#### The Semantic Web and Linked Data

**Globally Unique Identifiers** 

SPARQL - how to explore Linked Data

#### A basic beginners guide

#### Mark Wilkinson CBGP UPM-INIA Universidad Politécnica de Madrid Spain



RDF is a "framework" - a way of conceptualizing how to represent data and knowledge. There are a variety of ways to implement this conceptualization.

It is ~NOT a language (per se...)

It is NOT a file-format or data syntax (though its implementations generally are...)



#### RDF says that all data and knowledge can be represented through a combination of very simple statements called "Triples"

# Much like human language, the statements take the form:

Subject Predicate Object





#### "Triples"

S P O

Mark owns Cat

#### "Triples"

S P O

Mark owns Cat

Cat name "Stormy"

#### "Triples"



The "Object" of one triple becomes the "Subject" of another triple



#### "Graph"



### Resource Description Framework "Resources" "Literals" "Predicates"



There is nothing smaller than a Triple

Only a **Resource** can be the subject of a triple



There is nothing smaller than a Triple

Only a **Resource** can be the subject of a triple





## **Globally Unique Identifiers**

(GUIDs)

#### There are two "flavours" of GUID

- Those that are globally unique based on statistical (un)likelihood of a collision
- 2) Those that are globally unique based on convention/agreement

Network Working Group Request for Comments: 4122 Category: Standards Track P. Leach Microsoft M. Mealling Refactored Networks, LLC R. Salz DataPower Technology, Inc. July 2005

A Universally Unique IDentifier (UUID) URN Namespace

Status of This Memo

This document specifies an Internet standards track protocol for the Internet community, and requests discussion and suggestions for improvements. Please refer to the current edition of the "Internet Official Protocol Standards" (STD 1) for the standardization state and status of this protocol. Distribution of this memo is unlimited.

Copyright Notice

Copyright (C) The Internet Society (2005).

128-bit number, generated by one of 5 approaches

Grains of sand on earth: 75,000,000,000,000,000

Number of possible GUIDs: 340,282,366,920,938,463,463,374,607,431,770,000,000

#### The Point:

If you use a GUID to identify something, there is an infinitesimally small (~0%) chance that someone else has used that same GUID to name something else

#### THEREFORE! (the point!)

You can safely merge your dataset with someone else's dataset, and not "collide"

#### There are two "flavours" of GUID

 Those that are globally unique based on statistical (un)likelihood of a collision

2) Those that are globally unique based on convention/agreement

"Conventions" can be used to ensure that a set of characters is globally unique

For example, DOIs and Handles

A governing agency assigns a prefix (e.g. 10.xxxx for DOIs) to an identifier-generating authority

That prefix can only be (meaningfully) used to generate identifiers within that authority.

Unless the authority is mischievous, they can ensure that they never re-use an identifier, and nobody else "is allowed to" create an identifier with that prefix. "Conventions" can be used to ensure that a set of characters is globally unique

URLs are another example

http://www.wilkinsonlab.info

I own that!

I create identifiers within the *wilkinsonlab.info* "namespace" I would be crazy to create the same address for two different pages!

(and in fact, *I cannot*, as you intuitively know)

"Conventions" can be used to ensure that a set of characters is globally unique

The only time I *could* experience a "collision" with someone else's name is if they are using the string of characters

"http://wilkinsonlab.info"

to mean something other than the identifier for my website (which would be crazy)

So... Web addresses are pretty good GUIDs!

URLs (Web Addresses) are also resolvable!

That is, if I have a URL, there is a standard approach to finding the "thing" that URL is identifying.

# The protocol is embedded in the URL itself (HTTP $\rightarrow$ the Web!)

This is distinct from other kinds of GUIDs like DOIs, where the identifier is not associated with a protocol (and therefore might be resolved in a wide variety of ways)





Before the detour we were talking about my cat

(we were also talking about RDF... but mostly about my cat ;-))

# What is the connection between RDF and GUIDs?



RDF is used on the Web as follows:

Resources are assigned URLs

Predicates are assigned URLs



#### Benefits of RDF that uses URLs



#### Benefits of RDF that uses URLs



#### All of these are globally unique, SHARED\* and unambiguous

\*meaning anyone, anywhere, can use them... BETTER TO NOT THINK OF THEM AS "LOCATIONS" → NAMES!

#### Benefits of RDF that uses URLs



All of these can be automatically looked-up by a machine!

Get the page at "http://pet.registry.com/MDW\_112/pets/MDW\_112\_Cat"



#### **Benefits of RDF**



 $\blacksquare$  URLs are globally Unique!  $\rightarrow$  Two sites with statements **about the same thing**...

### **Benefits of RDF**

#### **Unified Graph**



A machine can do this data integration automatically

# No special technology was required to do that integration!

#### Because we use GUIDs + RDF, the integration "just happened"

You simply download and concatenate the files!

(of course, there are more powerful ways to do this than downloading the data)

#### The Relationship between RDF and the Web

#### Gene - Detail

0000

search list download trait classification

#### Detail of Gene

Click here f

Basic Information		
CGSNL Gene Symbol	AGR1	
Gene Symbol Synonym	agr1*, agr1	
CGSNL Gene Name	AGRAVITROPISM 1	
Gene Name Synonym	agravitropism1, agravitropism1, agravitropism-1	
Protein Name		
Allele		
Chromosome No.		
Explanation	PO:0009009; embryo ; PO:0009005; root ; PO:0009025; leaf. GRO:0007047; 02-seedling ; GRO:0007043; 01-germination.	
Trait Class	Seed - Morphological traits - Embryo	
Expression		
Sequence/Locus		
Accession No.	2 ·	1
RAP ID	(IRGSP 1.0 / Build5)	
INSD Accession List (Test version)	-	
Мар		
Locate (cM)		7
References		
Hong, S.K., T, Aoki, H. Kitano, H Phenotypic diversity of 188	I. Satoh and Y. Nagato (1995) Dev Genet. rice embryo mutants.	]
TextPresso Search	Search textpresso for AGR1 ( Recent references may be retrievable, but without any warranty )	Hyperlinks to
DB Reference		Functional
Gramene ID	GR:0060019	
Ontologies		Annotations
Gene Ontology	multicellular organismal development( GO:0007275 ) gravitropism( GO:0009630 ) anatomical structure morphogenesis( GO:0009653 ) embryonic development( GO:0009790 )	
Trait Ontology	root anatomy and morphology trait( TO:0000043 ) leaf anatomy and morphology trait( TO:0000419 )	

#### Hyperlinks carry no meaning

multicellular organismal development( GO:0007275 ) gravitropism( GO:0009630 )

anatomical structure morphogenesis( GO:0009653 ) embryonic development( GO:0009790 )

root anatomy and morphology trait( TO:000043 ) leaf anatomy and morphology trait( TO:0000419 )

> What will happen if I click this link? Will I see a movie? Will I download a file?
Humans design nice Web pages

so that other Humans

can make good guesses

about what they want to click

Humans design nice Web pages

so that other Humans

can make good guesses

about what they want to click

That is the only reason that the Web is so successful! Human Intuition!

0.000

### The Relationship between RDF and the Web

#### Gene - Detail

0000

search list download trait classification

#### Detail of Gene

Click here f

Basic Information		
CGSNL Gene Symbol	AGR1	
Gene Symbol Synonym	agr1*, agr1	
CGSNL Gene Name	AGRAVITROPISM 1	
Gene Name Synonym	agravitropism1, agravitropism1, agravitropism-1	
Protein Name		
Allele		
Chromosome No.		
Explanation	PO:0009009; embryo ; PO:0009005; root ; PO:0009025; leaf. GRO:0007047; 02-seedling ; GRO:0007043; 01-germination.	
Trait Class	Seed - Morphological traits - Embryo	
Expression		
Sequence/Locus		
Accession No.		1
RAP ID	( IRGSP 1.0 / Build5 )	
INSD Accession List (Test version)	-	
Мар		
Locate (cM)		
References		
Hong, S.K., T, Aoki, H. Kitano, H Phenotypic diversity of 188	I. Satoh and Y. Nagato (1995) Dev Genet. rice embryo mutants.	1
TextPresso Search	Search textpresso for AGR1 (Recent references may be retrievable, but without any warranty)	Hyperlinks to
DB Reference		Functional
Gramene ID	GR:0060019	Annotations
Ontologies		Annotations
Gene Ontology	multicellular organismal development( GO:0007275 ) gravitropism( GO:0009630 ) anatomical structure morphogenesis( GO:0009653 ) embryonic development( GO:0009790 )	
Trait Ontology	root anatomy and morphology trait( TO:0000043 ) leaf anatomy and morphology trait( TO:0000419 )	

### The Relationship between RDF and the Web

Brander Brander Brander Brander	e Gen	A	miGO		
Search	Browse	BLAST	Homolog Annotations	Tools & Resources	Неір
Search GO			] 🙃 terms 🍙 genes of	r proteins 🦵 exact m	atch Submit

#### gravitropism

Term information ↓ Term neighborhood ↓ External references ↓ 157 gene product associations →

Term Info	Term Information					
Accession	GO:0009630					
Ontology	Biological Process					
Synonyms	exact: geotropism					
Definition	ার The orientation of plant parts under the stimulation of gravity. Source: ISBN:0198547684					
Comment	None					
Subset	None					
Community	Add usage comments for this term on the GONUTS wiki.					

#### http://www.shigen.nig.ac.jp/rice/oryzabaseV4/gene/detail/24

	Rice Science Datab		Search			<ul> <li>for</li> <li>Last updated</li> </ul>	GO!	
bout Strains	distribute/deposite	Education Genome	Dev. Genes	Tools	Download	Mar 6, 2013		
ene - Det	ail	والمتعاد						
arch list	download trait cl	assification						
Detail of	f Gene						Click here for feedb	
Basic Infor	rmation							
CGSNL Ge	ne Symbol	AGR1						
Gene Symb	ool Synonym	agr1*, agr1						
CGSNL Ge	ne Name	AGRAVITR	OPISM 1					
Gene Name	e Synonym	agravitropi	sm1, agravit	ropism	1, agravi	tropism-1		
Protein Nar	me							
Allele								
Chromoson	ne No.							
Explanation	1					oot ; PO:0009 7043; 01-ge		
Trait Class		Seed - Mor	Seed - Morphological traits - Embryo					
Expression								
Sequence/	Locus							
Accession I	No.	-						
RAP ID		(IRGSP 1.	0 / Build5)					
INSD Acces (Test versio		-						
Мар								
Locate (cM)	)							
Reference:	s							
		H. Satoh and Y. Nag 8 rice embryo muta		v Gene	rt.			
TextPresso	Search		presso for A but without			eferences ma	iy be	
DB Refere								
Gramene ID	D	GR:006001	9					
Ontologies								
Gene Ontol	logy	gravitropisr anatomical	m( GO:00096	530) orphog	enesis( C	GO:0007275 GO:0009653 ) )		
Trait Optolo		2002 - 2004	ny and morp	holog	y trait( TO	:0000043)		

#### http://amigo.geneontology.org/amigo/term=GO%3A0009630

Name and American Strength Str	e Gen	А	miGO		
Search	Browse	BLAST	Homolog Annotations	Tools & Resources	Help
Search GO			] 🌀 terms 🎧 genes o	r proteins 🦵 exact m	atch Submit

#### gravitropism

Term information ♣ Term neighborhood ♣ External references ♣ 157 gene product associations ➡						
Term Information						
Accession	GO:0009630					
Ontology	Biological Process					
Synonyms	exact: geotropism					
<b>Definition</b> The orientation of plant parts under the stimulation of gravity.						
	Source: ISBN:0198547684					
Comment	None					
Subset	None					
Community	Add usage comments for this term on the GONUTS wiki.					

A human knows what this "link" means - it means that the rice gene is involved in gravitropism

But there are ~ 1.3 Trillion Gigabytes of data on the Web! Too much for a human to ever explore...

We need a Web that can be explored by machines!!

http://wy	ww.shigen.nig	<u>ı.ac.jp</u> /rice/oryzabaseV4/ge	ne/deta	ail/24	http://amig	<u>go.geneontology.org</u> /amigo/term=GO%3A	.0009630
но	Coryzabase spor Integrated Rice Science Databa ome About Strains distribute/deposite Gene - Detail search list download trait cla	aksrd by NBRP ve Education Genome/Dev. Genes Tools Download Last updated: Education Genome/Dev. Genes Tools Download Last updated: assification	sanese   English		attat or other		AmiGO Resources Help
	Detail of Gene						- exact match Submit
	Basic Information						
	CGSNL Gene Symbol	AGR1					
	Gene Symbol Synonym	agr1*, agr1				gravitropicm	
	CGSNL Gene Name	AGRAVITROPISM 1				gravitropism	
	Gene Name Synonym	agravitropism1, agravitropism1, agravitropism-1				•	
	Protein Name						
	Allele				Term in	formation 4 Term neighborhood 4 External references 4 157 gen	e product associations 🐳
	Chromosome No.						•
	Explanation	PO:0009009; embryo ; PO:0009005; root ; PO:0009025; leaf. GRO:0007047; 02-seedling ; GRO:0007043; 01-germination.			Term Info	rmation	
	Trait Class	Seed - Morphological traits - Embryo					
	Expression				Accession		
	Sequence/Locus		ι 💋		ACCESSION	GO:0009630	
	Accession No.	-			Ontology		
	RAP ID	(IRGSP 1.0 / Build5)			Untology	Biological Process	
	INSD Accession List (Test version)	*			Synonyms	exact: geotropism	
	Мар					CARGE: geoticpism	
	Locate (cM)				Definition	The orientation of plant parts under the stimulation of gravity.	
	References					en sen en ser en	
	Hong, S.K., T, Aoki, H. Kitano, H Phenotypic diversity of 188	H. Satoh and Y. Nagato (1995) Dev Genet. 3 rice embryo mutants.			NAME OF COMPANY OF	Source: ISBN:0198547684	
	TextPresso Search	Search textpresso for AGR1 ( Recent references may be retrievable, but without any warranty )			Comment	None	
	DB Reference				Subset	News	
	Gramene ID	GR:0060019				None	
	Ontologies				Community		
	Gene Ontology	multicellular organismal development( GC:0007275 ) gravitropism( GC:0009630 ) anatomical structure morphogenesis( GC:0009653 ) embryonic development( GC:0009790 )				Add usage comments for this term on the GONUTS wiki.	
0000		root anatomy and morphology trait( TO:0000043 )					

### The Relationship between RDF and the Web



#### ...and in my perfect world...

we'd just eliminate the human-readable Web entirely! ;-)



#### ...and in my perfect world...

we'd just eliminate the human-readable Web entirely! ;-)



(There are ways of embedding RDF into HTML, so the two technologies are not mutually exclusive! ...but I won't be teaching you how to do that.

If you want to learn how to do it, search for "RDFa")

### This is called "The Semantic Web"



### "Semantics" implies "meaning" Where is the "meaning" in this picture?



"Semantics" implies "meaning" Where is the "meaning" in this picture? Two parts of a Graph are able to carry meaning



Predicates...

"Semantics" implies "meaning" Where is the "meaning" in this picture? Two parts of a Graph are able to carry meaning



#### and "rdf:type" nodes

(http://www.w3.org/1999/02/22-rdf-syntax-ns#type)

# Where does "meaning" come from?



# Ontologies

the branch of metaphysics dealing with the nature of being.





http://purl.bioontology.org/ontology/NCBITAXON/9685

There is global agreement on what "taxon:9685" means

Details	Visualization	Notes (0)	Class Mappings (17)	
Preferred	Name	Felis	catus	
Synonyms	S	Felis	domesticus	
		cats		
		cat		
		Felis	silvestris catus	
ID		http://	purl.bioontology.org/ontology/NCBITAXON/9685	
altLabel		Felis	domesticus	
		cats		
		cat		
		Felis	silvestris catus	
cui		C0007	7450	
DIV		Mamn	nals	
notation		9685		
prefLabel		Felis	catus	
RANK		specie	es	
tui		T015		
subClass(	Df	Felis		

http://purl.bioontology.org/ontology/NCBITAXON/9685

There is global agreement on what "taxon:9685" means (and... you can also get the same information in 100% machine-readable RDF! Not just the Web page)

Details Visualization	Notes (0) Class Mappings (17)
Preferred Name	Felis catus
Synonyms	Felis domesticus cats cat Felis silvestris catus
ID	http://purl.bioontology.org/ontology/NCBITAXON/9685
altLabel	Felis domesticus cats cat Felis silvestris catus
cui	C0007450
DIV	Mammals
notation	9685
prefLabel	Felis catus
RANK	species
tui	T015
subClassOf	Felis







# Ontologizing like crazy!





# EFO:0005238



# breed

#### http://www.ebi.ac.uk/efo/EFO\_0005238

A breed is a specific group of domestic animals or plants having homogeneous appearance, homogeneous behavior, and other characteristics that distinguish it from other animals or plants of the same species and that were arrived at through selective breeding.

#### parents

population

# We hypothesise that a LOT of new discoveries

### will come by automatic "inference"

on this Global Graph of all human knowledge



## Link The World!

# **Ontologize The World!**

:-)

Imagine if 1.3 Trillion Gigabytes of data were represented in this format, and linked into ontologies... what the machines could discover for us!

# What does RDF "look like"

Remember, RDF is a "framework" - a way of thinking about things

To be USEFUL we have to find a way to represent it and pass it from machine-to-machine

There are several choices of "serialization":

- XML (RDF-XML)
- n3 (N-triples)
- Turtle
- RDFa



### Consider the BRCA1 gene

### Its URL is: http://identifiers.org/ncbigene/672

It would be tedious (and error-prone) to type this over and over again.

So we use "Namespaces" to make it easier

http://identifiers.org/ncbigene/672

# http://identifiers.org/ncbigene/672

ID

Namespace



http://identifiers.org/ncbigene/

672

### PREFIX ns: <http://identifiers.org/ncbigene/>

672

The prefix can be any series of characters; you can choose whatever you wish

PREFIX ncbi: <a href="http://identifiers.org/ncbigene/">http://identifiers.org/ncbigene/</a>

672

### PREFIX ncbi: <http://identifiers.org/ncbigene/>

ncbi:672
#### PREFIX ncbi: <http://identifiers.org/ncbigene/>

#### ncbi:672

And for the rest of the document you refer to BRCA1 as ncbi:672

easy!



@prefix rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#>.
@prefix foaf: <http://xmlns.com/foaf/0.1/> .
@prefix prov: <http://www.w3.org/ns/prov#> .
@prefix rdfs: <http://www.w3.org/2000/01/rdf-schema#> .

<https://linkeddata.systems/markdwilkinson.foaf>

a	<pre>prov:Person , foaf:Person ;</pre>
rdfs:label	"Mark D. Wilkinson" ;
foaf:account	

```
<https://orcid.org/0000-0001-6960-357X#orcid-id> ;
foaf:based_near [
```

```
a gn:Feature ;
gn:countryCode "ES" ;
gn:parentCountry <http://sws.geonames.org/2510769/] ;
foaf:familyName "Wilkinson" ;
foaf:givenName "Mark" ;
foaf:name "Mark D. Wilkinson" ;
```

#### N-Triples: ~~hard for humans!

<http://example.org/bob#me> <http://www.w3.org/1999/02/22-rdf-syntax-ns#type>

<http://xmlns.com/foaf/0.1/Person> .

<http://example.org/bob#me> <http://xmlns.com/foaf/0.1/knows> <http://example.org/alice#me> .

<http://example.org/bob#me> <http://schema.org/birthDate>

"1990-07-04"^^<http://www.w3.org/2001/XMLSchema#date>.

<http://example.org/bob#me> <http://xmlns.com/foaf/0.1/topic\_interest>

<a href="http://www.wikidata.org/entity/Q12418">http://www.wikidata.org/entity/Q12418</a>.

<http://www.wikidata.org/entity/Q12418> <http://purl.org/dc/terms/title> "Mona Lisa" .

<http://www.wikidata.org/entity/Q12418> <http://purl.org/dc/terms/creator>

<http://dbpedia.org/resource/Leonardo\_da\_Vinci>.

<http://data.europeana.eu/item/04802/243FA8618938F4117025F17A8B813C5F9AA4D619> <http://purl.org/dc/terms/subject> <http://www.wikidata.org/entity/Q12418> .

#### **RDF-XML**

<?xml version="1.0" encoding="utf-8"?> <rdf:RDF xmlns:dcterms="http://purl.org/dc/terms/" xmlns:foaf="http://xmlns.com/foaf/0.1/" xmlns:rdf="http://www.w3.org/1999/02/22-rdf-syntax-ns#" xmlns:schema="http://schema.org/"> <rdf:Description rdf:about="http://example.org/bob#me"> <rdf:type rdf:resource="http://xmlns.com/foaf/0.1/Person"/> <schema:birthDate rdf:datatype="http://www.w3.org/2001/XMLSchema#date">1990-07-04</schema:birthDate> <foaf:knows rdf:resource="http://example.org/alice#me"/> <foaf:topic interest rdf:resource="http://www.wikidata.org/entity/Q12418"/> </rdf:Description> <rdf:Description rdf:about="http://www.wikidata.org/entity/Q12418"> <dcterms:title>Mona Lisa</dcterms:title> <dcterms:creator rdf:resource="http://dbpedia.org/resource/Leonardo da Vinci"/> </rdf:Description> <rdf:Description rdf:about="http://data.europeana.eu/item/04802/243FA8618938F4117025F17A8B813C5F9AA4D619"> <dcterms:subject rdf:resource="http://www.wikidata.org/entity/Q12418"/> </rdf:Description> </rdf:RDF>

```
<?xml version="1.0" encoding="utf-8"?>
<rdf:RDF
xmlns:foaf="http://xmlns.com/foaf/0.1/"
xmlns:rdf="http://www.w3.org/1999/02/22-rdf-syntax-ns#">
```

<rdf:Description rdf:about="http://example.org/bob#me"> <rdf:type rdf:resource="http://xmlns.com/foaf/0.1/Person"/> <foaf:knows rdf:resource="http://example.org/alice#me"/>

#### Interpreted as:

foaf:knows ⇒ http://xmlns.com/foaf/0.1/knows rdf:Description ⇒ http://www.w3.org/1999/02/22-rdf-syntax-ns#Description @prefix contact: <http://www.w3.org/2000/10/swap/pim/contact#>.

<http://linkeddata.systems/markdwilkinson.foaf>
 a contact:Person;
 contact:fullName "Eric Miller";
 contact:mailbox <mailto:em@w3.org>;
 contact:personalTitle "Dr.".

Makes it easier to read (for humans).

(but readability ISN'T the reason that namespaces exist!! ...that's another topic...)

# How much data in the world is represented in RDF?

(hint... a lot!)



http://lod-cloud.net/versions/2014-08-30/lod-cloud.svg: Linking Open Data cloud diagram **2014**, by Max Schmachtenberg, Christian Bizer, Anja Jentzsch and Richard Cyga<u>niak. http://lod-cloud.net/</u>



# LOD Cloud Today

Last updated: 2017-02-20



### How do I get to the data?

SPARQL - the query language for RDF

You can query RDF Databases (RDF databases are called "Triplestores")

Many Triplestores include a "SPARQL Endpoint" where you can send queries in the SPARQL language.

With SPARQL you create a "graph pattern" and ask the Triplestore to look for everything that matches the pattern

SPARQL looks quite a bit like RDF!



# What are the names of all British Longs?



# What are the names of all British Longs?



# What are the names of all British Longs?





PREFIX pred: <http: relations="" some.predicates.org=""></http:> PREFIX rdfs: <http: 01="" 2000="" rdf-schema#="" www.w3.org=""> PREFIX foaf: <http: 0.1="" foaf="" xmlns.com=""></http:></http:>			
SELECT WHERE	•	· —	"Stormy" . ?breed . ?breedname . ?species

rdf:type is so commonly used on the Semantic Web that it has it's own special SPARQL abbreviation: "a"



Stormy a Cat ⇒ Pumpkin rdf:type Cat

# Now you.... Create the SPARQL for

"what predicate connects a cat and the cat's name?"



# Now you.... Create the SPARQL for

"what predicate connects a cat and the cat's name?"



# Now you.... Create the SPARQL for

"what predicate connects a cat and the cat's name?"



- You can use a URL explicitly in your query
- Predicates can also be variables that you SELECT

- You can use a URL explicitly in your query
- Predicates can also be variables that you SELECT

#### How do you find SPARQL endpoints

Google is your fastest option, but:

http://lod-cloud.net/ that diagram is clickable



(but only from April 15, 2016 until April 16, 2016)

(but only from April 15, 2016 until April 16, 2016)

(but only from April 15, 2016 until April 16, 2016)



(but only from April 15, 2016 until April 16, 2016)



(but only from April 15, 2016 until April 16, 2016)



(but only from April 15, 2016 until April 16, 2016)



#### (but only from April 15, 2016 until April 16, 2016)



#### (but only from April 15, 2016 until April 16, 2016)



(but only from April 15, 2016 until April 16, 2016)



(but only from April 15, 2016 until April 16, 2016)

What does that look like in RDF?



You *really* want to say something about this entire triple

A way to group triples into a shared context The **context** can become the subject of triples


Context is just another URI...

E.g. http://example.org/some/context/123456

Instead of Triples, you use Quads

- S: http://people.org/person/Mark
  - P: http://activity.org/teaches
    - O: http://technology.org/W3C/SPARQL
      - C: http://example.org/contexts/1234546

- S: http://example.org/contexts/1234546
  - P: http://daterange.com/begins
    - O: "April 15, 2015"

Instead of Triples, you use Quads

- S: http://people.org/person/Mark
  - P: http://activity.org/teaches
    - O: http://technology.org/W3C/SPARQL

C: http://example.org/contexts/1234546

S: http://example.org/contexts/1234546

The context URI = the name of the Named Graph

- P: http://daterange.com/begins
  - O: "April 15, 2015"

---- Null -----

Instead of Triples, you use Quads

- S: http://people.org/person/Mark
  - P: http://activity.org/teaches
    - O: http://technology.org/W3C/SPARQL
      - C: http://example.org/contexts/1234546



---- N11]] -----

O: "April 15, 2015"

The information about that context

Instead of Triples, you use Quads

- S: http://people.org/person/Mark
  - P: http://activity.org/teaches
    - O: http://technology.org/W3C/SPARQL

C: http://example.org/contexts/1234546

S: http://people.org/person/Mark

The triples in this graph share a context, and are in the same "Named Graph"

P: http://activity.org/resides

O: http://hotels.com/NH Utrecht

C: http://example.org/contexts/1234546

Browse to: http://training.fairdata.solutions/sparql



Browse to: http://training.fairdata.solutions/sparql



Typical uses include:

- Versioning
  - http://database.org/version/1.1
  - http://database.org/version/1.2
  - You query whichever version you need by putting the appropriate named graph in the box above the SPARQL query
- NanoPublications
  - Nanopubs use named graphs to attach citation/authorship information to scientific data and results. "This set of triples came from...."

Browse to: http://ltraining.fairdata.solutions/sparql

#### This is a "shortcut", since most of the time you will always want to query the same graph. This allows you to set the desired graph as default.



(Security restrictions of this server do not allow you to retrieve remote RDF data, see details.)

Browse to: http://linkeddata.systems:8890/sparql

This is a "shortcut", since most of the time you will always want to query the same graph. This allows you to set the desired graph as default.



(Security restrictions of this server do not allow you to retrieve remote RDF data, see details.)

Typical uses include:

- Versioning
  - http://database.org/version/1.1
  - http://database.org/version/1.2
  - You query whichever version you need by putting the appropriate named graph in the box above the SPARQL query
- NanoPublications
  - Nanopubs use named graphs to attach citation/authorship information to scientific data and results. "This set of triples came from...."

#### A way to attach citation information to data



In a few hours we will discuss the **World Wide Web Consortium's** proposal for the "**Linked Data Platform**", which uses Named Graphs to "simulate" a filesystem, with folders and files.

**Important Point:** We can do FAIR using ONLY W3C-approved standards.

FAIR invents NOTHING