

# Integration architecture

- Queryable sources that generate XML
- Distributed queries over sources
- Query Manager to manage queries
- Saved queries as sources that can also be queried
- Leads to potential for query networks

vSPARQL  
Service



# VSPARQL Demo

## Enter SparQL Query:

```
<pre>?structure fma:talairach ?structure_name .
?lateralized_structure fma:talairach ?lateralized_structure_name .
?part fma:talairach ?part_name .

FROM <http://sig.uw.edu/neurofma_v1.3>
WHERE
{
  ?structure rdf:type fma:BrainLateralized_structure .
  OPTIONAL { ?structure fma:BrainLateralized_structure fma:talairach ?lateralized_structure_name . }
  ?lateralized_structure rdf:type fma:BrainLateralized_structure .
  OPTIONAL { ?lateralized_structure fma:talairach ?l2 - ?l2 fma:talairach ?lateralized_structure_name . }
  ?lateralized_structure fma:part ?part .
  OPTIONAL { ?part fma:talairach ?l1 - ?l1 fma:talairach ?part_name . }
}
```

Submit Query

## Sample Queries:

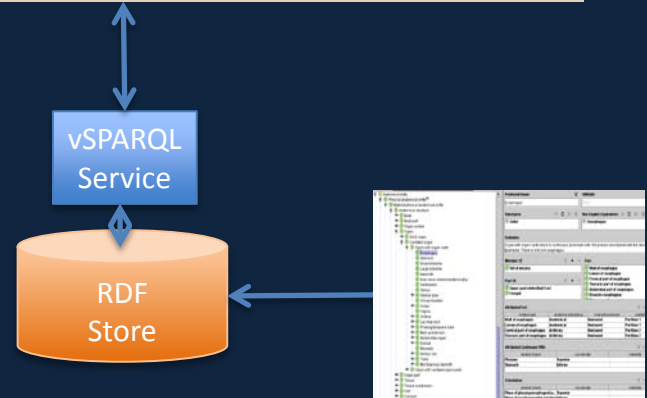
FMA Queries | NCI Thesaurus Queries | VSPARQL Queries

Sample Query 1: (select everything that is directly continuous with the Esophagus)

```
<pre>PREFIX fma:<http://sig.biostr.washington.edu/fma3.0#>
SELECT ?esoph_continuity
FROM <http://sig.biostr.washington.edu/fma3.0>
WHERE
{
  fma:Esophagus fma:continuity_with ?esoph_continuity .
}
```

Sample Query 2: (construct the regional part tree for Heart with PMIDs)

```
<pre>PREFIX fma:<http://sig.biostr.washington.edu/fma3.0#>
CONSTRUCT
{
  fma:Heart fma:regional_part ?object .
  fma:Heart fma:PMID ?heart_id .
}
```



```
CONSTRUCT {
?structure fma:Talairach ?structure_name .
?lateralized_structure fma:Talairach ?lateralized_structure_name .
?part fma:Talairach ?part_name}

FROM <http://sig.uw.edu/neurofma_v1.3>
WHERE
{
?structure apf:assign fma:Dorsolateral_prefrontal_cortex .
OPTIONAL {?structure fma:Talairach ?t1 . ?t1 fma:name ?structure_name }.
?lateralized_structure rdfs:subClassOf ?structure .
OPTIONAL {?lateralized_structure fma:Talairach ?t2 . ?t2 fma:name ?lateralized_structure_name } .
?lateralized_structure gleen:OnPath ( '([fma:regional_part]|[fma:constitutional_part])*' ?part ) .
OPTIONAL {?part fma:Talairach ?t3 . ?t3 fma:name ?part_name }

}
```

```
2-rdf-syntax-ns#"
ofunction.library."
n.edu/tmp#"
leen."
w1#"
rdf-schema#"

w.edu/fma#Brodmann_area_46_of_right_middle_frontal_gyrus">
w.w3.org/2001/XMLSchema#string">Right Cerebrum.Frontal Lobe.Middle Frontal Gyrus.Gray Matter.Brodmann area 46</f

w.edu/fma#Brodmann_area_46_of_left_middle_frontal_gyrus">
w.w3.org/2001/XMLSchema#string">Left Cerebrum.Frontal Lobe.Middle Frontal Gyrus.Gray Matter.Brodmann area 46</f

w.edu/fma#Brodmann_area_9_of_left_middle_frontal_gyrus">
w.w3.org/2001/XMLSchema#string">Left Cerebrum.Frontal Lobe.Middle Frontal Gyrus.Gray Matter.Brodmann area 9</f

w.edu/fma#Brodmann_area_9_of_left_superior_frontal_gyrus">
w.w3.org/2001/XMLSchema#string">Left Cerebrum.Frontal Lobe.Superior Frontal Gyrus.Gray Matter.Brodmann area 9</f

w.edu/fma#Brodmann_area_9_of_right_superior_frontal_gyrus">
w.w3.org/2001/XMLSchema#string">Right Cerebrum.Frontal Lobe.Superior Frontal Gyrus.Gray Matter.Brodmann area 9</f

w.edu/fma#Brodmann_area_9_of_right_middle_frontal_gyrus">
w.w3.org/2001/XMLSchema#string">Right Cerebrum.Frontal Lobe.Middle Frontal Gyrus.Gray Matter.Brodmann area 9</f
```

# VSPARQL Demo

## Enter SparQL Query:

```
<pre>?structure fma:talairach ?structure_name .
?lateralized_structure fma:talairach ?lateralized_structure_name .
?part fma:talairach ?part_name .

FROM <http://sig.wv.edu/neurofma_v1.3>
WHERE
{
  ?structure rdf:type fma:Dorsolateral_gro[frontal_cortex .
  OPTIONAL (??structure fma:talairach ?l - ?l fma:same ?structure_name ) .
  ?lateralized_structure rdf:type fma:O ?structure .
  OPTIONAL (??lateralized_structure fma:talairach ?l2 - ?l2 fma:same ?lateralized_structure_name ) .
  ?lateralized_structure rdfs:subClassOf ( (fma:regional_part || fma:constititucional_part) ?part ) .
  OPTIONAL (?part fma:talairach ?l3 - ?l3 fma:same ?part_name )
}
</pre>
```

Submit Query

## Sample Queries:

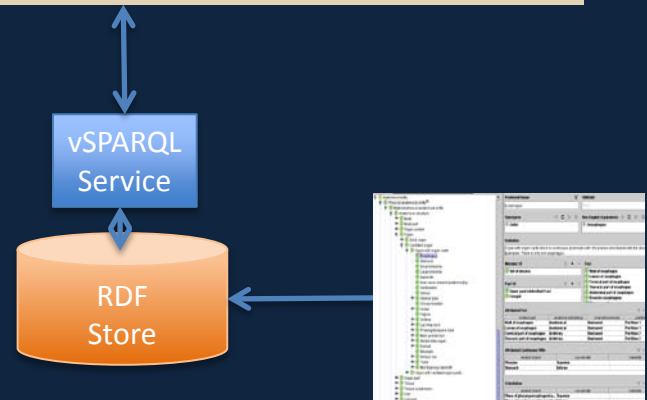
FMA Queries | NCI Thesaurus Queries | VSPARQL Queries

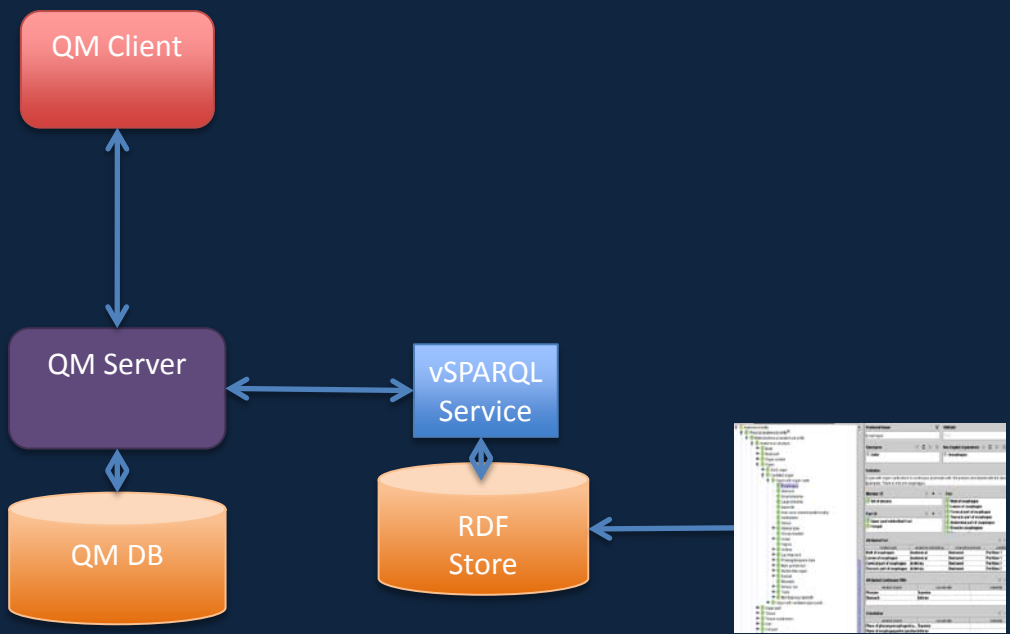
Sample Query 1: (select everything that is directly continuous with the Esophagus)

```
<pre>PREFIX fma:<http://sig.biostr.washington.edu/fma3.0#>
SELECT ?esoph_continuity
FROM <http://sig.biostr.washington.edu/fma3.0>
WHERE
{
  fma:Esophagus fma:continuity_with ?esoph_continuity .
}
</pre>
```

Sample Query 2: (construct the regional part tree for Heart with PMIDs)

```
<pre>PREFIX fma:<http://sig.biostr.washington.edu/fma3.0#>
CONSTRUCT
{
  fma:Heart fma:regional_part ?object .
  fma:Heart fma:PMID ?heart_id .
}
</pre>
```







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## Title

74: AMIA 2010: Talairach: Regional or constitutional parts of a structure that has lateralized subclasses: Dorsolateral\_prefrontal\_cortex

## Description

First find direct subclasses, then transitive regional parts, output those that have Talairach labels.  
To change structure substitute different structure name for Dorsolateral\_prefrontal\_cortex

## Query

```
PREFIX rdfs: <http://www.w3.org/2000/01/rdf-schema#>
PREFIX rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#>
PREFIX owl: <http://www.w3.org/2002/07/owl#>
PREFIX gleen: <java:edu.washington.sig.gleen.>
PREFIX fma: <http://sig.uw.edu/fma#>
PREFIX tmp: <http://sig.biostr.washington.edu/tmp#>
PREFIX apf: <java:com.hp.hpl.jena.query.pfunction.library.>

CONSTRUCT {
?structure fma: Talairach ?structure_name .
?lateralized_structure fma: Talairach ?lateralized_structure_name .
?part fma: Talairach ?part_name }

FROM <http://sig.uw.edu/neurofma_v1.3>
WHERE
{
?structure apf: assign fma: Dorsolateral_prefrontal_cortex .
OPTIONAL { ?structure fma: Talairach ?t1 . ?t1 fma: name ?structure_name } .
?lateralized_structure rdfs: subclassOf ?structure .
OPTIONAL { ?lateralized_structure fma: Talairach ?t2 . ?t2 fma: name ?lateralized_structure_name } .
?lateralized_structure gleen: OnPath ( '([fma: regional_part]|[fma: constitutional_part])*' ?part ) .
OPTIONAL { ?part fma: Talairach ?t3 . ?t3 fma: name ?part_name }

}
```

Language **VSparQL**

[Execute](#)

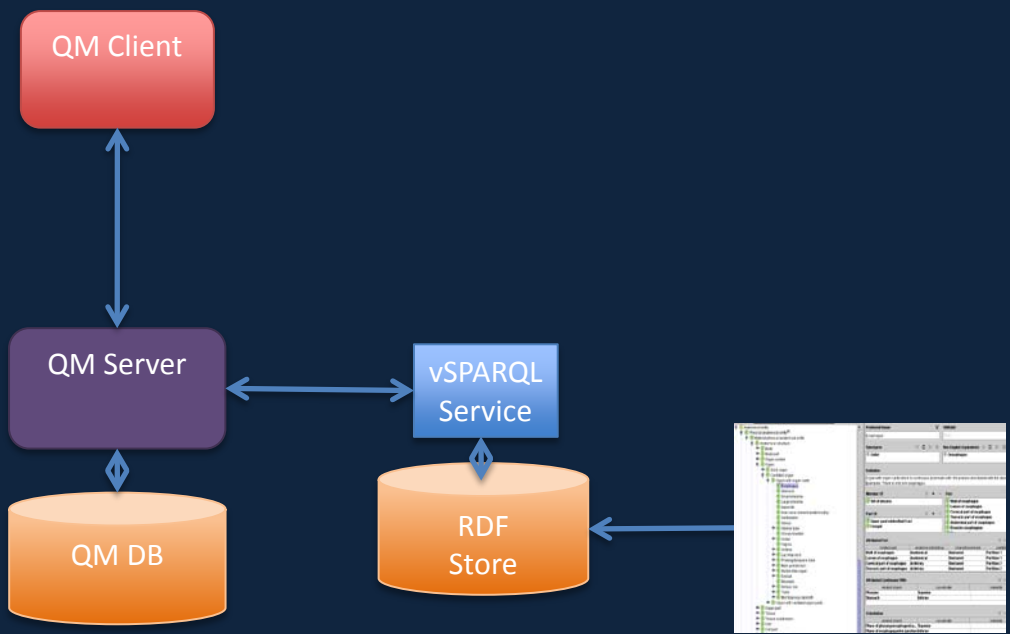


## Query Results



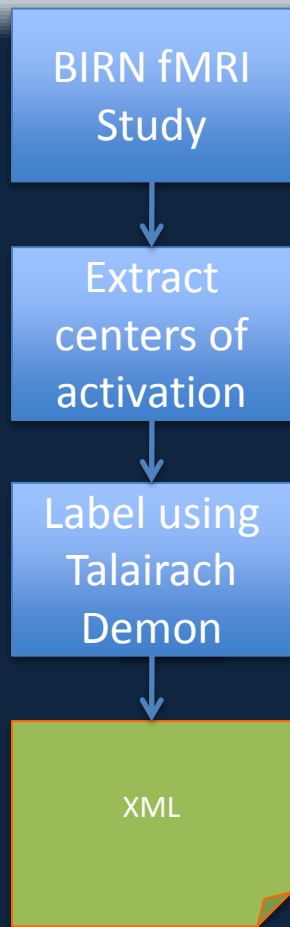
```
<rdf:RDF
  xmlns:rdf="http://www.w3.org/1999/02/22-rdf-syntax-ns#"
  xmlns:apf="java:com.hp.hpl.jena.query.pfunction.library."
  xmlns:tmp="http://sig.biostr.washington.edu/tmp#"
  xmlns:gleen="java:edu.washington.sig.gleen."
  xmlns:owl="http://www.w3.org/2002/07/owl#"
  xmlns:rdfs="http://www.w3.org/2000/01/rdf-schema#"
  xmlns:fma="http://sig.uw.edu/fma#" >
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_46_of_right_middle_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Right Cerebrum.Frontal Lobe.Middle Frontal Gyrus.Gray Matter
  </rdf:Description>
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_46_of_left_middle_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Left Cerebrum.Frontal Lobe.Middle Frontal Gyrus.Gray Matter.
  </rdf:Description>
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_9_of_left_middle_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Left Cerebrum.Frontal Lobe.Middle Frontal Gyrus.Gray Matter.
  </rdf:Description>
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_9_of_left_superior_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Left Cerebrum.Frontal Lobe.Superior Frontal Gyrus.Gray Matte
  </rdf:Description>
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_9_of_right_superior_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Right Cerebrum.Frontal Lobe.Superior Frontal Gyrus.Gray Matt
  </rdf:Description>
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_9_of_right_middle_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Right Cerebrum.Frontal Lobe.Middle Frontal Gyrus.Gray Matter
  </rdf:Description>
</rdf:RDF>
```





# BIRN fMRI Study

Schizophrenics vs Healthy  
Controls



```
- <results>
  - <site>
    <site_label>301882920_11.4</site_label>
    <BIRN_ID>301882920</BIRN_ID>
    <Diagnostic_group>h</Diagnostic_group>
    <Voxels>475</Voxels>
    <Max_Z>11.4</Max_Z>
    <P>5.96E-8</P>
    <log10P>7.22</log10P>
    <Distance>4</Distance>
  - <Tal>
    Right Cerebrum.Temporal Lobe.Middle Temporal Gyrus.Gray Matter.Brodmann area 22
  </Tal>
  <right_coord>52</right_coord>
  <sup_coord>0</sup_coord>
  <ant_coord>-44</ant_coord>
</site>
- <site>
  <site_label>301882920_11.3</site_label>
  <BIRN_ID>301882920</BIRN_ID>
  <Diagnostic_group>h</Diagnostic_group>
  <Voxels>7287</Voxels>
  <Max_Z>11.3</Max_Z>
  <P>0</P>
  <log10P>54.8</log10P>
  <Distance>11</Distance>
- <Tal>
  Right Cerebrum.Frontal Lobe.Precentral Gyrus.Gray Matter.Brodmann area 6
</Tal>
<right_coord>44</right_coord>
<sup_coord>26</sup_coord>
<ant_coord>0</ant_coord>
</site>
```



XML

XML

QM Client

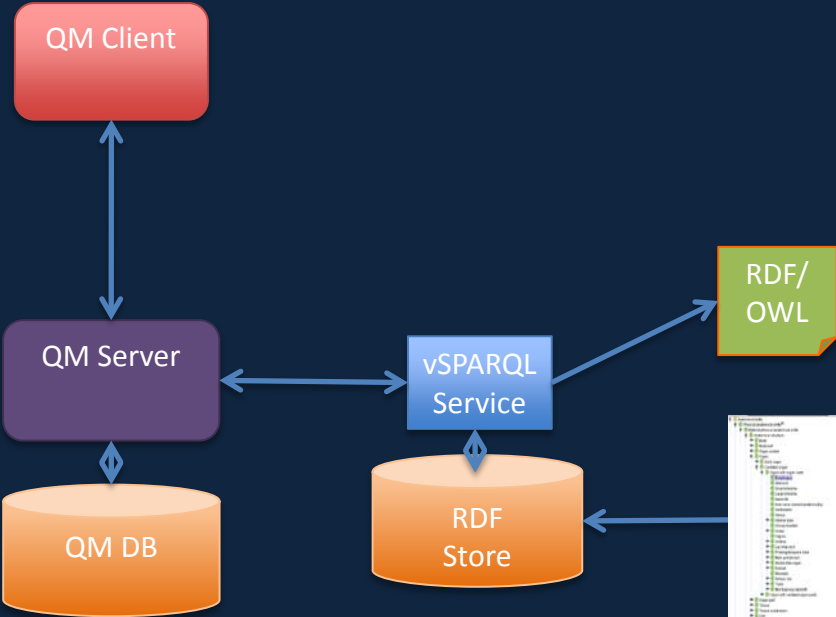
QM Server

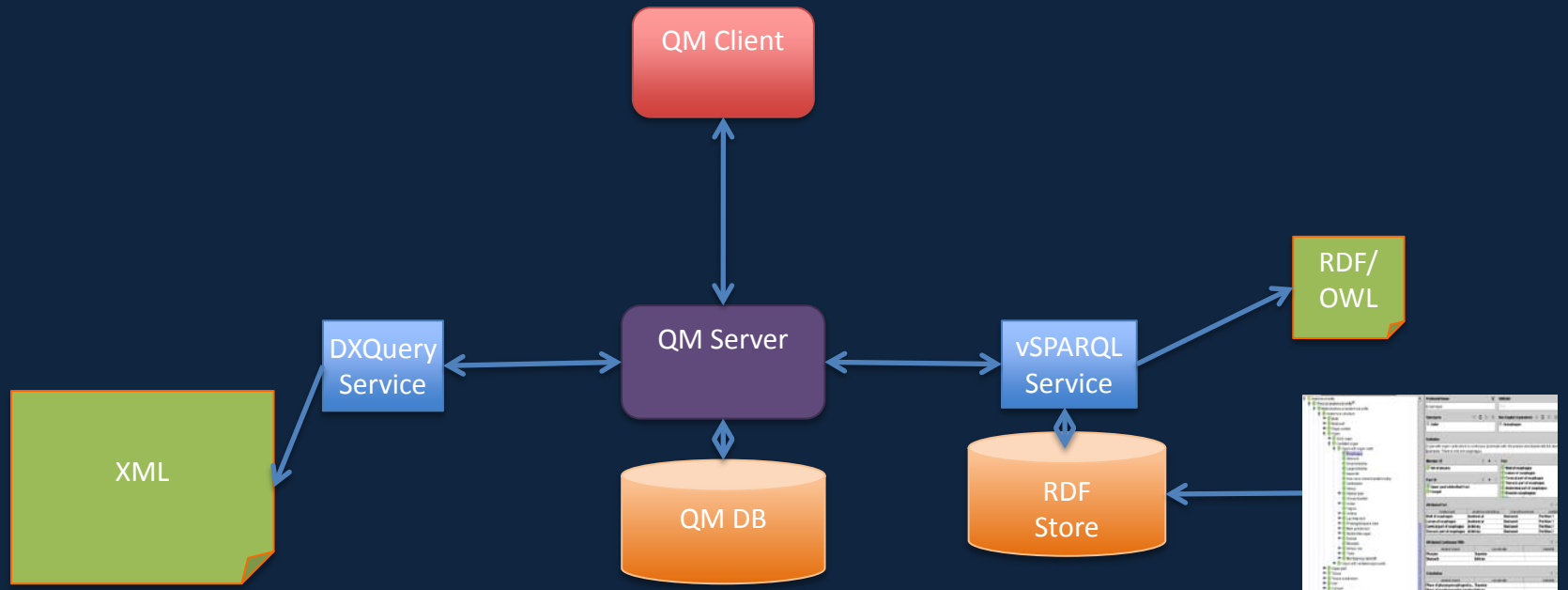
QM DB

vSPARQL Service

RDF Store

RDF/OWL







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## Title

Group JessMerged into Sz and Healthy

## Description

Create two lists

## Query

```
let $act_doc := doc("http://synapse.biostr.washington.edu/~brinkley/dxdemos/JessMerged.xml")
```

```
(: now filter by healthy vs sz :)
```

```
let $healthy_act := $act_doc/site[Diagnostic_group/text()='h']
```

```
let $schizo_act := $act_doc/site[Diagnostic_group/text()='s']
```

```
(: add color for display of both groups :)
```

```
let $h_color := 'Red'
```

```
let $s_color := 'Yellow'
```

```
let $shape := 'BIG_SPHERE'
```

```
let $size := '1.75'
```

```
let $healthy_sites :=
```

```
  for $el in $healthy_act
```

```
  return
```

```
    <site>
```

```
      {$el/site_label}
```

```
      {$el/BIRN_ID}
```

```
      {$el/Diagnostic_group}
```

```
      {$el/Voxels}
```

```
      {$el/Max_Z}
```

```
      {$el/P}
```

```
      {$el/log10P}
```

```
      {$el/Distance}
```

```
      {$el/Tal}
```

```
      {$el/right_coord}
```

```
      {$el/sup_coord}
```

```
      {$el/ant_coord}
```

```
      <Color>{$h_color}</Color>
```

Language

**DXQuery**

**Execute**



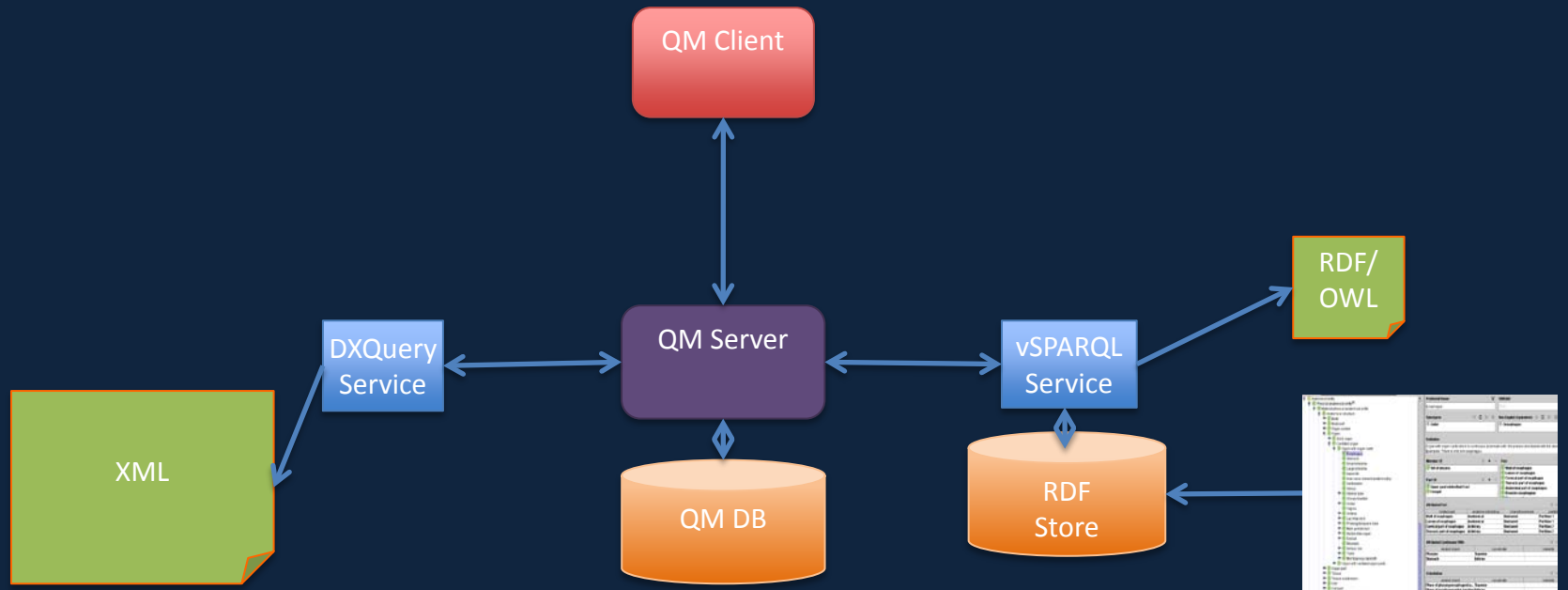
## Query Results

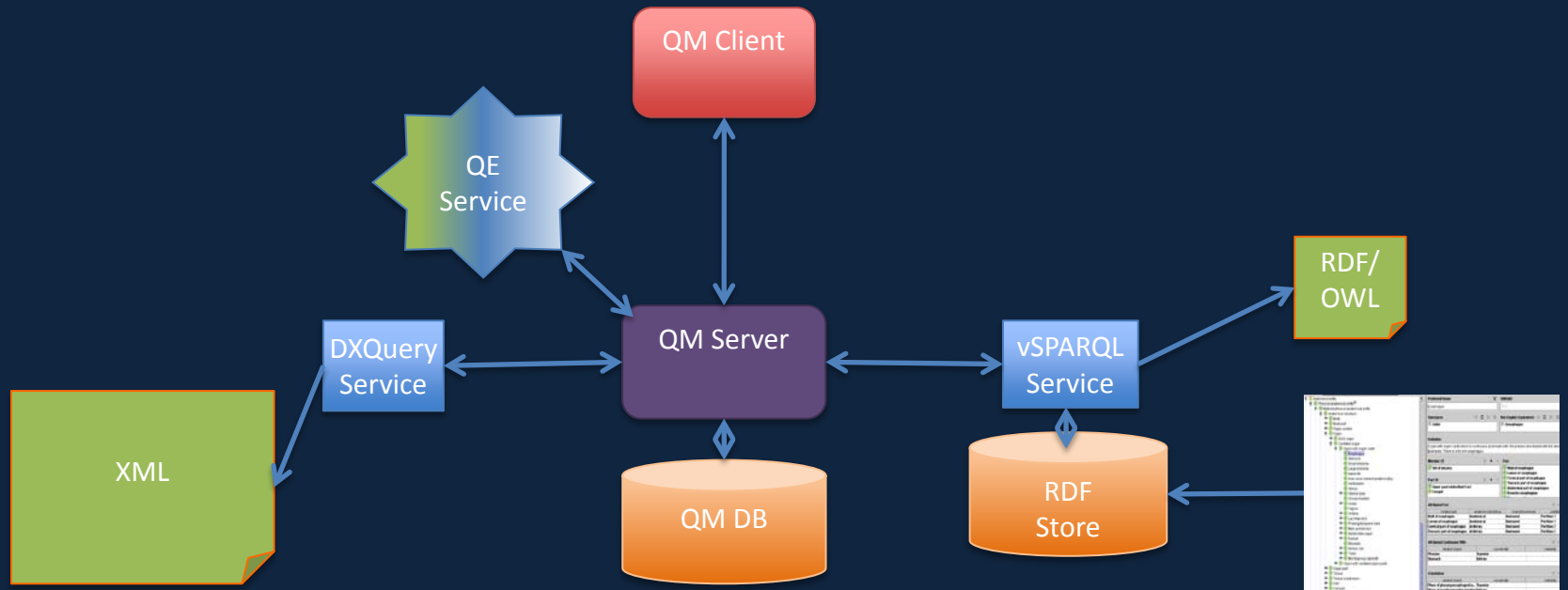


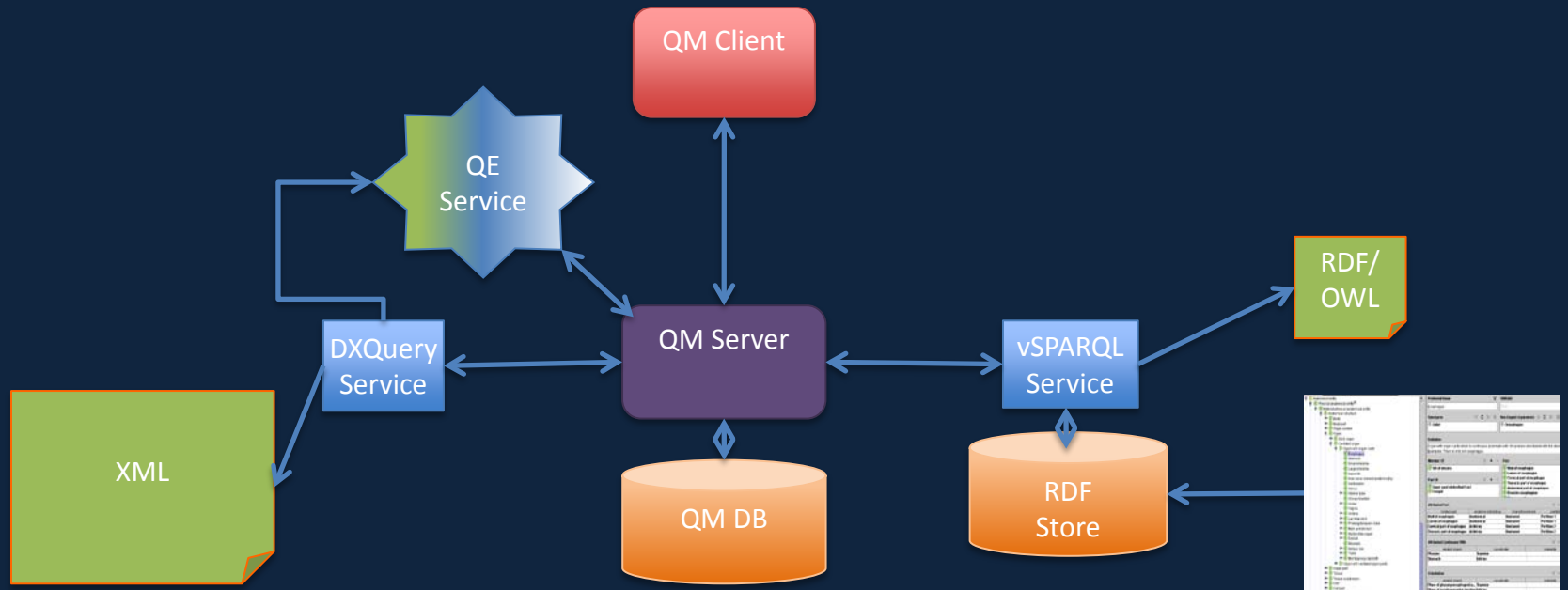
```
<?xml version="1.0" encoding="UTF-8"?>
<results>
  <patient>
    <pnum>MNI</pnum>
    <site>
      <site_label>301882920_11.4</site_label>
      <BIRN_ID xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">301882920</BIRN_ID>
      <Diagnostic_group xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">h</Diagnostic_group>
      <Voxels xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">475</Voxels>
      <Max_Z xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">11.4</Max_Z>
      <P>5.96E-8</P>
      <log10P>7.22</log10P>
      <Distance>4</Distance>
      <Tal> Right Cerebrum.Temporal Lobe.Middle Temporal Gyrus.Gray Matter.Brodmann area 22</Tal>
      <right_coord>52</right_coord>
      <sup_coord>0</sup_coord>
      <ant_coord>-44</ant_coord>
      <Color>Red</Color>
      <Shape>BIG_SPHERE</Shape>
      <Size>1.75</Size>
    </site>
    <site>
      <site_label>301882920_11.3</site_label>
      <BIRN_ID xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">301882920</BIRN_ID>
      <Diagnostic_group xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">h</Diagnostic_group>
      <Voxels xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">7287</Voxels>
      <Max_Z xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">11.3</Max_Z>
      <P>0</P>
      <log10P>54.8</log10P>
      <Distance>11</Distance>
      <Tal> Right Cerebrum.Frontal Lobe.Precentral Gyrus.Gray Matter.Brodmann area 6</Tal>
      <right_coord>44</right_coord>
      <sup_coord>26</sup_coord>
      <ant_coord>0</ant_coord>
      <Color>Red</Color>
      <Shape>BIG_SPHERE</Shape>
      <Size>1.75</Size>
    </site>
    <site>
      <site_label>301882920_10.1</site_label>
      <BIRN_ID xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">301882920</BIRN_ID>
      <Diagnostic_group xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">h</Diagnostic_group>
      <Voxels xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">232</Voxels>
      <Max_Z xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">10.1</Max_Z>
      <P>1.39E-4</P>
      <log10P>3.86</log10P>
```

Upload to

Save to File







## Query Search



Field:

title



Type:

contains



Pattern:

Talairach

Language:

VSparQL



User:

brinkley



Search

## Query Search



User	Title	Description	Language	isPublic
brinkley	AMIA 2010: Talaira	List all structures that have one or more	VSparQL	true
brinkley	75: AMIA 2010: T	First find direct subclasses, then transiti	VSparQL	true
brinkley	77: Test AMIA 201	First find direct subclasses, then transiti	VSparQL	true
brinkley	74: AMIA 2010: T	First find direct subclasses, then transiti	VSparQL	true

New Search

Delete

Select



Structural  
Informatics  
Group

# Query Manager

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## Title

74: AMIA 2010: Talairach: Regional or constitutional parts of a structure that has lateralized subclasses: Dorsolateral\_prefrontal\_cortex

## Description

First find direct subclasses, then transitive regional parts, output those that have Talairach labels.  
To change structure substitute different structure name for Dorsolateral\_prefrontal\_cortex

```

<rdf:RDF
  xmlns:rdf="http://www.w3.org/1999/02/22-rdf-syntax-ns#"
  xmlns:apf="java:com.hp.hpl.jena.query.pfunction.library."
  xmlns:tmp="http://sig.biostr.washington.edu/tmp#"
  xmlns:gleen="java:edu.washington.sig.gleen."
  xmlns:owl="http://www.w3.org/2002/07/owl#"
  xmlns:rdfs="http://www.w3.org/2000/01/rdf-schema#"
  xmlns:fma="http://sig.uw.edu/fma#" >
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_46_of_right_middle_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Right Cerebrum.Frontal Lobe.Mid
  </rdf:Description>
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_46_of_left_middle_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Left Cerebrum.Frontal Lobe.Mid
  </rdf:Description>
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_9_of_left_middle_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Left Cerebrum.Frontal Lobe.Mid
  </rdf:Description>
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_9_of_left_superior_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Left Cerebrum.Frontal Lobe.Supe
  </rdf:Description>
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_9_of_right_superior_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Right Cerebrum.Frontal Lobe.Sup
  </rdf:Description>
  <rdf:Description rdf:about="http://sig.uw.edu/fma#Brodmann_area_9_of_right_middle_frontal_gyrus">
    <fma:Talairach rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Right Cerebrum.Frontal Lobe.Mid
  </rdf:Description>
</rdf:RDF>

```



```
let $sub_regions_tals := doc("http://xiphoid.biostr.washington.edu:8080/ValueSetService/ValueSet?qid=74")
```

```
(: now gather results from activation data :)
```

```
let $act_doc := doc("http://synapse.biostr.washington.edu/~brinkley/dxdemos/JessMerged.xml")
```

```
(: first filter by Talarach label that is also in list returned rom vSPARQL query :)
```

```
let $act_by_tal :=
```

```
  for $act in $act_doc//site
```

```
  for $fmatal in $sub_regions_tals
```

```
  return if (fn:contains($fmatal,$act/Tal))
```

```
  then $act
```

```
  else ()
```

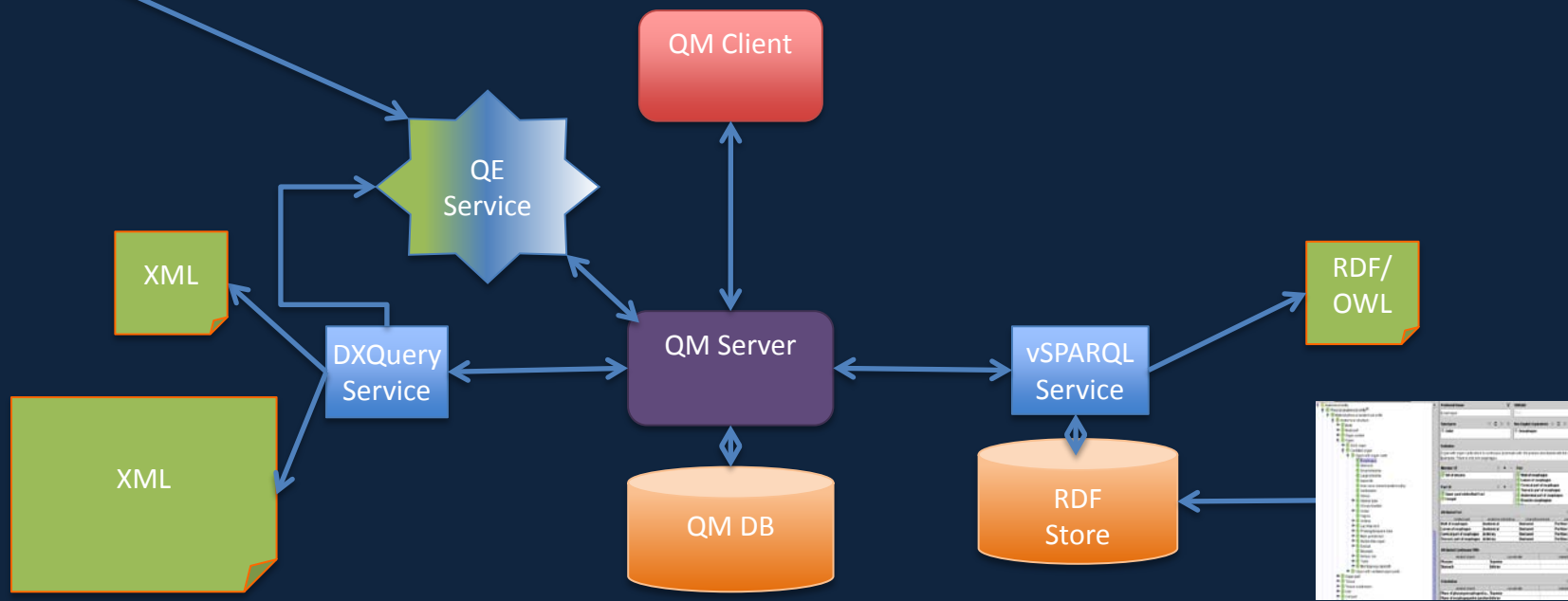
```
(: now filter by healthy vs sz :)
```

```
let $healthy_act := $act_by_tal[Diagnostic_group/text()='h']
```

```
let $schizo_act := $act_by_tal[Diagnostic_group/text()='s']
```

## Query Results

```
<?xml version="1.0" encoding="UTF-8"?>
<results>
  <patient>
    <pnum>MNI</pnum>
    <site>
      <site_label>310542253_5.55</site_label>
      <BIRN_ID xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">310542253</BIRN_ID>
      <Diagnostic_group xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">h</Diagnostic_group>
      <Voxels xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">651</Voxels>
      <Max_Z xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">5.55</Max_Z>
      <P>3.74E-4</P>
      <log10P>3.43</log10P>
      <Distance>18</Distance>
      <Tal> Right Cerebrum.Frontal Lobe.Middle Frontal Gyrus.Gray Matter.Brodmann area 46</Tal>
      <right_coord>50</right_coord>
      <sup_coord>30</sup_coord>
      <ant_coord>40</ant_coord>
      <Color>Red</Color>
      <Shape>BIG_SPHERE</Shape>
      <Size>1.75</Size>
    </site>
  </patient>
  <site>
    <site_label>311987949_7.95</site_label>
    <BIRN_ID xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">311987949</BIRN_ID>
    <Diagnostic_group xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">h</Diagnostic_group>
    <Voxels xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">268</Voxels>
    <Max_Z xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">7.95</Max_Z>
    <P>0.0306</P>
    <log10P>1.51</log10P>
    <Distance>1</Distance>
    <Tal> Left Cerebrum.Frontal Lobe.Middle Frontal Gyrus.Gray Matter.Brodmann area 9</Tal>
    <right_coord>-26</right_coord>
```

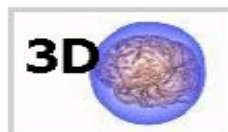
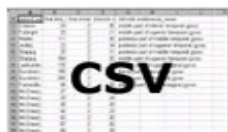


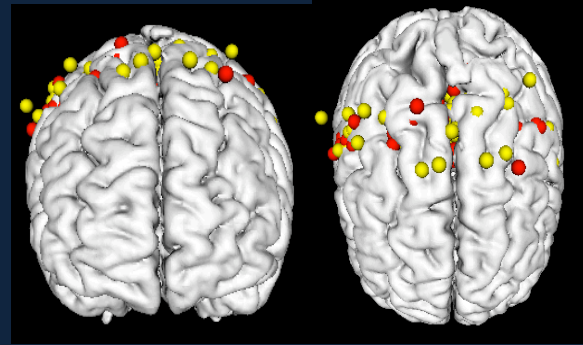
**Title (max 200 chars):****Description (max 400 chars):****Query**

```
let $jess4 := doc("http://xiphoid.biostr.washington.edu:8080/ValueSetService/ValueSet?qid=68")
return
<results>
{$jess4//patient}
</results>
```

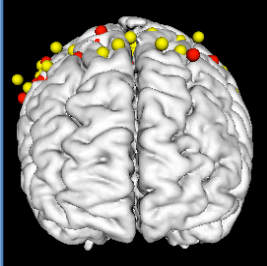
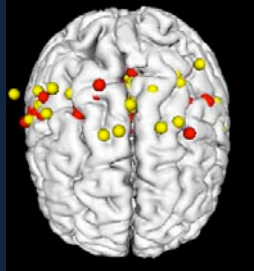
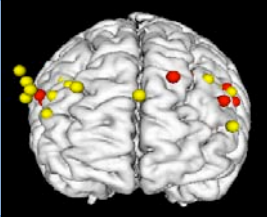
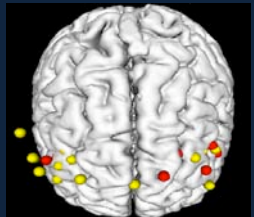
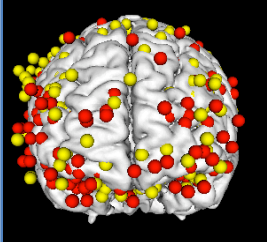
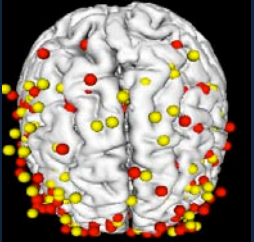
Public  Private   

Select one of the following output formats.





N=1741

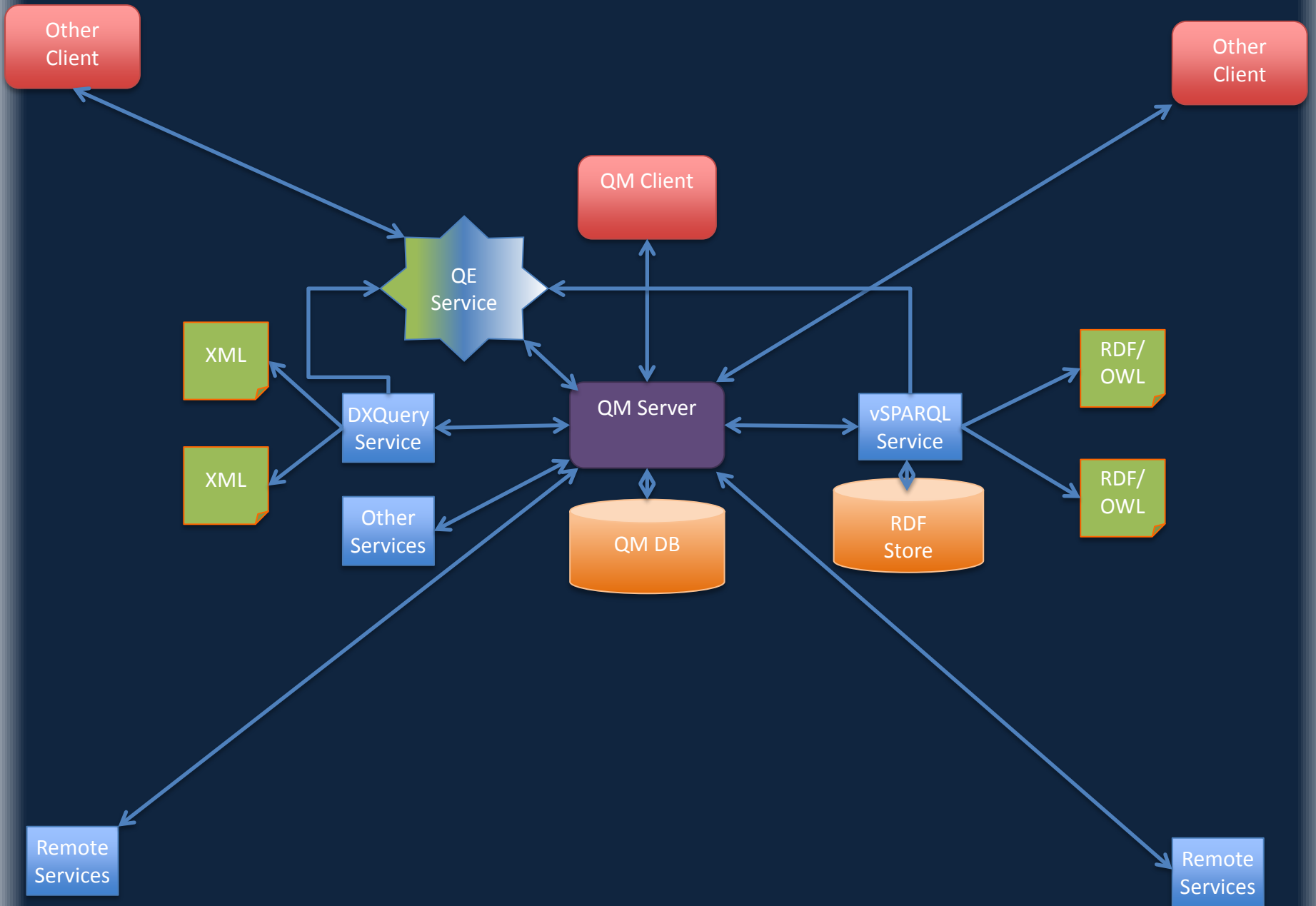
Region	Filtered	Healthy (Red)	Schizophrenic (Yellow)		
Dorso_lateral prefrontal cortex	55	25	30		
Brodmann area 6	213	107	106		
Frontal lobe	575	288	287		



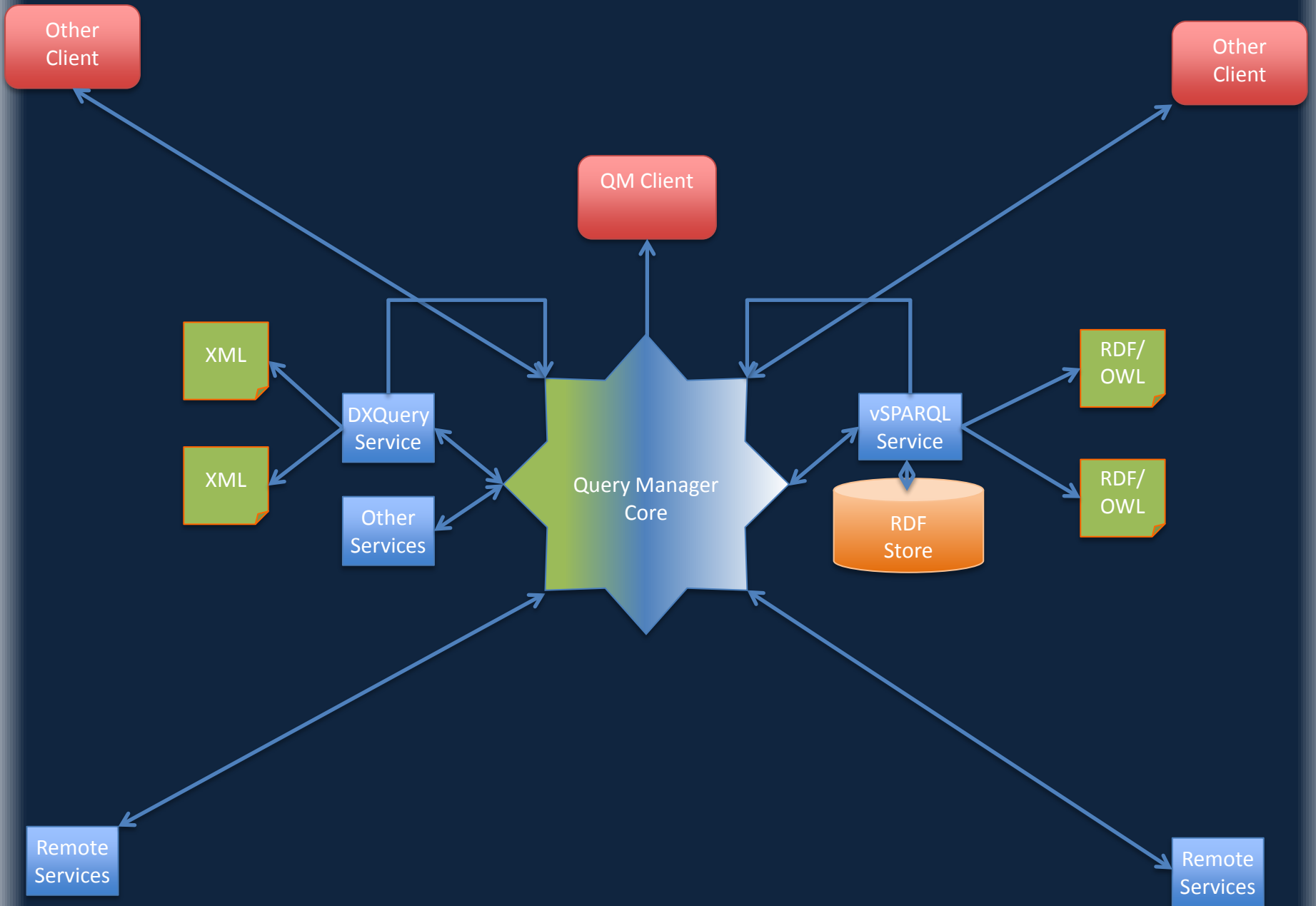
```
graph TD
    subgraph "graph TD"
        QM_Client[QM Client] <--> QM_Server[QM Server]
        QM_Server <--> QM_DB[(QM DB)]
        QM_Server <--> vSPARQL_Service[vSPARQL Service]
        vSPARQL_Service <--> RDF_Store[(RDF Store)]
        QM_Server <--> QE_Service{QE Service}
        QE_Service <--> DXQuery_Service[DXQuery Service]
        DXQuery_Service <--> QM_Server
    end
```

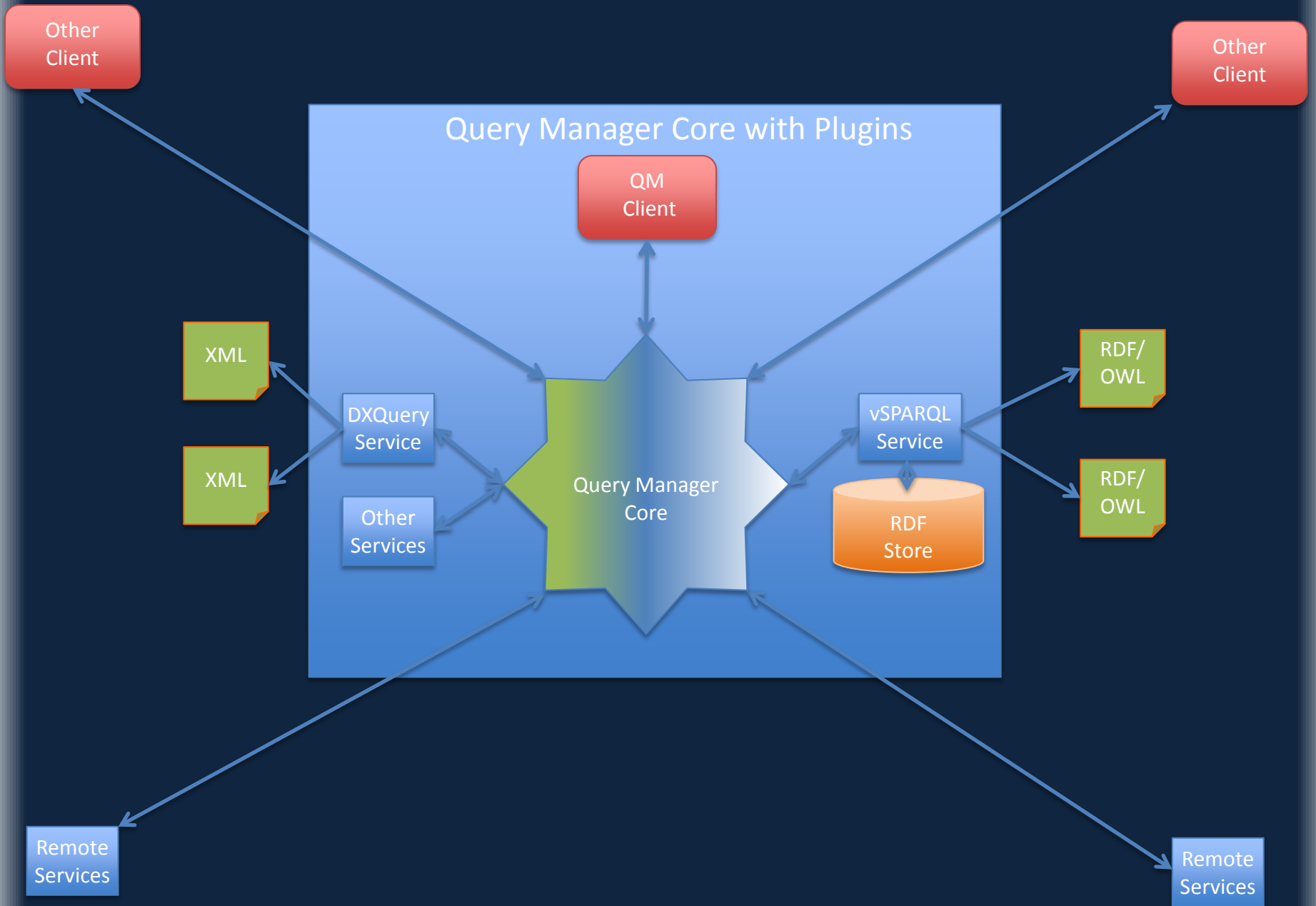
```
graph TD
    subgraph "graph TD"
        QM_Client[QM Client] <--> QM_Server[QM Server]
        QM_Server <--> QM_DB[(QM DB)]
        QM_Server <--> vSPARQL_Service[vSPARQL Service]
        vSPARQL_Service <--> RDF_Store[(RDF Store)]
        QM_Server <--> QE_Service{QE Service}
        QE_Service <--> DXQuery_Service[DXQuery Service]
        DXQuery_Service <--> QM_Server
    end
```











Other Client

Other Client

XML

XML

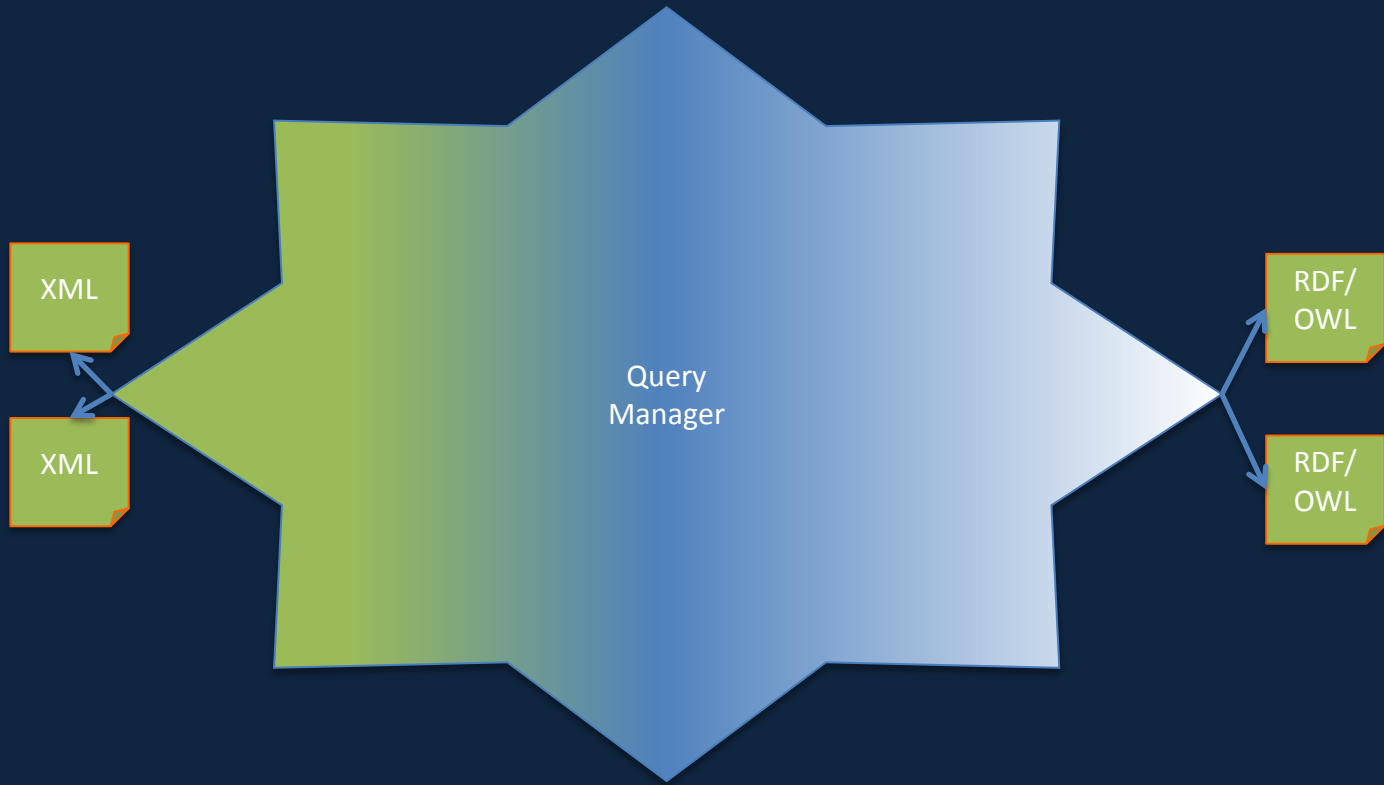
RDF/  
OWL

RDF/  
OWL

Query  
Manager

Remote  
Services

Remote  
Services







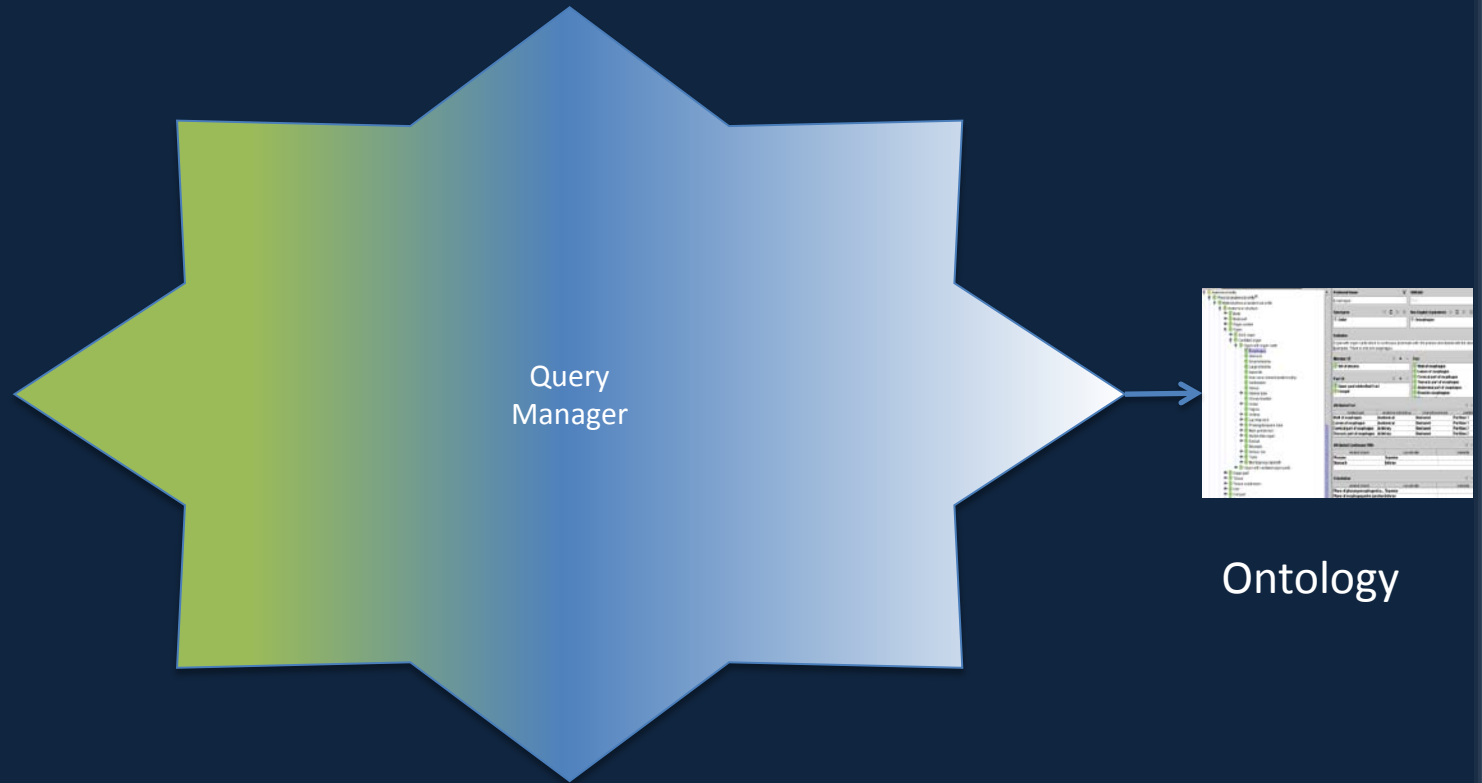




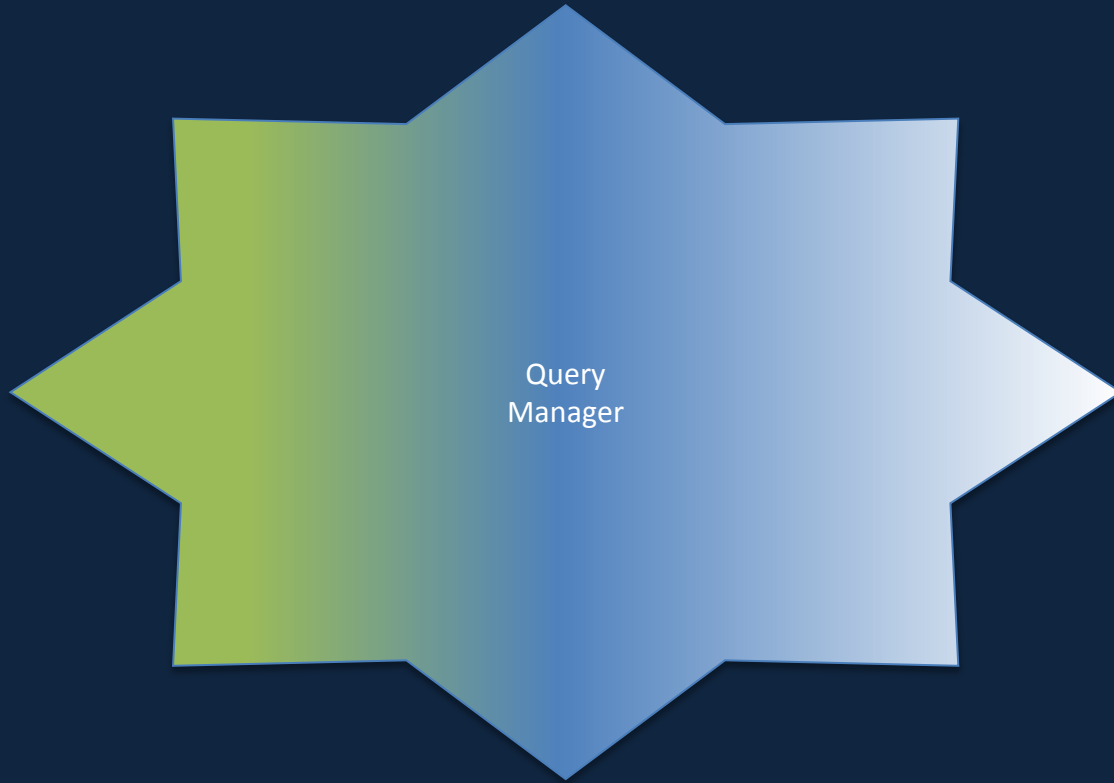




# Ontology Views



# Summary

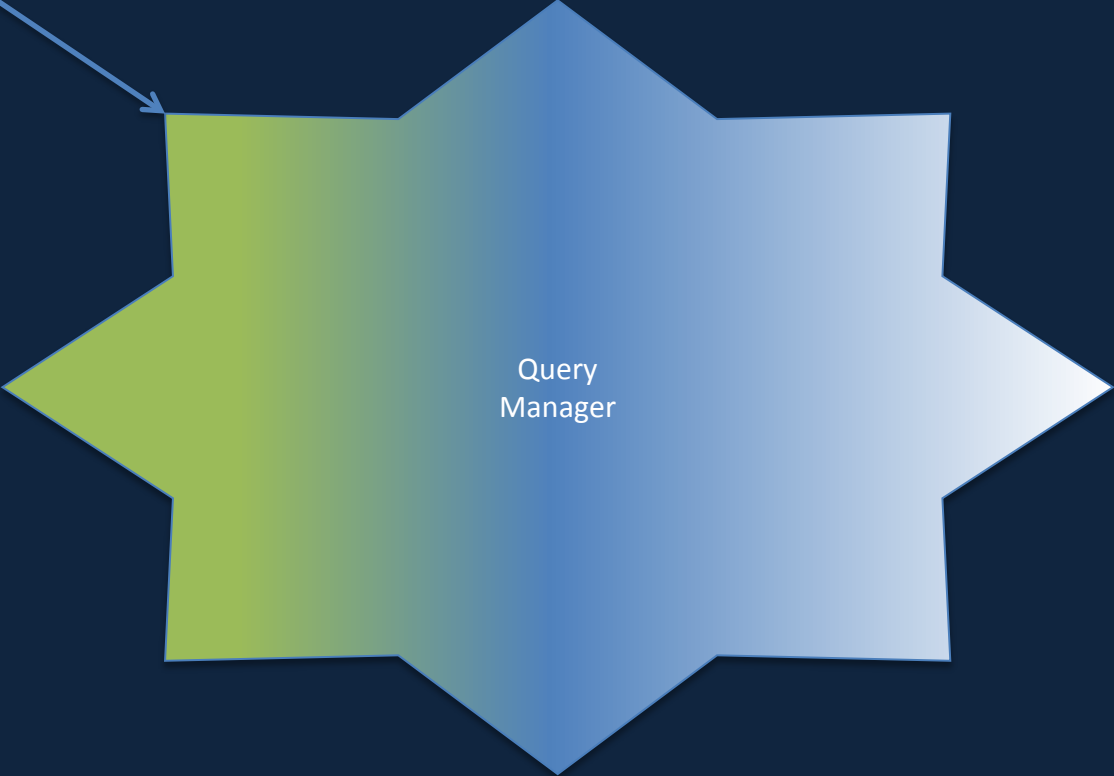


# Issues

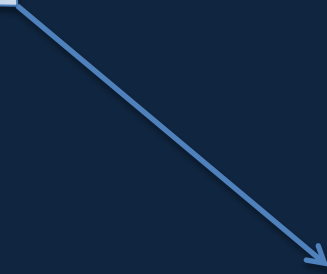
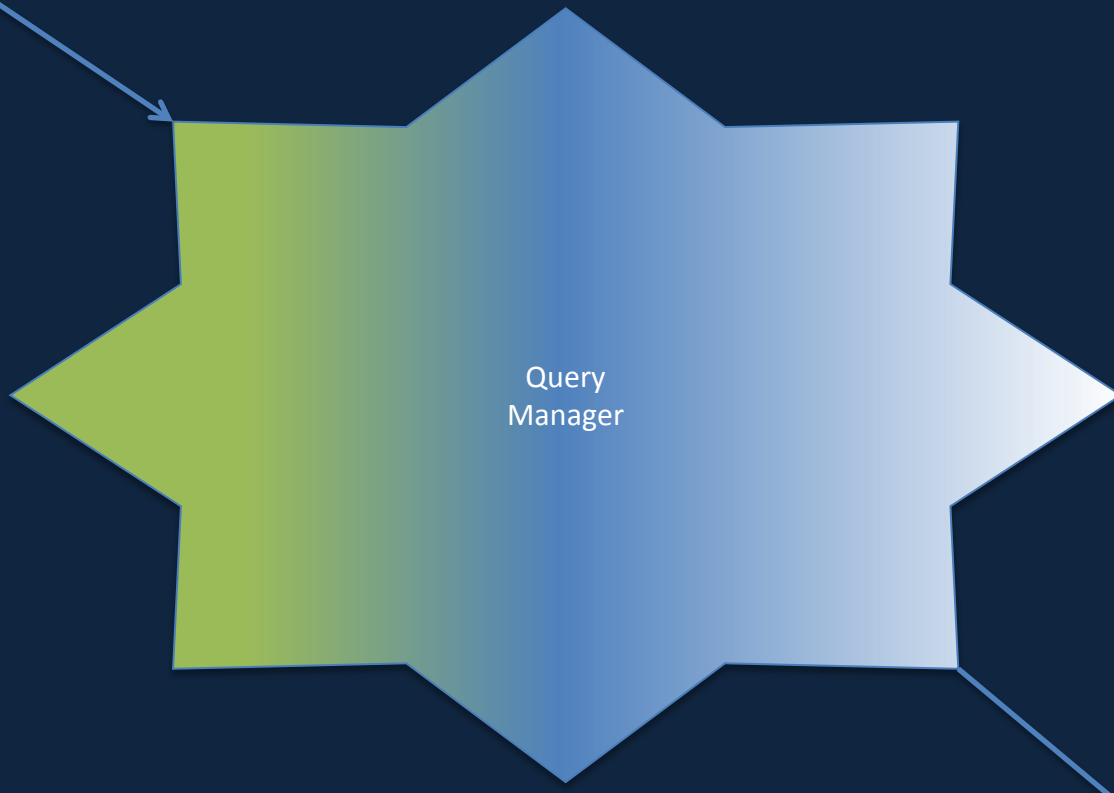
- Response time
- Reliability
- Security
- Discoverability
- Usability
- Integration with other systems and methods

# Future Work

Query  
Manager



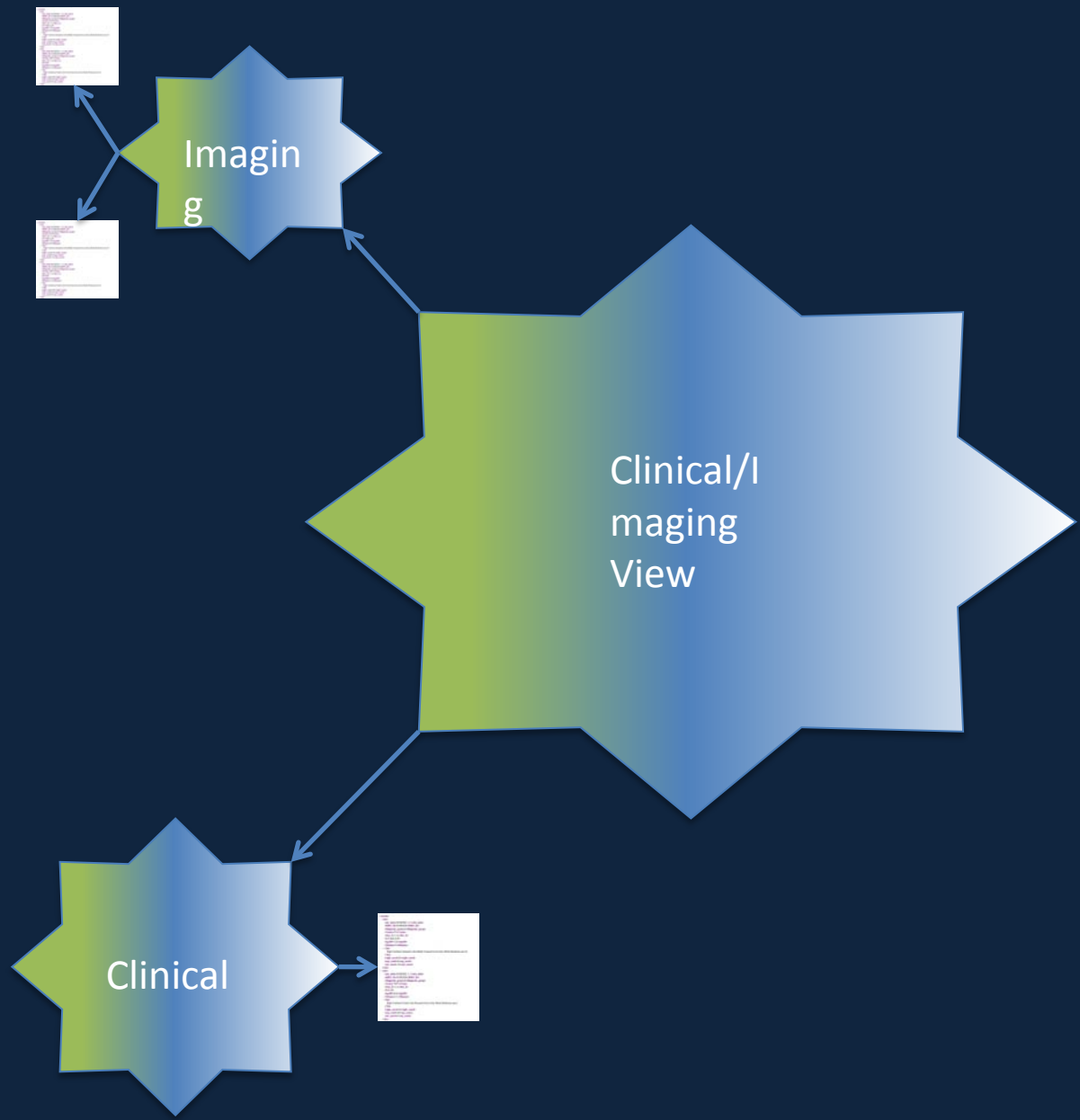
Query  
Manager



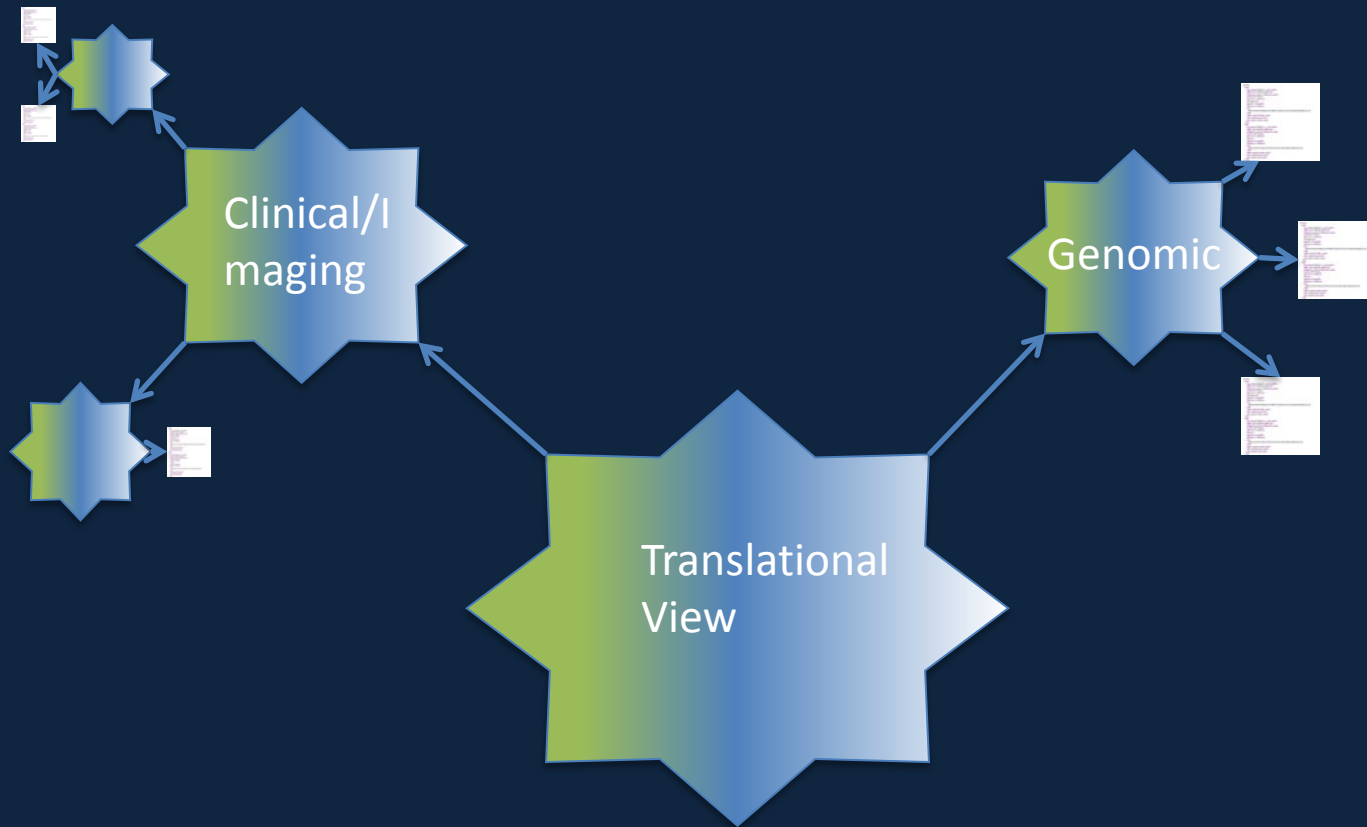


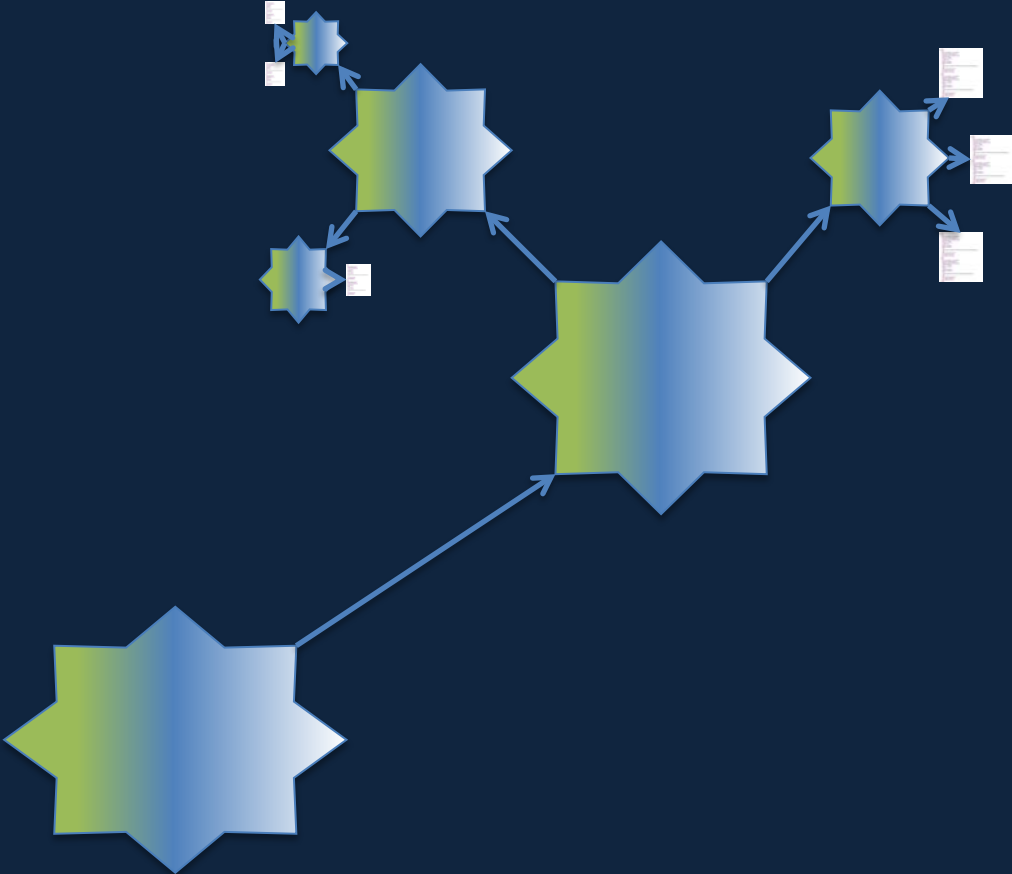


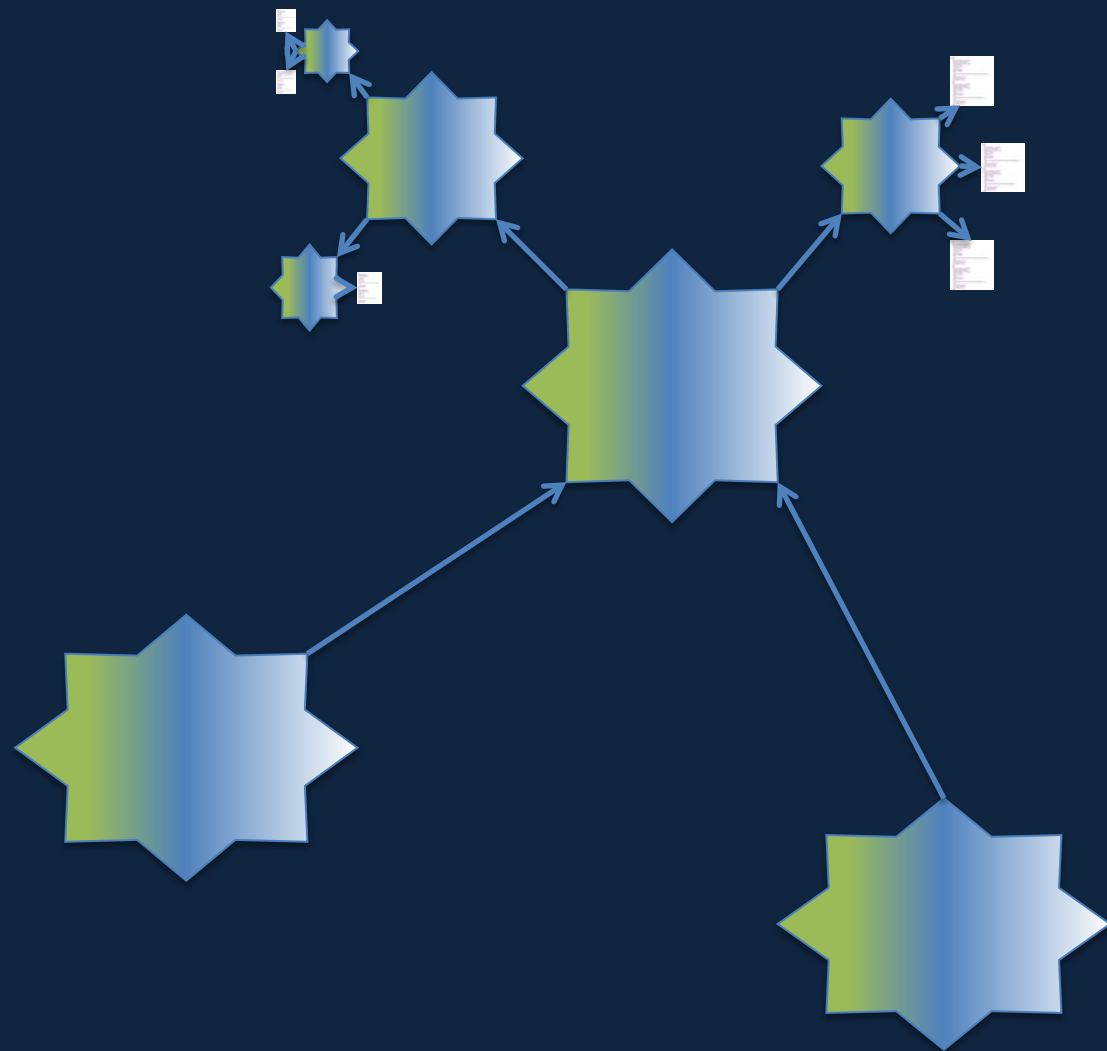




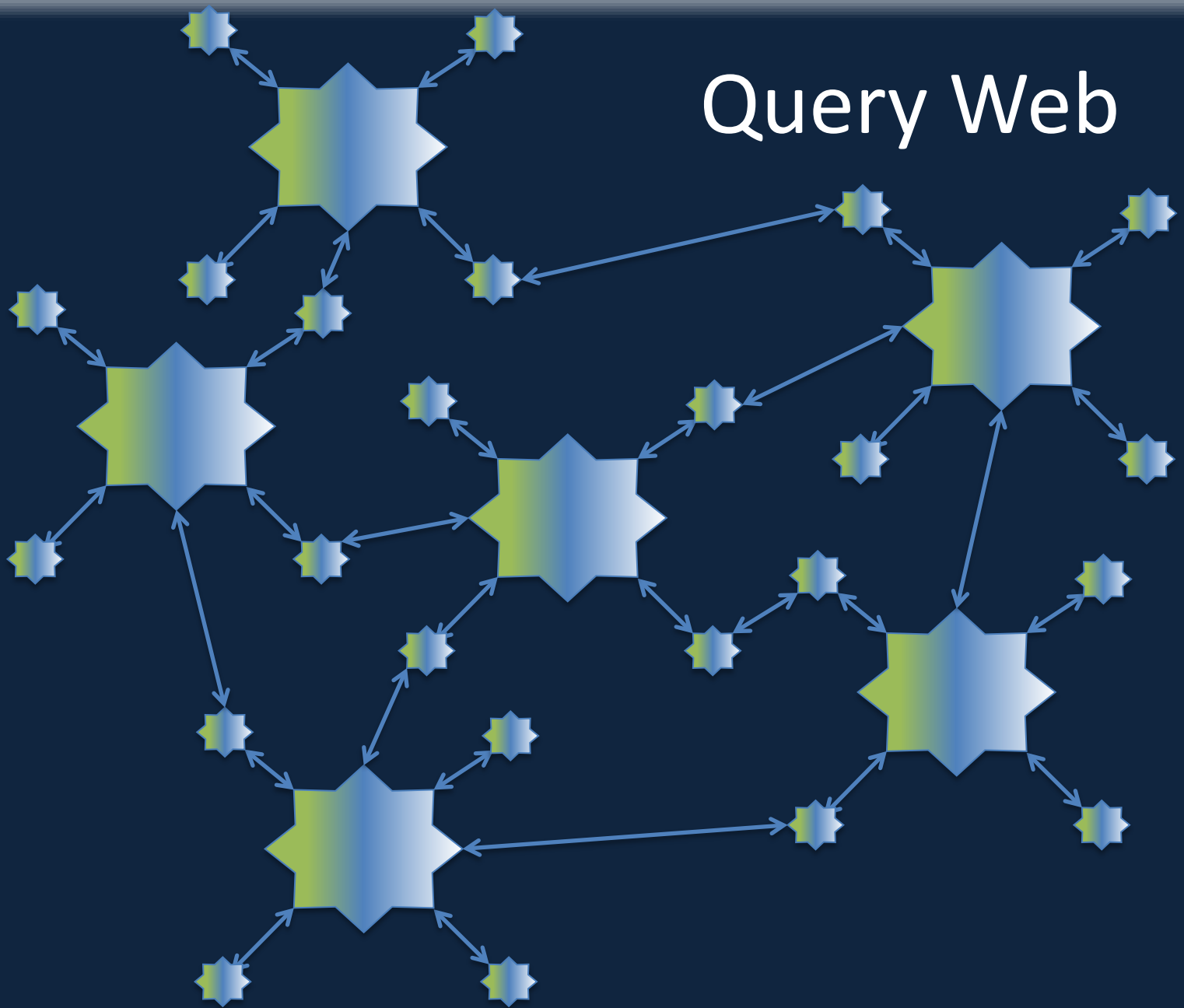




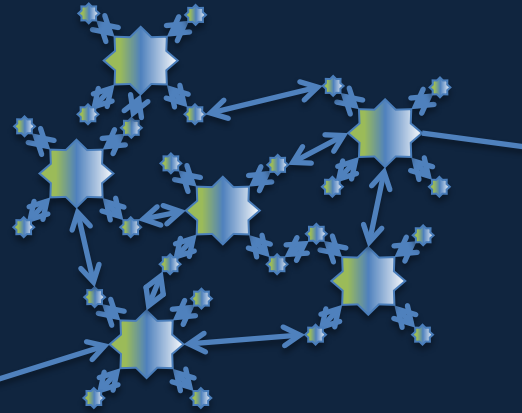




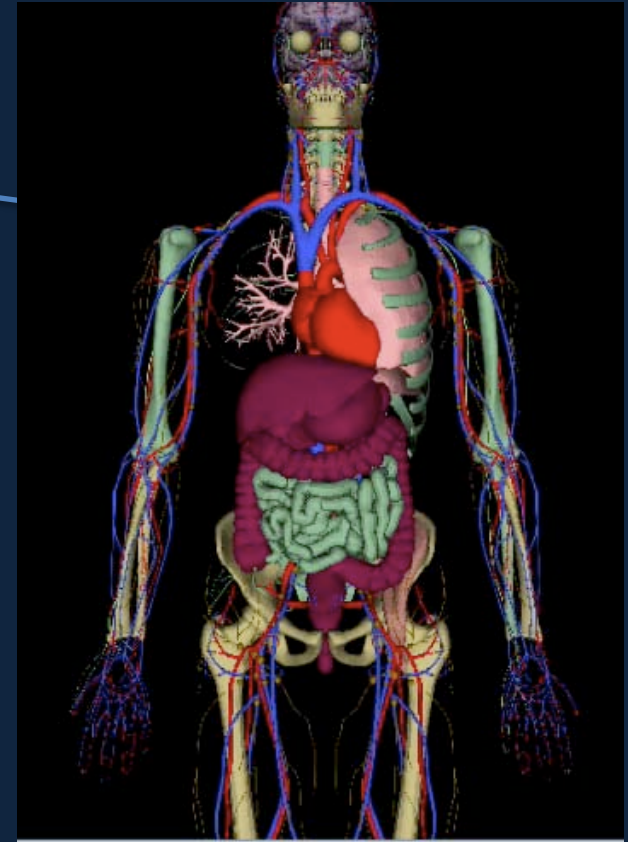
# Query Web



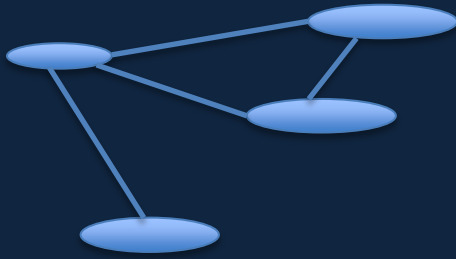
# Translational Medicine



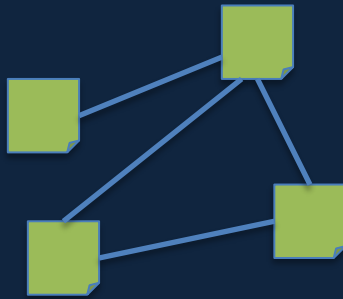
Query Web



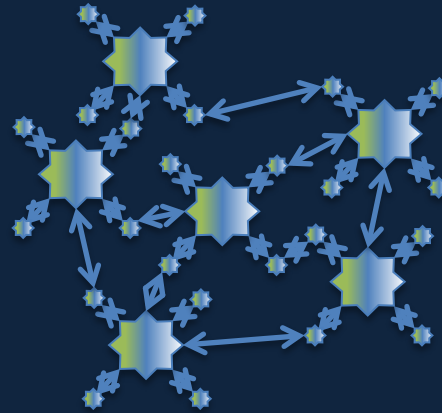
# A Network-centric view of information systems



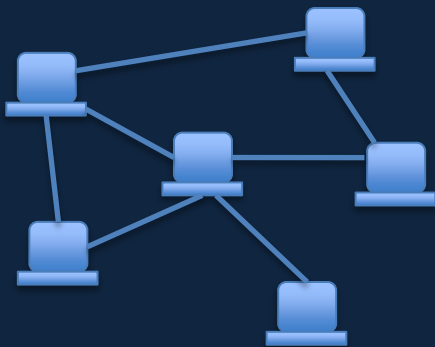
Semantic Web



Web



Query Web



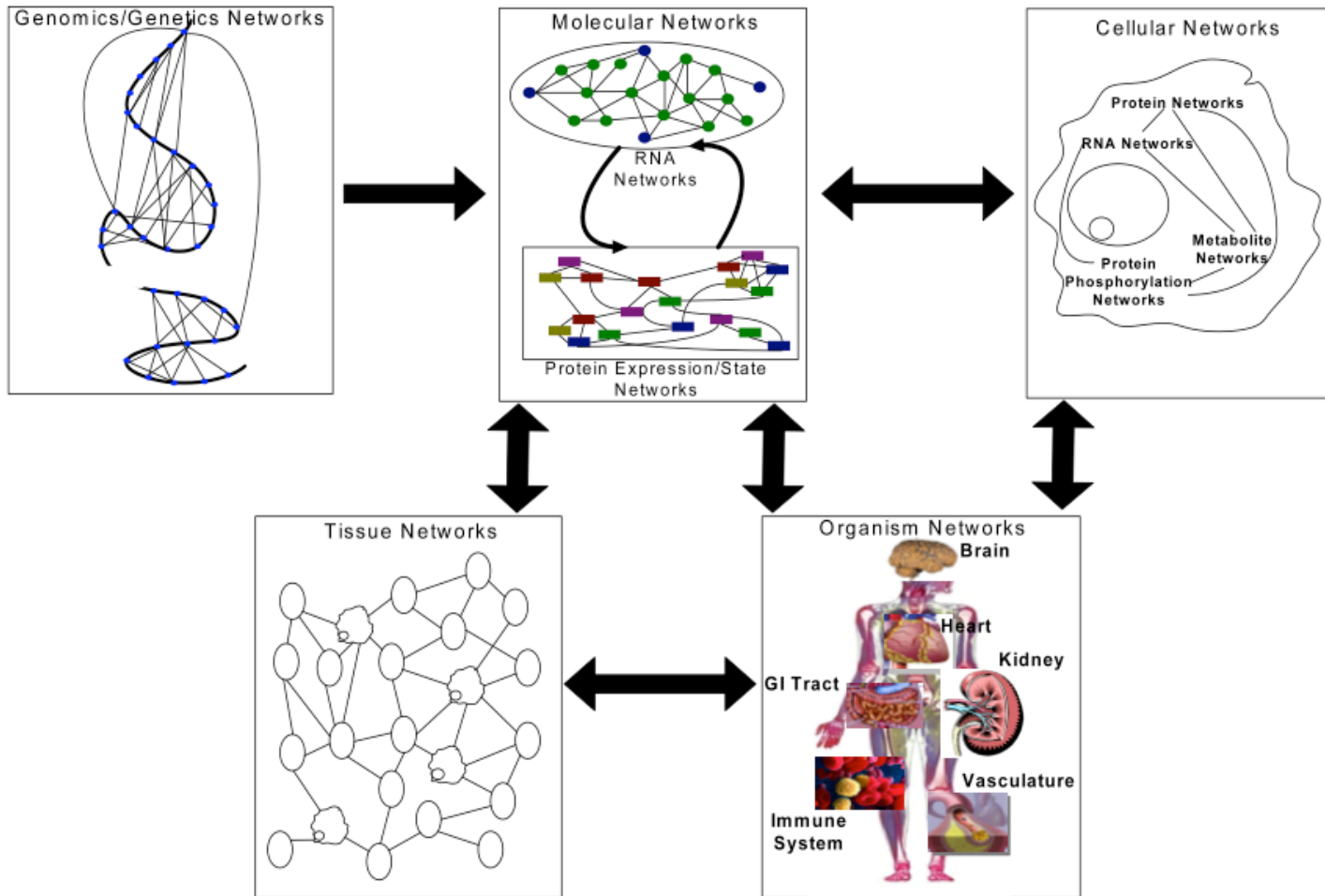
Internet





# A Network-centered View of Biological Systems

Bin Zhang, PhD



# A convergence?

- Biology as an information science
- Information science as biology
- Networks as the link

# Credits

- Todd Detwiler
- Onard Mejino
- Joshua Franklin
- Ron Shaker
- Xenia Hertenberg
- Nolan Nichols
- Marianne Shaw
- Wayne Warren
- Linda Shapiro
- Dan Suciu
- Cornelius Rosse