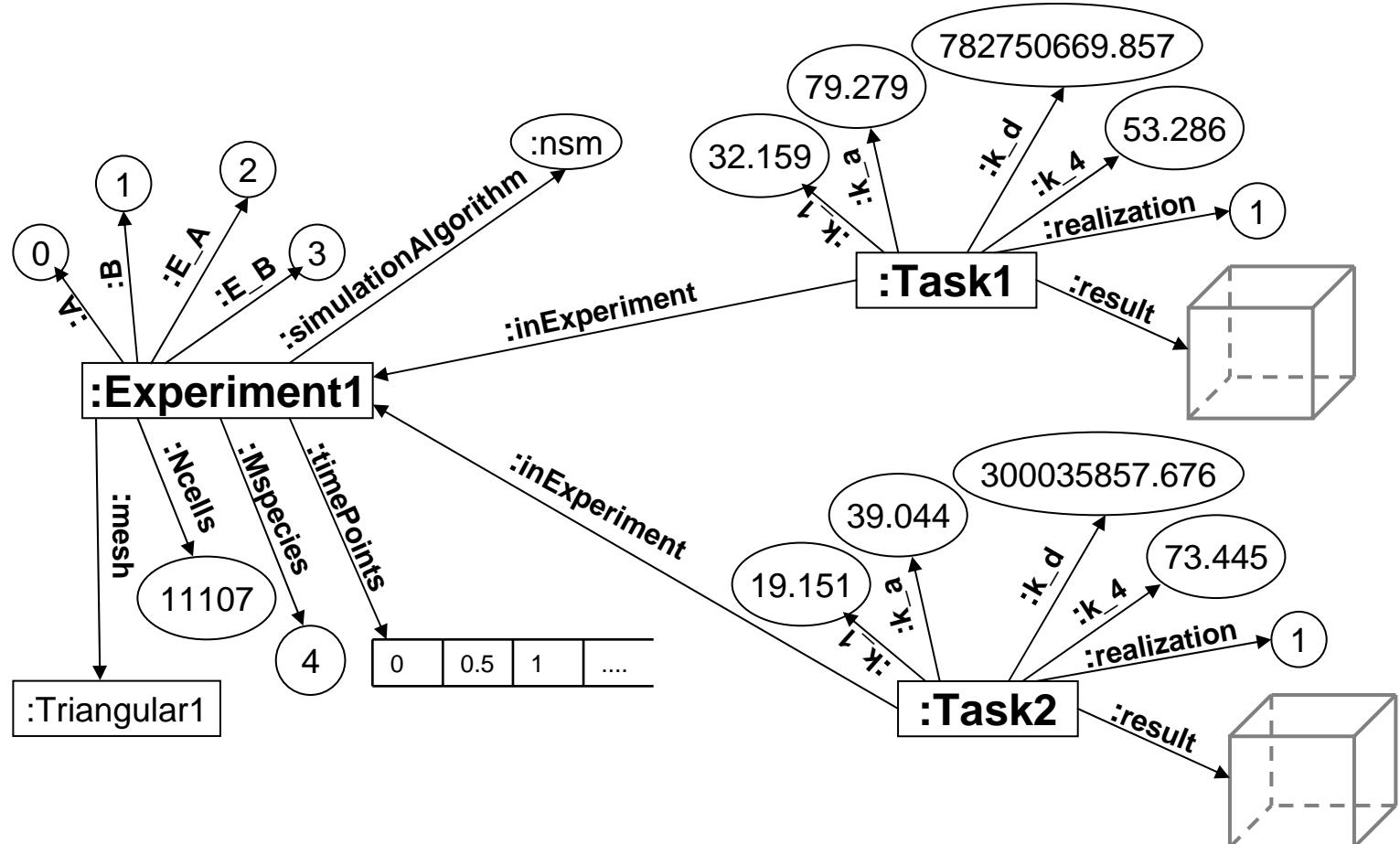
(array of 11107 x 4 x 201 integers)



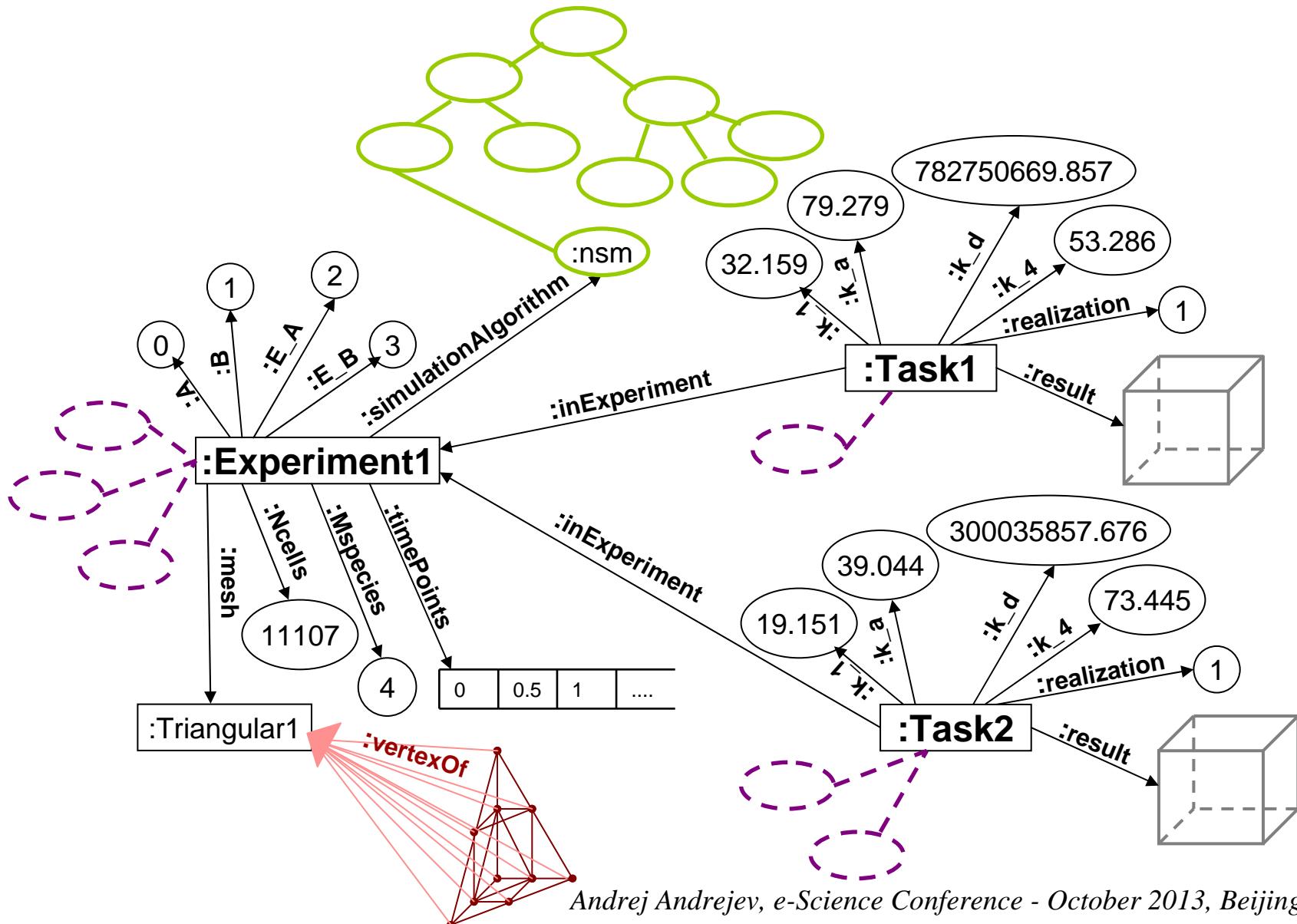
# SSDM Representation: RDF with Arrays

## EXAMPLE

of an RDF database describing BISTAB scientific experiments



# SSDM Representation: RDF with Arrays

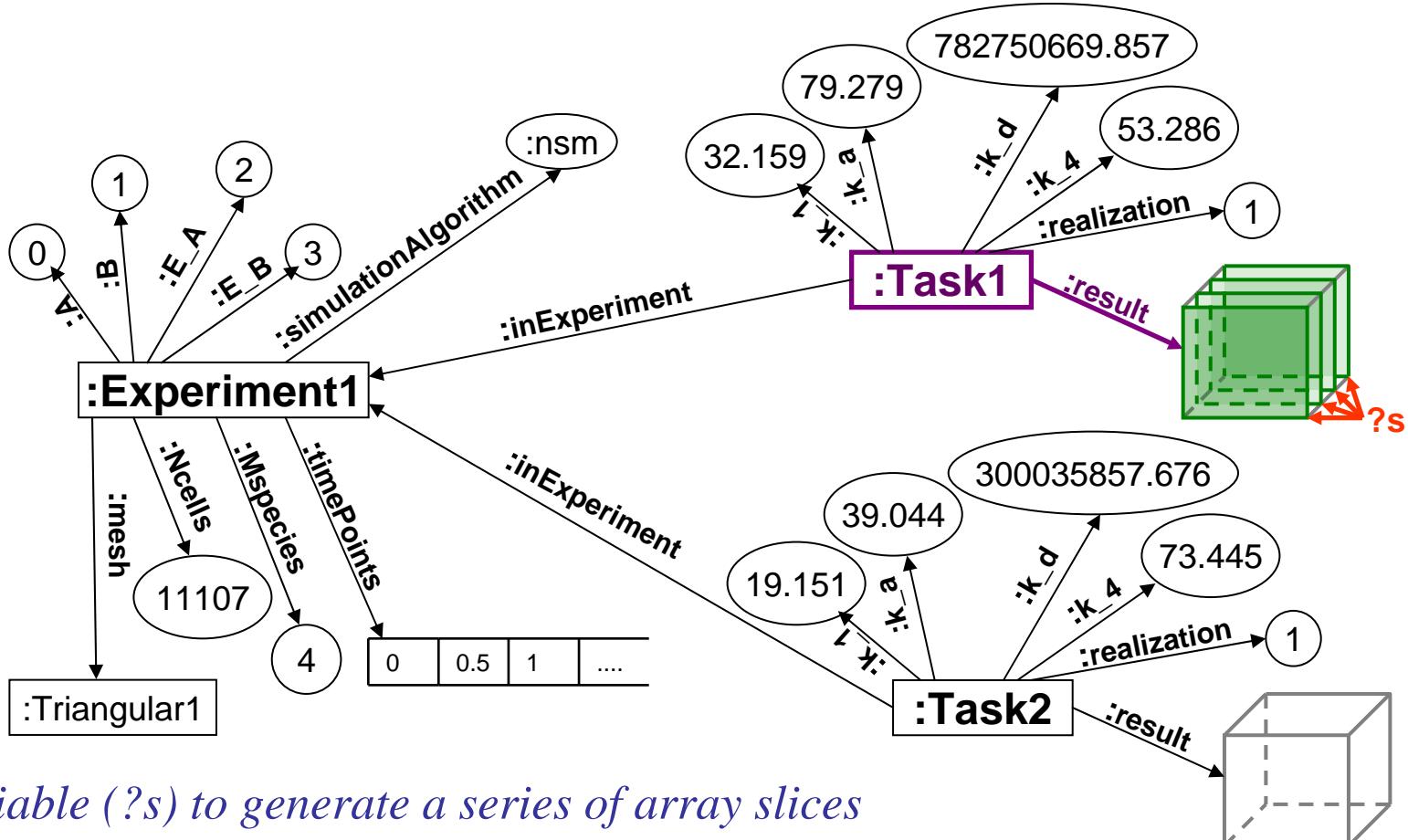




# More Scientific SPARQL examples

# SciSPARQL Query Language

```
SELECT (AVG(?result[:, :, ?s]) AS ?specAvarage)
WHERE { :Task1 :result ?result }
```

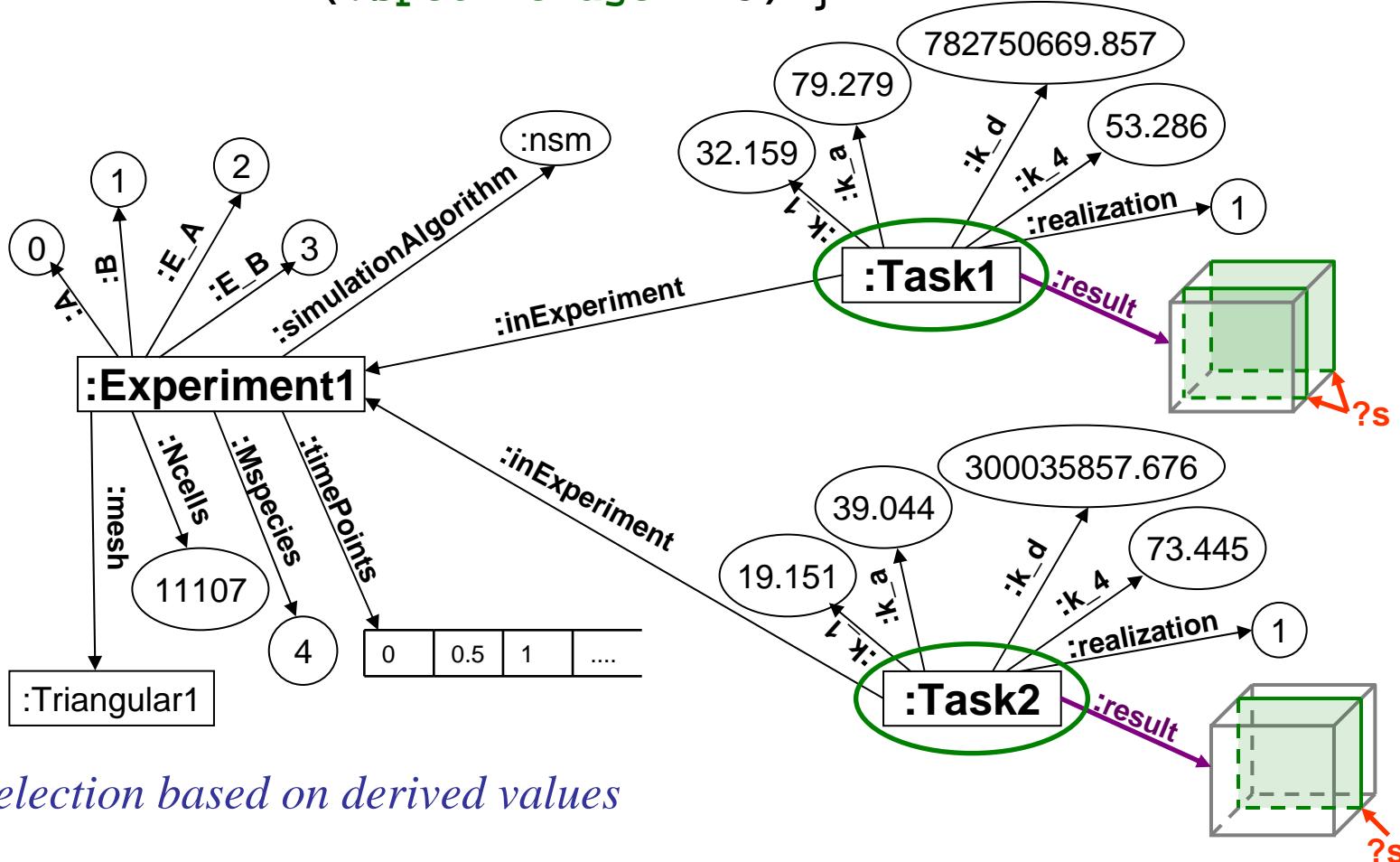


- Use free variable (?s) to generate a series of array slices

# SciSPARQL Query Language

```

SELECT ?task ?s ?specAverage
WHERE { ?task :result ?result .
        BIND ( AVG(?result[::,:,:,?s]) AS ?specAvarage ) .
        FILTER (?specAverage > 5) }
    
```



- Filter data selection based on derived values



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# Our Contribution

**SSDM shows performance on par with MATLAB,  
with added value of**

## MATLAB

Programs  
implementing analysis algorithms

No metadata management  
user manually manages files

## SciSPARQL

High-level queries

Uniform management of  
both data and metadata

# Our Contribution

**SSDM shows performance on par with MATLAB,  
with added value of**

MATLAB

```
sum_of_A = [];
load('input.mat'); % parameters, tspan 'metadata'
t = find(tspan==10);
a = 1; % this 'metadata' is not stored anywhere
mspecies = 8;
for ii=1:100 % amount of files should be known!
    if parameters(1,ii) >= 50
        && parameters(1,ii) <= 90
        && parameters(3,ii) >= 1.0E8
        && parameters(3,ii) <= 1.0E9
    realization = strcat( % construct filenames
        'C:/DATA/bistab2f/realization_',
        int2str(ii),'_1.mat');
    load(realization); % load matrices 1-by-1
    sum_of_A = [sum_of_A
                sum(UU(a:mspecies:end, t))];
end
sum_of_A;
```

SciSPARQL Q2

```
SELECT (array_sum(?U[?a-1::?mspecies,?j]) AS ?res)
WHERE { ?task :U ?U ; # retrieve data
        :k_a ?k_a ; # retrieve metadata
        :k_d ?k_d ;
        :inExperiment ?experiment .
?experiment :A ?a ;
            :MSpecies ?mspecies ;
            :tspan ?tspan .
FILTER (?tspan[?j] = 10 &&
       1.0E8 <= ?k_d && ?k_d <= 1.0E9 &&
       50 <= ?k_a && ?k_a <= 90 ) };
```



# Our Contribution

SSPM shows performance on par with MATLAB,

```
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        load(realization); % load matrices 1-by-1
        sum_of_A = [sum_of_A
                    sum(UU(a:mspecies:end, t))];
    end
end
sum_of_A;
```

?res)

ta

0E9 &&  
};



# Our Contribution

SSDM shows performance on par with MATLAB,  
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```
SELECT (array_sum(?U[?a-1::?mspecies,?j]) AS ?res)
WHERE { ?task :U ?U ; # retrieve data
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         :k_d ?k_d ;
         :inExperiment ?experiment .
?experiment :A ?a ;
             :MSpecies ?mspecies ;
             :tspan ?tspan .
FILTER (?tspan[?j] = 10 &&
        1.0E8 <= ?k_d && ?k_d <= 1.0E9 &&
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data
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sum_of_A =
load('input
t = find(ts
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  && p
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reali
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sum_of_A = [sum_of_A
            sum(UU(a:mspecies:end, t))];
end
end
sum_of_A;
```

# SSDM Performance

7GB database, query execution times (in seconds) with all data on disk

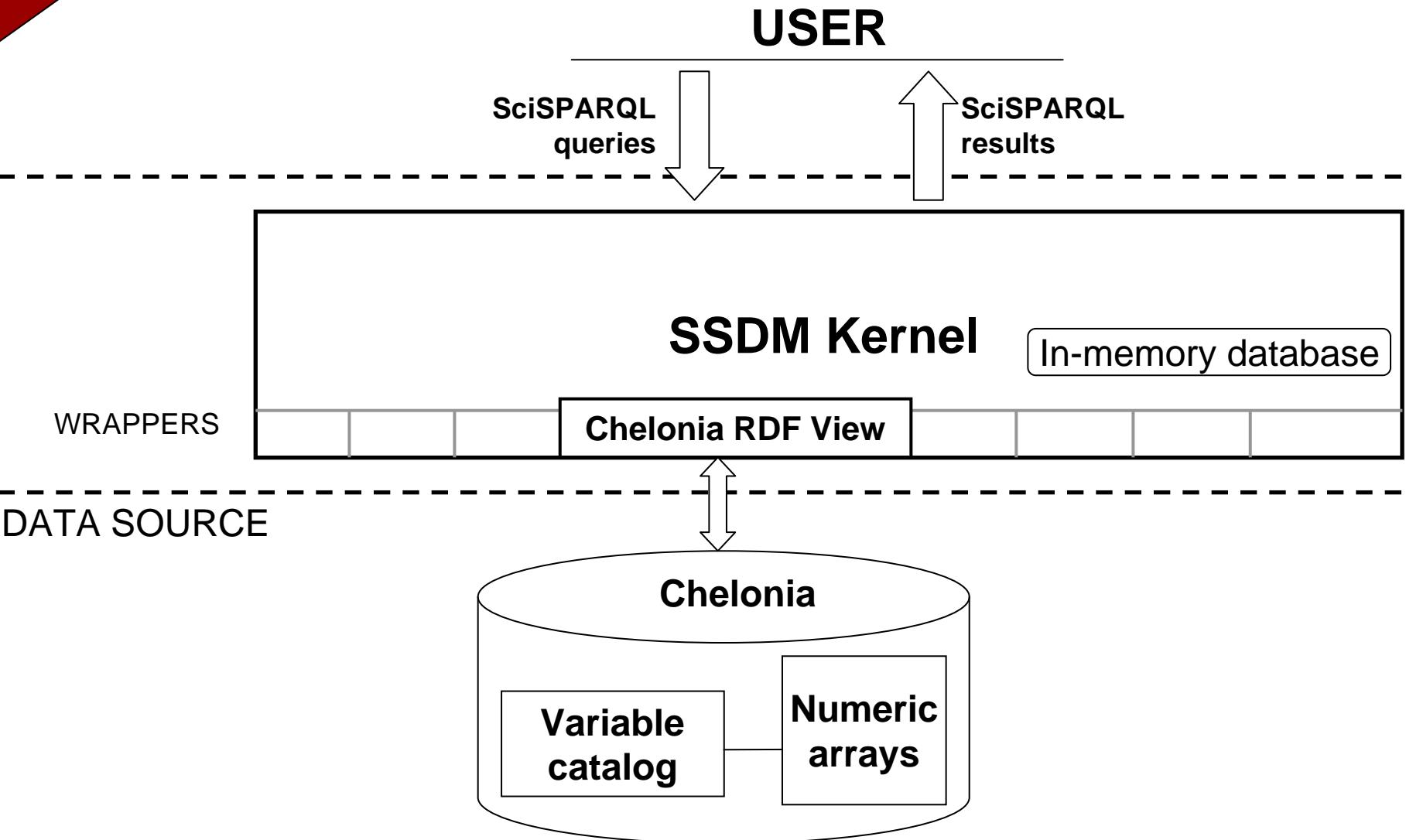
Task	Data retrieved	SSDM with back-end		MATLAB script
		MySQL	MS SQL Server	
<b>Q1: (selective query)</b> Compute an aggregate value over 1 big matrix, every 8th row	<b>18MB</b>	<b>1.748</b>	<b>2.15</b>	<b>1.826</b>
<b>Q2: (SSDM worst case)</b> Select 36 matrices, access one column × every 8th row	<b>642MB</b>	<b>80.703</b>	<b>44.512</b>	<b>30.042</b>
<b>Q3: (database scan)</b> Compute AGRMAX of Q1 across all matrices, 25% rows	<b>1785MB</b>	<b>187.073</b>	<b>192.365</b>	<b>133.279</b>

=> SSDM provides desired functionality with competitive performance

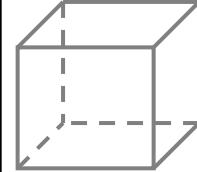
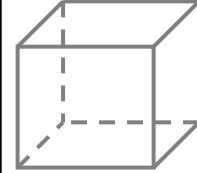


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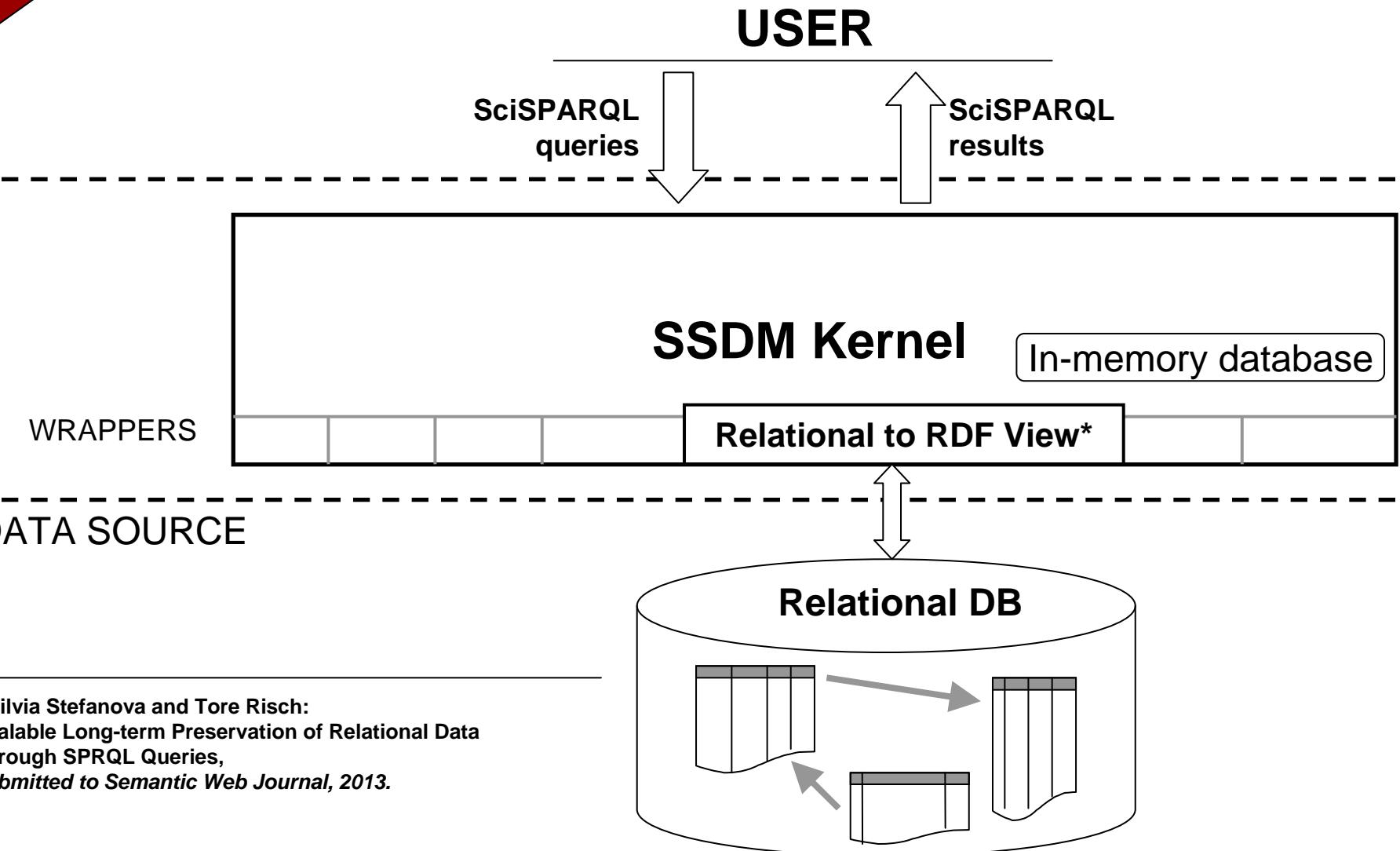
# RDF views over external storage systems



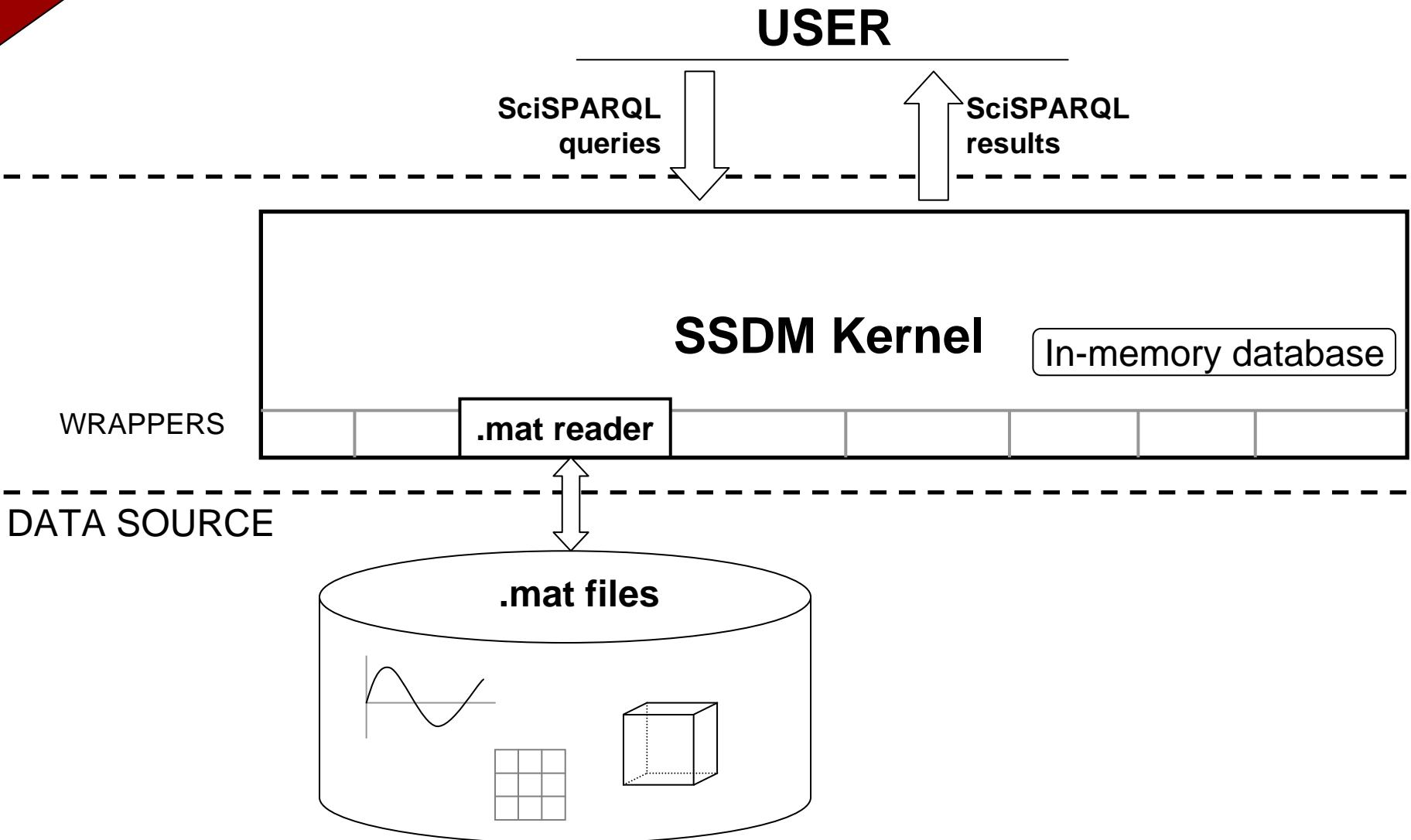
# Chelonia Native Schema

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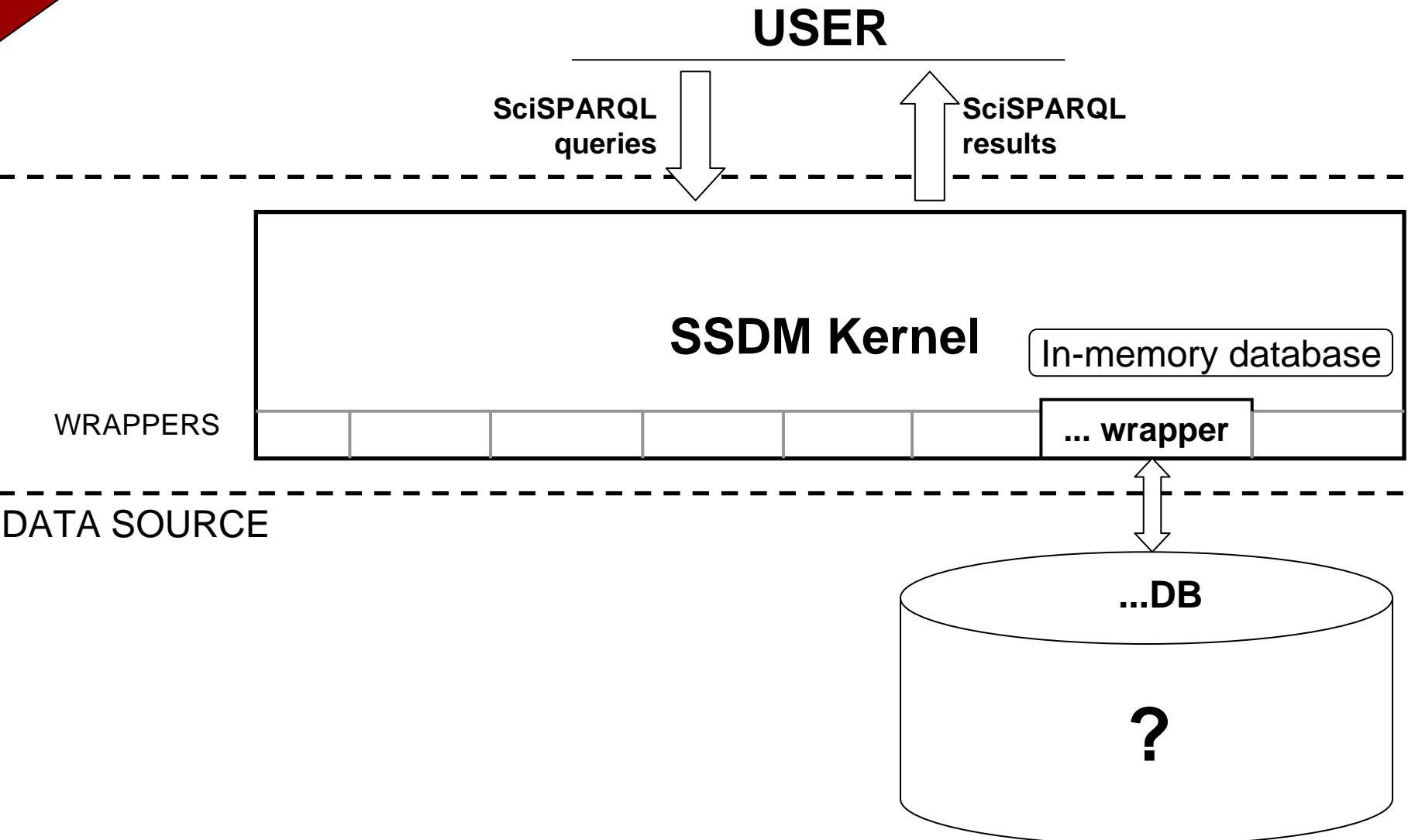
# RDF views over external storage systems



# RDF views over external storage systems



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# Related approaches

## Databases

- High-level metadata descriptions (schemas)
- Scalable data representation
- High-level query languages

## RDF

- Designed for metadata in general
- Voluntary schema
- Weak support numeric applications

## Files and programs

- No explicit metadata
- Many storage formats and APIs
- Numerical libraries maintained since 1960:s
- Extensively used in scientific computing

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- No explicit metadata
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## SSDM and SciSPARQL

- Full database support
- Flexibility of RDF
- Reuse of existing libraries

# Related query languages

## Extending RDBMS with array semantics

- AQuery [Lerner & Shasha, 2003]
- SciQL [Kersten et.al, 2011]
- Storing arrays as BLOBs
  - RasQL [Furtado & Baumann, 1999]
  - UDFs in T-SQL [Dobos et.al., 2011]
- Specialized array databases and languages
  - AQL [Libkin et.al. 1996]
  - SciDB [Cudre-Mauroux et.al. 2009]
- Foreign Functions in SPARQL
  - SESAME
  - CORESE



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# Summary

## **SSDM (SciSPARQL Database Manager) provides**

- Efficient storage of RDF with arrays
- Back-end relational database storage and various data file formats
- Access to external databases

## **SciSPARQL provides**

- support of numeric multidimensional arrays and operations
- extensibility with foreign functions in C, Java, Python, and MATLAB



**The software, documentation, and examples  
are available at**

**<http://www.it.uu.se/research/group/udbl/SciSPARQL>**

**This work was supported by**

