

MASTER'S THESIS

CONFIGURABLE SCHEMA-AWARE RDF DATA INPUT FORMS

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Abstract

Ontologies describing biological research processes are getting prevalent nowadays. They are used to define the rules of the investigation, and establish a basis for their standardized. The goal is to replace textual documentation of the processes, with structured data. The advantage of such representation that information can be exchanged and software agents can perform reasoner algorithm. The data describing the investigations the ontologies classes for the entities, measurement analysis steps etc. Furthermore the rules of these processes are described by means of restrictions. As these ontologies are based on formal logic, and they are machine readable format, it is possible to build such applications that are building their interfaces upon them. However the data structure itself is not sufficient to build GUI-s that allows to convenient digitalisation of the research processes. It is an additional problem and time consuming and tedious. The thesis aims to build such application framework that allows the definition of the input processes on a higher level logic the abstract from the implementation details. The system is supposed to accelerate the development and should be able to define.

Kurzfassung

Kurzfassung auf Deutsch

Contents

Abstract	II
Kurzfassung	III
List of Tables	X
1 Introduction	1
1.1 RDFBones Project	1
1.2 Goal of the thesis	1
1.3 Thesis outline	2
2 Preliminaries	3
2.1 Semantic Web	3
2.1.1 RDF	3
2.1.2 RDF Schema	5
2.1.3 OWL	6
2.1.4 SPARQL	8
2.2 Applied Ontologies	10
2.2.1 Foundational Model of Anatomy - <i>FMA</i>	10
2.2.2 Ontology for Biomedical Investigations - <i>OB</i> <i>I</i>	11
2.3 Web applications	12
2.3.1 Client-sever architecture	12
2.3.2 Data driven web applications	14
2.3.3 Applications with RDF Data	17

3	Problem Statement	21
3.1	Modeling anthropological research activity	21
3.1.1	Data on skeletal remains	21
3.1.2	Investigation process	23
3.1.3	Ontology Extensions	25
3.2	RDF Data input	27
3.2.1	Dynamic data entry forms	27
3.2.2	Adoption form elements to the ontology	28
3.2.3	Editing form data	30
3.3	Solution Scheme	33
3.3.1	Model for RDF data input processes	33
3.3.2	Software framework outline	34
4	Vocabulary for web application domain	35
4.1	Elements of the vocabulary	35
4.1.1	Data definitions	35
4.1.2	Form definition	37
4.2	Use-cases of the <i>RDFBones</i> project	38
4.2.1	Skeletal Inventories	38
4.2.2	Study Design Execution	41
5	Framework functionality	43
5.1	Main software modules and tasks	43
5.1.1	Descriptor data processor	43
5.1.2	Web interface generation and user action handling . .	43
5.1.3	Saving and querying the RDF data	43
5.2	Implementation	43
5.2.1	Server side	43
5.2.2	Client side	46
5.2.3	Data dependency	50
5.2.4	Form validation	50
6	Evaluation	51
6.1	Expressivity of the framework	51
6.2	Reduced amount of code	51
6.3	Limitations of the system	51

7	Conclusion and future work	52
7.1	Summary of achievements	52
7.2	Future work	52
A	Glossary	54
B	Appendix	59
B.1	Something you need in the appendix	59

List of Figures

2.1	Main structure of the RDFS vocabulary	5
2.2	RDFS domain and range definition	6
2.3	RDFS domain and range definition	6
2.4	A subset of OWL vocabulary	7
2.5	OWL object properties	8
2.6	Properties for qualified cardinalities	8
2.7	Ontology structure for skeleton	11
2.8	Client server communication	12
2.9	HTML document is interpreted by the browser	13
2.10	Navigation through the web application	14
2.11	Data flow	15
2.12	Flow of information from DB to client	15
2.13	SQL query with parameter	16
2.14	Links to data items	16
2.15	Form layout and HTML document	16
2.16	Request with parameters	17
2.17	Example Java routine for data storage	17
2.18	Ontology and data in RDF	17
2.19	VIVO Profile page	18
2.20	Triples representing a new instance	19
2.21	Data input scheme by RDF	20
3.1	Ontology and triples of the skull	22
3.2	Bone segment in RDFBones	22

3.3	RDFBones as extension of OBI	23
3.4	Custom bone segment example	23
3.5	Applied subset of OBI ontology	24
3.6	Glabella and its expressions	24
3.7	Study design execution dataset	25
3.8	Ontology extension for skeletal inventories	25
3.9	Ontology extension for sex estimation	26
3.10	Multi dimensional form layout	27
3.11	Multi dimensional RDF dataset	28
3.12	Initial state of the form	29
3.13	Loading form data through AJAX	30
3.14	Two directions of the data flow	30
3.15	Multi dimensional data example	31
3.16	Vocabulary for RDF data input processes	33
3.17	Implementation workflow	34
4.1	Complete workflow of data input process	36
4.2	Variable types and their attributes	36
4.3	Triple types	37
4.4	Form definition	38
4.5	Input form for skeletal inventories	39
4.6	Skeletal inventory data triples	40
4.7	Complete data definition	40
4.8	Form layout definition	40
4.9	Input form for study design execution	41
4.10	Instance selector for existing bone segment	42
4.11	Complete data model	42
5.1	Overview of the mechanism on the server	44
5.2	UML Diagram of the classes for the form	44
5.3	Form descriptor JSON object	45
5.4	Graph UML	45
5.5	JSON vs graph model	46
5.6	Example problem	46
5.7	Elements of the simplified notation	47
5.8	Two restriction directions	47

5.9	Getting dependent data	48
5.10	Difference between the submissions and edit data on the form	48
7.1	Scheme of scheme	52

List of Tables

3.1	SPARQL result table	31
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Chapter 1

Introduction

1.1 RDFBones Project

The master thesis is written in the frame of the project called *RDFBones*. The goal of the project is to develop a data standard, as well a web application for documenting research activity related to biological anthropology. In anthropology each institute has different set of skeletal remains and they conduct their investigations differently, thus it is not possible to cover during the project period the problem of other institute as well. Therefore the core idea of the project is that to make possible the definition of custom processes through ontology extensions. This is only possible by means of RDF data, because the ontologies in OWL can be extended, and specific rules can be defined through restrictions. We are using Semantic Web application framework called *VIVO*, which is capable of editing not only the RDF data itself but the ontology as well. So the idea of the project that not only the data will be exchangeable but the extensions as well. The task of the application is to generate the data input forms based on the ontology extensions.

Web application so that they do not

1.2 Goal of the thesis

The data input forms and the data model is tightly coupled. Which means that the application has limited tasks of exercises. The The goal is to implement a such vocabulary that allows abstracts from the low implementation.

This would allow the developers the implementation of data input processes of people without programming experience. This

The vocabulary incorporates the definition of the input form layout, and the underlying data structure. Based on this compact definition the framework has to be able to generate the interfaces and perform the necessary data operations. T

1.3 Thesis outline

Chapter 2

Preliminaries

2.1 Semantic Web

2.1.1 RDF

In RDF, abbreviation for Resource Description Framework, the information of the web is represented by means of triples. Each triple consists of a subject, predicate and object. The set of triples constitute to an RDF graph, where the subject and object of the triples are the nodes, the predicates are the edges of the graph. An RDF triple is called as well statement, which asserts that there is a relationship defined by the predicate, between subject and the object. The subjects and the objects are RDF resources. A resource can be either an IRI (Internationalized Resource Identifier) or a literal or a blank node (discussed later). A resource represents any physical or abstract entity, while literals hold data values like string, integer or datum. Basically there are two types of triples, the one that links two entities to each other, and the other that links a literal to an entity. The former expresses a relationship between two entities, and the latter in turn assign an attribute to the entity. Common practice is to represent IRI with the notation prefix:suffix, where the prefix represents the namespace, and the expression means the concatenation of the namespace denoted by the prefix, with the suffix. This convention makes the RDF document more readable. The namespace of RDF is the `http://www.w3.org/1999/02/22-rdf-syntax-ns#`, whose prefix is in most cases "rdf". This is defined on the following way:


```
@prefix rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#>.
```

Literals are strings consisting of two elements. The first is the lexical form, which is the actual value, and the second is the data type IRI. RDF uses the data types from XML schema. The prefix (commonly xsd) is the following :

```
@prefix xsd: <http://www.w3.org/2001/XMLSchema#>.
```

So a literal value in RDF looks as follows:

```
"Some literal value"^^xsd:string
```

The RDF vocabulary provides some built-in IRIs. The two most important are, the `rdf:type` property, and the `rdf:Property` class. The meaning of the triples, where the predicate is the property `rdf:type` is that the subject IRI is the instance of the class denoted by the object. Therefore the following statement holds in the RDF vocabulary:

```
rdf:type rdf:type rdf:Property.
```

It is maybe confusing that an IRI appears in a triple as subject and predicate as well, but we will see by the RDFS vocabulary that it is inevitable to express rules of the language. To be able to represent information about a certain domain, it is necessary to extend the RDF vocabulary with properties and classes. The classes will be discussed in the next section, but here it is explained how custom properties can be defined. The namespace of the example is the following:

```
@prefix eg: <http://example.org#>.
```

The example dataset intends to express information about people, which university they attend and how old are them. To achieve this two properties are needed:

```
eg:attends rdf:type rdf:Property .  
eg:age rdf:type rdf:Property .
```

The actual data about a person:

```
eg:JanKlein eg:attends eg:UniversityOfFreiburg .  
eg:JanKlein eg:age "21"^^xsd:integer .
```

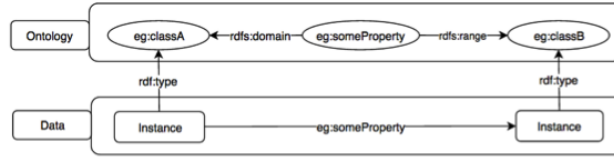



Figure 2.2: RDFS domain and range definition

the class `rdfs:Class`. The property `rdf:subPropertyOf` expresses the relationship between two properties. If property `P2` is sub property of `P1` and two instances are related by `P2` then they are related by `P1` as well. Its domain and range is the class `rdf:Property`. Now everything is given to define the ontology for the example of the previous section.

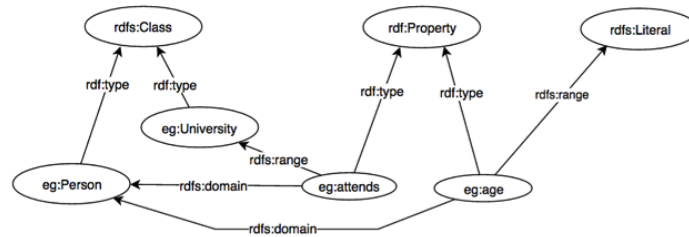


Figure 2.3: RDFS domain and range definition

2.1.3 OWL

OWL, abbreviation for Ontology Web Language is an extension of the RDFS vocabulary. OWL allows expressing additional constraints on the data, above the range and domain definitions. These constraints are called restrictions. Restrictions are conventionally expressed by blank nodes. Blank nodes do not have IRIs, but it is defined through the triples in which they participate as a subject. For example a restriction stating that the instances of the class `eg:FootballTeam` can build a triple through the `eg:hasPlayer` property only with the instances of `eg:FootballPlayer` class can be expressed the following way:

```
eg:FootballTeam rdfs:subClassOf [
  rdf:type      owl:Restriction ;
  owl:onProperty eg:hasPlayer ;
```

```
owl:allValuesFrom eg:FootballPlayer .
]
```

Listing 2.1: OWL restriction in N3 format

owl:Restriction is class and owl:onProperty and owl:allValuesFrom are properties. It can be seen that class, on which the restriction applies is the subclass of the restriction blank node. Furthermore OWL is capable of expressing qualified cardinality restriction. For example the statement that a basketball team has to have exactly five players, look as follows in OWL:

```
PREFIX rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#>
PREFIX eg: <http://example.org>

eg:BasketballTeam rdfs:subClassOf [
  rdf:type owl:Restriction ;
  owl:onProperty eg:hasPlayer ;
  owl:onClass eg:Player ;
  owl:qualifiedCardinality "5"^^xsd:nonnegativeInteger
] .
```

Listing 2.2: OWL restriction in N3 format

These two examples cover the thesis related features of OWL. The next image depicts the OWL vocabulary.

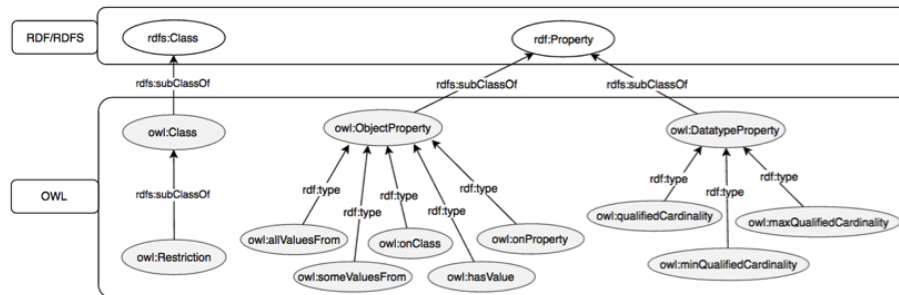


Figure 2.4: A subset of OWL vocabulary

There are two new class types are the owl:Class and the owl:Restriction. The rdf:Property has two subclasses, the owl:ObjectProperty and owl:DatatypeProperty. owl:ObjectProperty represent the properties that links instances to instances, and the owl:DatatypeProperty is those that link instances to literals. The

following two images shows the domain and range definitions of the OWL properties used to describe restrictions.

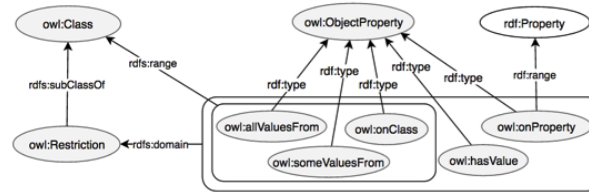


Figure 2.5: OWL object properties

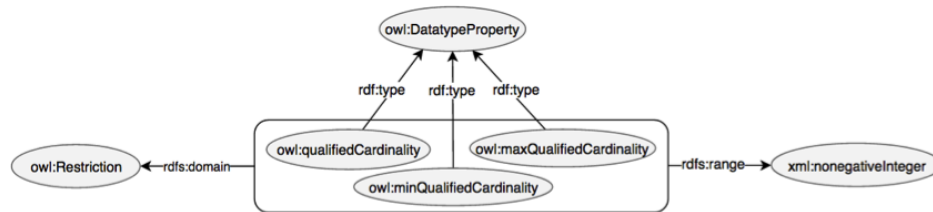


Figure 2.6: Properties for qualified cardinalities

2.1.4 SPARQL

SPARQL is a query language for querying data in RDF graphs. A SPARQL query is a definition of a graph pattern through variables and constants. The following example query returns all IRIs that represent a football player:

```
SELECT ?player
WHERE {
  ?player    rdf:type    eg:FootballPlayer .
}
```

Listing 2.3: SPARQL Query I.

In the example the query consist of only one triple. The subject is a variable and the predicate and the object are constant. Therefore the triple store looks all the triples and checks the predicate is `rdf:type` and the object is `eg:FootballPlayer`. It is well possible to not just ask the IRI of the players but further information by adding additional triples to the query in order to ask the name for example of the player:

```
SELECT ?player ?name
WHERE {
  ?player    rdf:type      eg:FootballPlayer .
  ?player    eg:name       ?name .
}
```

Listing 2.4: SPARQL Query II.

The result table in this case will contain two columns, one with the IRI of the person and one with their name. Important that it is as well possible to query blank nodes by introducing a variable for it. So if we want to list all the instances that are coming into question as player to a football team we can formulate the following query:

```
SELECT ?person ?name
WHERE {
  eg:FootballTeam rdfs:subClassOf ?restriction .
  ?restriction    rdf:type          owl:Restriction .
  ?restriction    owl:onProperty  eg:hasPlayer .
  ?restriction    owl:allValuesFrom ?playerType .
  ?player         rdf:type          ?playerType .
  ?player         eg:name           ?name .
}
```

Listing 2.5: SPARQL Query III.

2.2 Applied Ontologies

Ontologies are used to describe types, relationships and properties of objects of a certain domain. It is a common practice to use already defined ontologies rather than developing an own. The first reason is, that the development of an ontology is a complex and a tedious process, and requires a lot of resource. Secondly, it is reasonable to use standardized vocabularies, in order to make data from same domain but different sources inter-operable.

2.2.1 Foundational Model of Anatomy - *FMA*

The foundational Model of Anatomy ontology is an open source ontology written in OWL. FMA is a fundamental knowledge source for all biomedical domains, and it provides a declarative definition of concepts and relationships of the human body for knowledge based applications. It contains more than 70 000 classes, and 168 different relationships, and organize its entities into a deep subclass tree [4]. All types of anatomical entities are represented in FMA, like molecules, cells, tissues, muscles and of course bones. In our project we use only the subset of the FMA. The taken elements are the subclasses of the following two classes and the three properties:

- Classes

Subdivision of skeletal system - fma:85544

Bone Organ – fma:5018

- Properties

fma:systemic_part_of

fma:constitutional_part_of

fma:regional_part_of

The class *Bone Organ* is the superclass of all bones in the human skeleton. Each bone belongs to a skeletal subdivision and a skeletal subdivision can be a part of another skeletal subdivision. This relationship in both cases is expressed by the property *fma:systemic_part_of*. To define which bone organ belongs to which skeletal subdivision FMA uses OWL restrictions (see Figure 2.7). The properties *fma:constitutional_part_of* and *fma:regional_part_of* are discussed later.

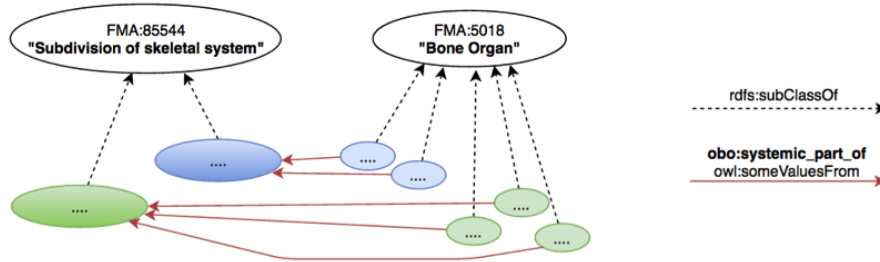


Figure 2.7: Ontology structure for skeleton

Finally the advantage of using the FMA ontology is that, if in the future further elements of the human body have to be addressed by the research processes, i.e. muscles, then these classes can be easily integrated to the currently applied subset.

2.2.2 Ontology for Biomedical Investigations - *OBI*

The aim of OBI ontology, is to provide the formal representation of the biomedical investigation in order to standardize the processes among different research communities. It is a result of a collaborative effort of several working groups, and it continuously evolving as new research methods are being developed. Its main function to describe the rules how biological and medical investigations have to be performed. OBI reuses terms from BFO *Basic Formal Ontology* IAO *Information Artifact Ontology* and OBO *Open Biological and Biomedical Ontologies*[3]. To define processes OBI uses the following three general classes:

- *Information Content Entity* - obo:IAO_0000030
- *Material Entity* - obo:BFO_0000040
- *Process* - obo:BFO_0000015

Information Content Entity represent results of a specific measurement, while Material Entity stands for the objects, on which the measurements have been performed. The Process could mean any kind of step within an investigation, from the planning, through execution till the conclusion.

- *Planning* - obo:OBI_0000339
- *Study Design Execution* - obo:OBI_0000471
- *Drawing a conclusion* - obo:OBI_0000338

In our project the following three properties are used:

- *has part* - obo:BFO_00000051
- *has specified input* - obo:OBI_00000293
- *has specified output* - obo:OBI_00000299

2.3 Web applications

This chapter contains practical information about how web applications work. In section 2.3.1 the basic mechanism of data driven applications are discussed, like navigation between page, data display and creation. Section ?? then focuses on the applications that are using semantic technologies, and addresses what kind of architectural changes that means.

2.3.1 Client-sever architecture

A web application is program that runs on a machine, which is accessible through the web. The machine is called server, because its main purpose is to server request that are coming from the web browser. Web browsers are as well programs, but they run on personal computers, tablets, etc, and they are capable of sending request through web to the servers. The response to these requests are HTML document, which can be displayed by the browser.

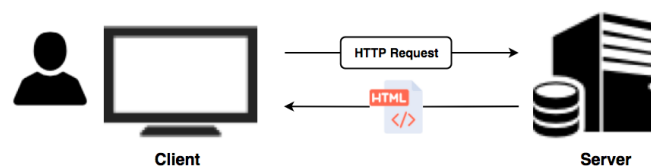


Figure 2.8: Client server communication

An HTML document contains definition of the elements of the pages, such as tables, buttons, etc. It contains as well so called CSS documents

(Cascading Style Sheet), which is responsible for the definition of the style of the elements. Moreover to make the web pages more interactive, JavaScript (JS) can be embedded to HTML as well.

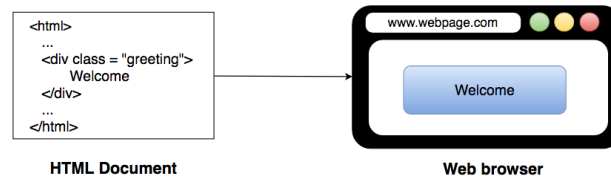


Figure 2.9: HTML document is interpreted by the browser

Initially web pages were static, which means that their only function was to show certain set of information. These applications usually web applications do not consist of one single page, but of several different pages. Like a web page for news, have normally a main page, and different sub pages for the particular topics. In order to navigate between the pages of the application, the HTML document contains links that trigger further HTTP requests. Links in HTML can be defined by means of the `<a/>` tag. The most important parameter of this tag is `href`, whose value contains the URL of the HTTP request. Let assume that an application's main page is accessible through the URL `http://newsPortal.com`. Common practice that sub-pages of the application can be called through various url-mappings, which means the main URL is extended with a keyword that denotes the page to be requested.

```
<a href="http://newsPortal.com/politics"> Politics </a>
<a href="http://newsPortal.com/sport"> Sport </a>
```

Listing 2.6: Example link definitions

If the user clicks on of these link (with the label 'Politics' and 'Sport') then these request will be sent to the news portal page. Each such request has to be served, differently to each mapping some routine has to be assigned. For example by Java web applications, the classes of the server that process the request are called servlets. On the next image it is shown, how the XML file defines, which class is responsible for the the mapping '/politics'.

```
<servlet-mapping>
  <url-pattern>/politics</url-pattern>
```

```
<servlet-class>servlets.PoliticsController</servlet-class>
</servlet-mapping>
```

Listing 2.7: Java servlet mapping definition

Then the responsibility of the class *servlets.PoliticsController* is to respond the corresponding HTML page for the client. Figure 2.10 show the main structure of the applications, where the rectangles on the client side represent the different pages of the application.

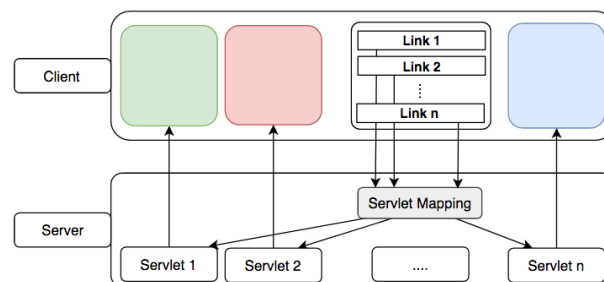


Figure 2.10: Navigation through the web application

2.3.2 Data driven web applications

This section aims to present the fundamentals of the web technologies that allows to build application for browsing and creating data. Modern web applications do not store the information in HTML documents. So the page loading process is not just the sending the HTML document, but a retrieval of a particular dataset, and the substitution into a web page. First of all the task of responding requires a query that retrieves that data from the database. By applications using relational data model, the tables and attributes are always modeled by classes of the used object oriented programming (OOP) language. So the data retrieval is the instantiation of the classes in scope.

Let assume that articles of a news portal is stored in a table with the attributes, id, type, title, summary and text. Then there has to be a class defined in the server code with the same attributes. To instantiate instances of the class, it is necessary to perform an SQL query.

The query results not only one instance of the News class, but a list (*List<News> newList*). To generate from this a HTML page that shows the

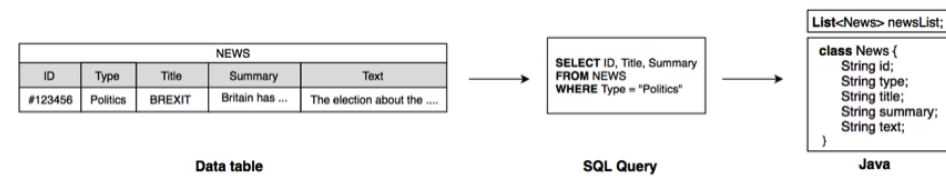


Figure 2.11: Data flow

articles, normally so called template engines are used. Templateing enables to define the HTML documents parametric, and passing them data, and they generate the result page automatically.

```
<#list newsList as news>
  <h3>  ${news.title} </h3>
  <p>  ${news.summary} </p>
  <a href = "http://newsPortal.com/wholeNews?id=${news.id}">
    Read more
  </a>
</#list>
```

Listing 2.8: Template file example

The template file is a description of how the data has to be converted into HTML document. It can be seen that it is possible for instance to declare a list on the input variable `newsList`. Then the template engine iterates through the `News` objects and by accessing its fields (title, summary, id) and generates the HTML for each element. So the complete flow of data from the database to the client looks as follows:

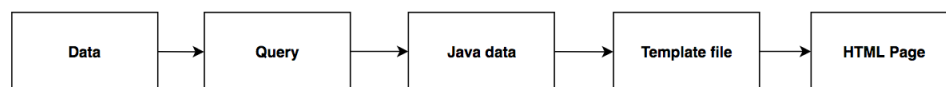


Figure 2.12: Flow of information from DB to client

The template shows only the summary of the article, but offers the following link:

```
http://newsPortal.com/wholeNews?id=${news.id}
```

The new feature is that after the url mapping there is a parameter *id*, and its value will be the database id of the web application. The idea is

that this link redirects to the page where the whole article can be seen. So there has to be a servlet class defined to the mapping `/wholeNews`, which to perform the following query where the `id` is the input.

```
SELECT Text  
FROM NEWS  
WHERE ID = ${id}
```

Figure 2.13: SQL query with parameter

Thus it is achieved that different links are programmed to get access not only to different other pages, but to specific data items.

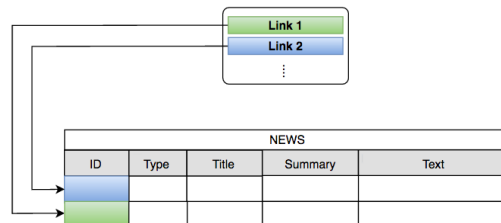


Figure 2.14: Links to data items

Web applications do not only just display existing data, but they allow the users to enter their new data. In HTML the element used for data input is called form. Form is a container, and it consists of particular form elements according to the data to be added.

Form layout

HTML Document

```
<html>
<form action="/webApp/newArticleController">
  Title <input type = "text" name = "title"> </br>
  Type <input type = "text" name = "type"> </br>
  Summary <input type = "text" name = "summary"> </br>
  Text <input type = "text" name = "text"> </br>
  <input type = "submit" value = "Submit">
</form>
</html>
```

Figure 2.15: Form layout and HTML document

Submitting the form to the server send an HTTP request with multiple parameters, where they are divided through the `&` character.

By the data entry creation the task of the controller is to get the values from the request and instantiate the class representing the data to be created.

```
"http://newsPortal.com/newArticleController?title=France won the EC&type=Sport&summary= ...."
```

Figure 2.16: Request with parameters

Then initialized class instance is passed to the database where the entered data will be persistently stored.

```
String title = request.getParameter("title");
....
News news = new News(title, type, summary, text);
DatabaseConnector.insert(news);
```

Figure 2.17: Example Java routine for data storage

2.3.3 Applications with RDF Data

This section aims to give an insight to web application that are based on RDF data. It will be covered what kind of requirements do the software have on the server side to create RDF data, and what is the difference between the RDF model based applications and the relational ones. The most important feature of RDF that the data scheme, namely the ontology is stored in RDF triples too, thus can be queried.

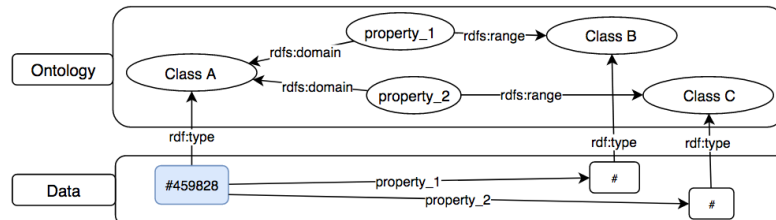


Figure 2.18: Ontology and data in RDF

Therefore it is possible to generate web pages that can adapt to the ontology. The following query demonstrates that how it is possible to get all the instances, which are connected to a particular instance (#459828).

```
SELECT ?property ?relatedInstance
WHERE {
    ?instance      rdf:type      ?class .
    ?property      rdfs:domain   ?class .
    ?instance      ?property     ?relatedInstance .
```

```

FILTER ( ?instance = #459828 ) .
}

```

Listing 2.9: Dynamic SPARQL query

The first two lines of the query defines the properties whose domain class is the type of the input instance, while the third asks for all triples with the possible properties. If the result of the query is then grouped based on properties, then the dataset can be displayed by a template using two lists. The outer list ceates fields for the properties, and the inner show all the instances with that property.

```

<#list properties as property>
  <#list property.dataSet as instance>
    <#list >
</#list>

```

Listing 2.10: Ontology adaptive template file

The VIVO framework applied in the RDFBones project generates the pages for instances this way.

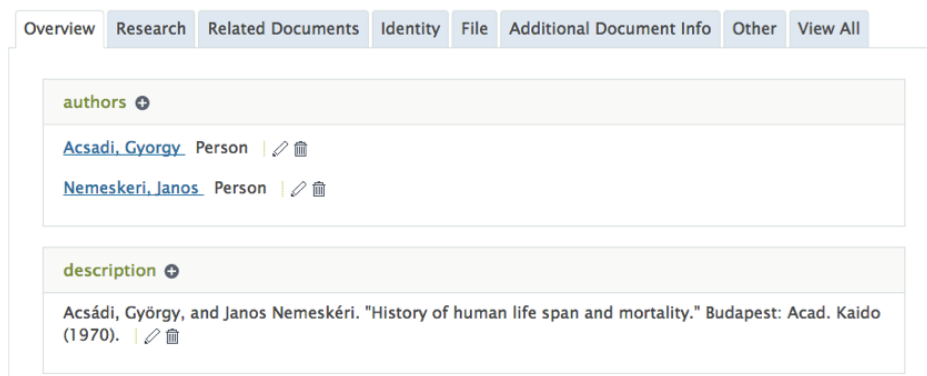


Figure 2.19: VIVO Profile page

Above the display of the existing data, RDF based applications are different as well in the data input mechanism. First of all, due to the fact that ontologies can contain thousand of class it is not an option to represent them all as classes of the server application language as well. It is time consuming and the system would loose its flexibility in the cases when new ontology subsets are supposed to be loaded. Therefore there are neither for

each type of the database an entry form with a unique controller servlet, but more generic approaches are used. To define what dataset has to be created, semantic web based applications simply define them as a set of triples, with variables like in SPARQL, just the data flows the other way around.

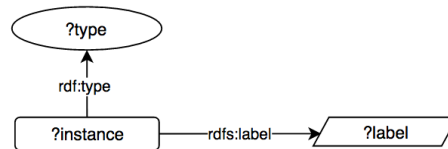


Figure 2.20: Triples representing a new instance

Figure 2.20 simple set of triples that have to be created by new data entry generation. The value of the variable `?instance` will be IRI that have not been used. This is an essential part of every triple store implementation that they provide unused uris for the server application for the new set of triples. The values of the *type* and label, are coming from the input form. The label is just like in the previous example, it is a string typed by the user it is stored as an attribute of the new entity. But the type of the instance is class IRI. The point is that the options of the selector field, from which the type value is coming, is filled with the results of a SPARQL query on the ontology. So for example if the entry form provide the possibility to create any type of processes from the OBI ontology then the before the form loading the following query has to be executed.

```

SELECT ?class ?label
WHERE {
  ?class      rdf:subClassOf    obo:OBI_0000339 .
  ?class      rdf:label        ?label .
}
  
```

Listing 2.11: SPARQL query for the input form

will be a new unused IRI of the triple store, while the `?type` variable come from the client from a selector field. The following image depicts this simple scheme of the data input process.

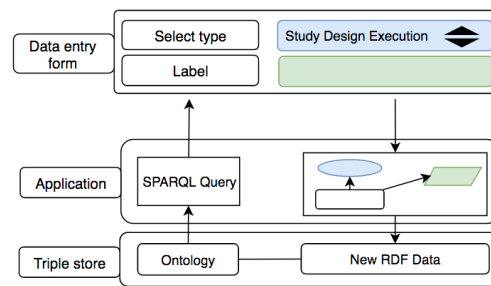


Figure 2.21: Data input scheme by RDF

Chapter 3

Problem Statement

This chapter is divided into three sections. As the application is highly dependent on the underlying data model, the first section is dedicated to the data scheme describing the investigations in scope. The second chapter in turn addresses the problem of web applications that allows the creation of RDF data explained in the first section. It covers the issues of both the client and server side implementation and their communication. Finally section 3.3 outlines the scheme of the solution proposed and implemented by the thesis work.

3.1 Modeling anthropological research activity

This section consist of three subsections. The first two (3.1.1 and 3.1.2) describes how the RDFBones ontology (developed during the project) integrates the *FMA* and *OB* ontologies for describing research processes related to anthropology. While the third section (3.1.3) discusses how can the core ontology be extended to define custom bone segments and processes.

3.1.1 Data on skeletal remains

We have seen in section 2.2.1 the base structure of the human skeleton. The most important point is that not only individual bones will be represented in the data we create, but the skeletal regions as well, like skull or vertebral column. The institute where these investigations are conducted posses mainly skeletal remains of skulls. The skull has the peculiarity that it does

not consists directly of bones organs, but from two sub skeletal divisions, and these two subdivisions contain the bone organs. Figure 3.1 shows the ontology subset for the skull and the data instances (each denoted with #). The red arrows denotes restrictions on the properties *fma:systemic_part_of*.

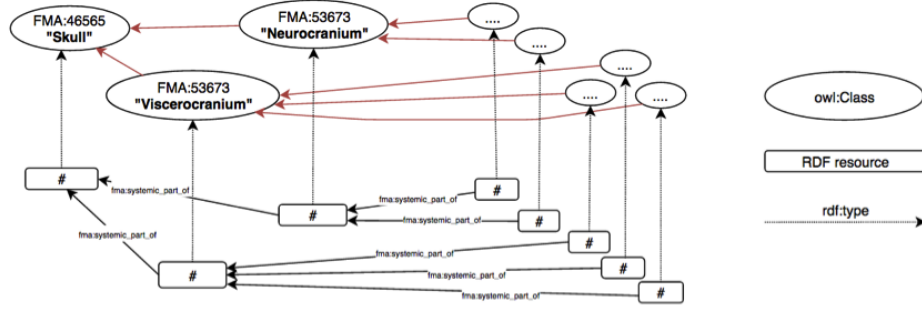


Figure 3.1: Ontology and triples of the skull

So we know how skull and its subdivisions and bone organs are represented by RDF, but there are processes where specific bone segments have to be addressed as well. Therefore RDFBones have the class *rdfbones:SegmentOfSkeletalElement*. This instances of this class is connected to the instances of the class *Bone Organ* with the property *fma:regional_part_of*.

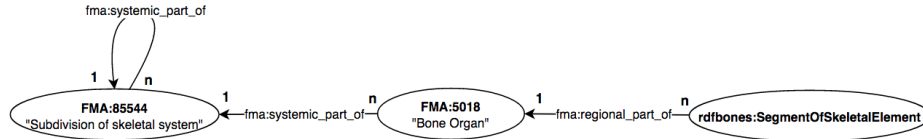


Figure 3.2: Bone segment in RDFBones

Furthermore these instances representing skeletal remains do not stay in the database individually. If a researcher takes a specific skull from the collection of the institute, it makes a so-called skeletal inventory, which records what bones segments are complete, partly present or missing. To store these information in the RDFBones ontology introduces three further classes (above the one for bone segment), which are all the subclasses of *OBI* classes.

Figure 3.4 illustrates the dataset through a simple problem. The upper left part of the figure shows that a specific bone is divided into three bone

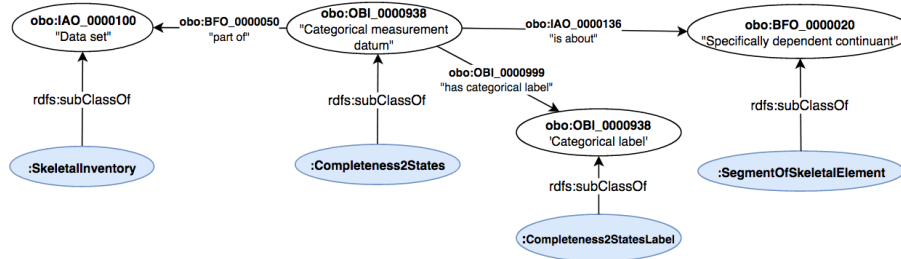


Figure 3.3: RDFBones as extension of OBI

segments, while the right shows the existing bone from which the data has to be stored. It can be seen the section I. is complete, section II. is just partly present and the III. is missing. The lower part of the figure then shows that the three segment are represented as subclass of the *:SegmentOfSkeletalElement* class (denoted with blue), and from the third type there are no instance have been created, while the first two are connected to the skeletal inventory and completeness label instances through *:Completeness2States* instances.

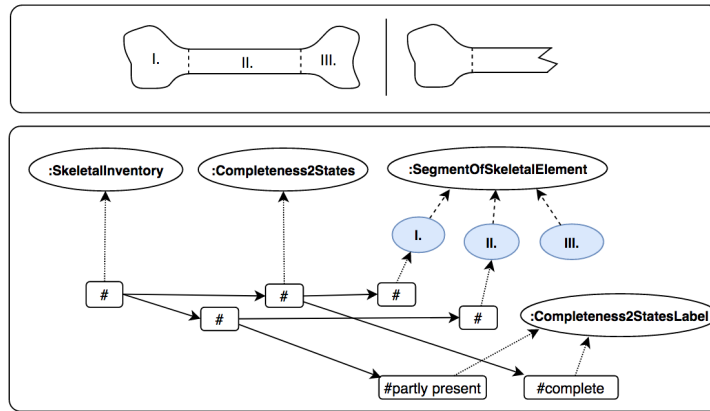


Figure 3.4: Custom bone segment example

This is the way how the information is stored about skeletal remains using the RDFBones ontology.

3.1.2 Investigation process

An investigation is done by execution of the a study design. It has three parts, the assay, the data transformation and the drawing of a conclusion.

3.1. Modeling anthropological research activity

From assays and data transformations there can be more in one execution, but there is only one conclusion. The inputs of the assays are always segments of skeletal elements, and their output is always a measurement datum, while the data transformation's input and output are both measurement datums.

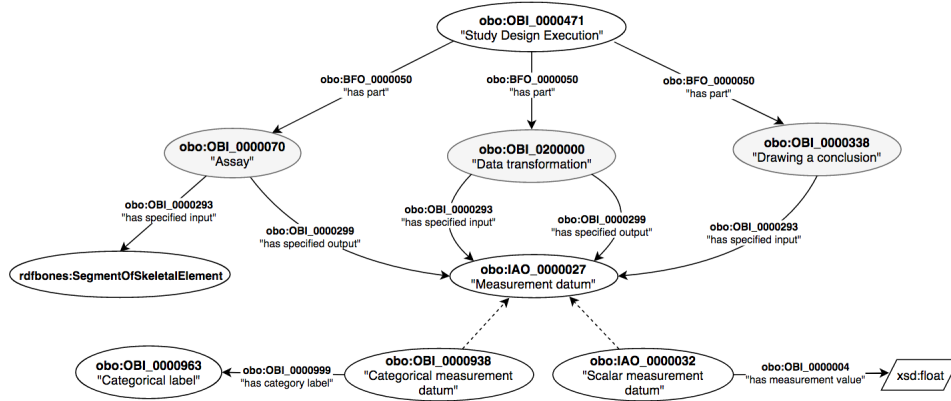


Figure 3.5: Applied subset of OBI ontology

To understand a bit more what this data model can be actually used, let us take the example of an investigation, whose goal is to determine if a taken skull belonged to a male or female. The basis is that the male and female skeleton has different peculiarities that can be quantified, how expressed they are. Figure 3.6 illustrates an example the token *Glabella*, which is on the *Nasal bone*, and its expressions.

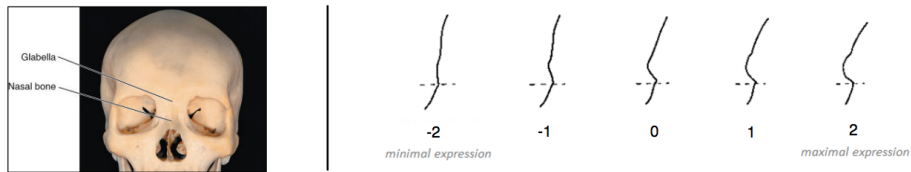


Figure 3.6: Glabella and its expressions

The larger the numbers for masculine and the lower are feminine expression. An assay in this case is an assignment of a scalar value to a bone segment. The investigation process does not take only one bone segment but several different ones, to reduce the possibility of the erroneous output. Fig-

ure 3.7 shows a dataset of a study design execution where the green arrows stand for *has specified input* and the blue ones for the *has specified output* predicates. The idea is simple, the output of the assays are aggregated, and if the output is smaller then zero then it was a male, otherwise a female.

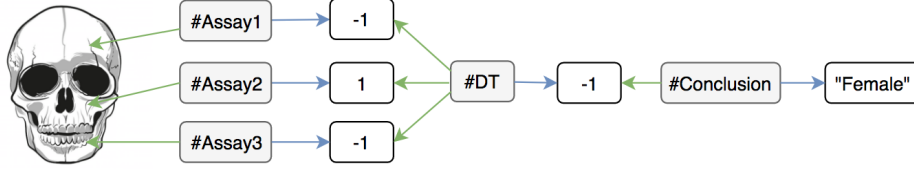


Figure 3.7: Study design execution dataset

Where the light grey boxes are the instances of *Assay*, *Data Transformation* and *Drawing conclusion* classes respectively.

3.1.3 Ontology Extensions

The previous two sections introduced the data scheme of the problems of the project. This part provides a more detailed explanation about how exactly these ontologies can be extended to tackle custom problems. By skeletal inventories *RDFBones* ontology allows to define custom bone segments of the bone organs in order to enable more fine-grained representation of the research activity. However by most of the cases it is sufficient to address the bone as a whole. For such cases *RDFBones* has the class *:PrimarySkeletalInventory* which encompasses all the entire bone organs. The definition of this inventory can be seen on the upper part of Figure 3.8.

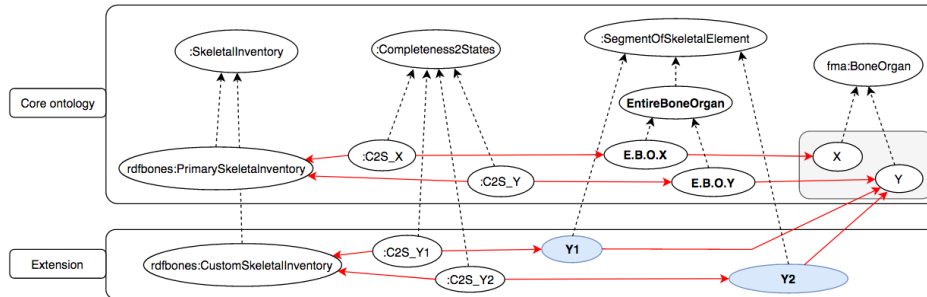


Figure 3.8: Ontology extension for skeletal inventories

The class *:EntireBoneOrgan* has as many subclass as many bone organs are there. The classes *X* and *Y* represents the set of all bone organs. Furthermore each of the entire bones are connected to the custom subclasses of *:Completeness2States*, to establish the connection to the primary skeletal inventory. The reason why there is not only one entire bone class for all bone organs, is that the input of the assays must be a bone segment, and it must be possible to address each of them individually. The lower part of Figure 3.8 shows a custom inventory definition, by new completeness and bone segment classes.

By the study design execution only the assay part will be covered, because the data transformation has exactly the same structure. As it was already mentioned through the example from Figure 3.7, each assay must address a bone segment an input, and a measurement datum as an output. To define a custom study design execution (Figure 3.9) at first a subclass is needed (*:SexExstimation*), and the different custom assays (denoted with blue) that are connected to each other again with restrictions. Then the outputs of the assays are custom scalar measurement datum classes.

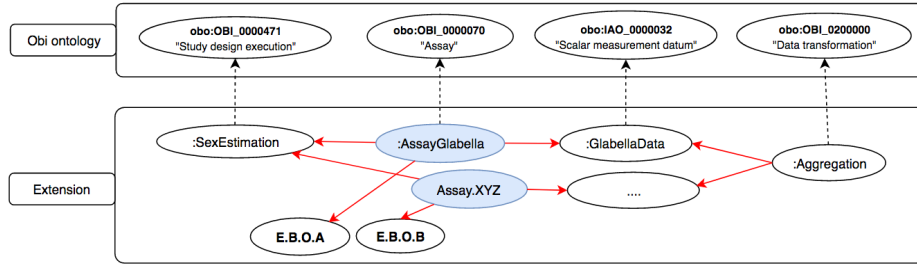


Figure 3.9: Ontology extension for sex estimation

3.2 RDF Data input

3.2.1 Dynamic data entry forms

In section ?? it was explained that the simplest data input process by RDF data is the substitution of the values coming from the client into a predefined set of RDF triples. In this case the interface is a static HTML form, the input data is a set of key-value pairs, while the RDF triples are defined on the server simply by a string. However in the previous section we have seen that the data models of the problems contain relationships with 1 to n cardinalities. This means that the data input process may contain, above a selection of the type, and setting some of the literal values of one new RDF instance, the dynamic adding of sub forms that represent further instances that are connected to the main instance with cardinality n.

Figure 3.10: Multi dimensional form layout

Figure 3.10 depicts a form which allows to add new sub forms that represent the sud subdivisions of the selected skeletal subdivision. Such functionality can be implemented by means of JavaScript routines embedded into the HTML document. The idea is that there is a main form, which consist of the same type of elements like the static forms, and one of its selector field is equipped with a button that initiates the loading of a new sub form. Then the sub form consists of again the standard elements, and maybe of further sub form adders. In the example the sub sud subdivision form contains an other sub form adder field for the bone organs.

The task of the JavaScript code running on the client is handle these click events and add the new form elements to the HTML document dynamically.

Moreover it has to access the input elements and insert the values into the form data object with defined keys. As there can be multiple sub forms added to the forms their data object stored by means of arrays. Listing 3.1 shows the JSON data (JavaScript Object Notation) of the form in Figure 3.10.

```
var formData = {
  subdivisionType : "fma:5018",
  subDivisionLabel : "Skull_5733FS2",
  systemic_parts : [
    {
      subsubdivisionType : "fma:45720",
      subSubdivisionLabel : "Neurocranium:_93KE43",
    }, { ... } ]
}
```

Listing 3.1: JSON object generated by the form

The server then in turn has to be prepared that a particular subgraph of the data model has to be created multiple times, and the values are arriving in arrays. The following image shows the data model of the entry form.

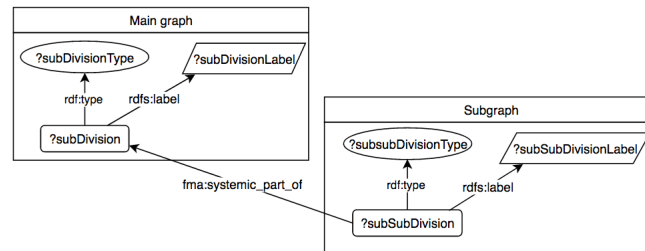


Figure 3.11: Multi dimensional RDF dataset

3.2.2 Adoption form elements to the ontology

The previous section showed the basics of forms for multi dimensional data input. An important issue which was just partly addressed in 2.3.3, is how the options of the particular selector fields are loaded. In order to load the options of the initial selector can template files be used, that gets the list of the classes coming from a SPARQL query on the ontology. In the example these are all skeletal subdivisions. But there are elements, for example the second selector field in Figure 3.17, whose values are dependent on the pre-

vious elements, which means their options must be loaded dynamically only after selection of its ancestor element.

Figure 3.12: Initial state of the form

Therefore their values are not part of the initial HTML document like possibly in the case of the independent skeletal subdivision class selector, but they have to be set as values of JS variables, so that they can be loaded after the selection. This can be achieved the same way by template file routines, that gets the set of sub subdivisions grouped by skeletal division.

```
var subSubDivisions = {
  skull : [{ neurocranium } ... { .. }],
  vertebralColumn : [{ cervical vertebra } ... { .. }],
  ...
}
```

Listing 3.2: Form option data structure in JSON

The data set on the listing 3.2 stores the lists of the sub subdivision with the corresponding keys (the keys are real IRIs in practice). Such data can be written to JS with template routines by means of two lists, like in listing 2.10. Then if the user selects the first field, JS can access based on the selected value the array to be loaded as options.

But as the example on Figure Figure 3.10 shows that after adding the sub subdivision the bone organs are supposed to be selected to. Therefore the client has to have the same data structure for bone organs as well grouped by the sub subdivision. The problem is that this data on the form can easily become too large, which may lead to performance issues. Additionally if the whole dataset is queried by the page load then it can be as well inefficient. Therefore it is common a practice that data of the form is loading dynamically through AJAX (Asynchronous JavaScript and XML). AJAX is technology that allows the client to communicate with the server without reloading the whole page. This makes the pages more interactive and as well

more efficient.

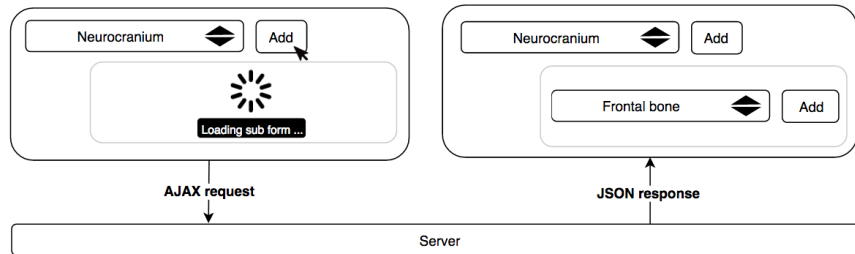


Figure 3.13: Loading form data through AJAX

The idea is that JS is programmed so that upon certain events, a particular request data is assembled and sent to the server. The response then in this case is not a whole HTML page but a dataset in XML or JSON format. After the response arrives, a JS routine continues its. This more elegant solution addresses further JS programming, and of course new server routines, that handles the requests, perform the queries with the incoming parameters, and respond the results.

3.2.3 Editing form data

The dataset created by the forms have be browsed and edited as well. This means that the application has to be able to restore the form to the same state as it submitted for initial data creation. This is the other direction of the data flow, because here the existing data will be retrieved by SPARQL, whose result comes into a JSON object, based on which the form can reset the its state.

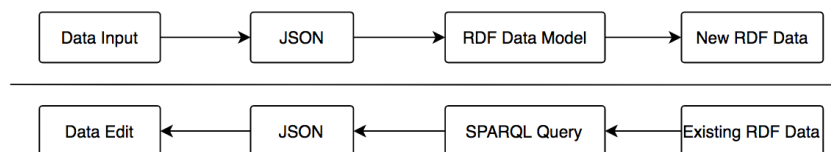


Figure 3.14: Two directions of the data flow

Important issue regarding the server implementation is that in such multi dimensional dataset, it is not sufficient to perform only one query for the whole form data. For example there can be more elaborate assays that have

?boneSegment	?measurementDatum
BS1	MD1
BS1	MD2
BS2	MD1
BS2	MD2

Table 3.1: SPARQL result table

multiple inputs and output (i.e Figure 3.15).

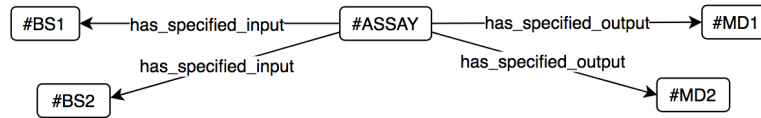


Figure 3.15: Multi dimensional data example

This means if the following query is executed on the dataset, results such a table ((Table 3.1)) that is hard to process.

```

SELECT ?boneSegment ?measurementDatum
WHERE {
  ?assay      :has_specified_input      ?boneSegment .
  ?assay      :has_specified_output     ?measurementDatum .
  FILTER ( ?assay = ${inputParameter} )
}

```

Listing 3.3: SPARQL query for the form data

Therefore the data object of the form has to be retrieved gradually, by dividing the data model graph by the predicates, whose cardinality is larger than one. The process have to start with the main graph, and the results of the sub graph queries then has to be stored in arrays of the form data JSON object. The task of the client is to iterate on the arrays and reload the sub forms. Important part of the form reloading is that, not only the values of the literal fields have to be restored, but the selector field options as well. So that for example if the existing dataset contains a skeletal subdivision, it is necessary to load the possible bone organs into the selector, so that the user can add additional bone segments conveniently. Finally if a value of a

literal field has been edited by the user, or new sub forms has to be added or removed, the saving operation should be done immediately through AJAX. To achieve this JS has to assign events to each literal fields and sub form container, and be able to send the appropriate request to the server.

3.3 Solution Scheme

3.3.1 Model for RDF data input processes

The task of the implementation of a data input forms addresses two main questions. Firstly is what dataset has to be created, and secondly how this dataset is represented on the form. The vocabulary which is capable of defining the whole input process, has to be able to answer both these question. Therefore the model is divided into two sections, to the data definition and the form definition. The data definition consist of a set of RDF nodes and triples, while the interface definition consist of the forms and the data input fields.

Nevertheless we have the previous chapters we have seen that the scheme of the ontology plays an important role of the data entry form process. Namely if a set restriction statement assign multiple classes to one, like i.e. bone organs to a skeletal subdivisions, then the web user interface has to be adapted to this scheme, and it has to of the possibility to add sub forms dynamically. So to make the vocabulary complete is it inevitable to expresses what triples may be added multiple times and of which form elements adds further sub forms.

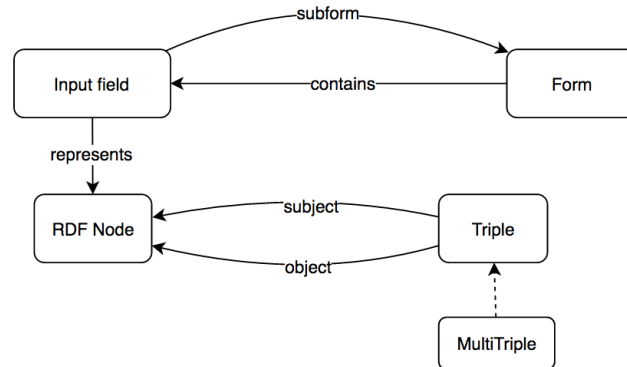


Figure 3.16: Vocabulary for RDF data input processes

Figure Figure 3.16 shows the main elements and relationships of the vocabulary. It can be seen that the connection between the data and the form definition is established through the represents predicate that connects form elements to RDF nodes. Moreover the MultiTriple is a subclass of the

triple, and it that allows to expresses the scheme of the ontology subset, for which the RDF data will be created.

3.3.2 Software framework outline

The above introduced vocabulary is represented currently by means of Java classes. Thus the definition of the problems is the creation of Java objects from the classes.

```
//Data definition
List<Triple> triples = new ArrayList<Triple>();
RDFNode subdivision = new RDFNode("subdivision");
RDFNode boneOrgan = new RDFNode("boneOrgan");
triples.add(new MultiTriple(boneOrgan, "fma:systemic_part_of",
    subdivision));
...
//Form definition
Form form = new Form();
form.addElement(new LiteralField("label"));
...
//FormConfiguration
FormConfig skeletalInventory = new FormConfig(form, triples);
```

Listing 3.4: Input process definition in Java

The idea is that framework operates completely upon this definition. The first task is to process the definition and generate JSON objects that describes the forms layout, and further Java objects for the data operations.

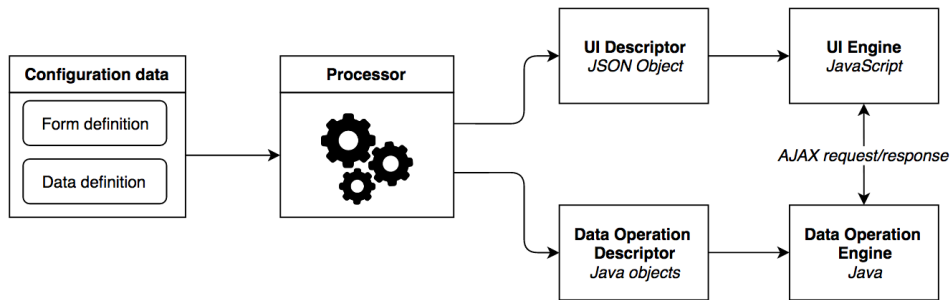


Figure 3.17: Implementation workflow

Chapter 4

Vocabulary for web application domain

4.1 Elements of the vocabulary

4.1.1 Data definitions

To understand the necessity of certain elements of the vocabulary, further details of web the applications have to be explained. In VIVO, the display of the existing and the creation of the new data happens in individual pages. The display is done by so-called profile pages, that show the information about one particular instance. As it was in section 2.3.3, the information is grouped by predicates, and each predicate field contains a link that can call the data input pages. The link contains three parameters, *subjectUri*, *predicateUri* and the *rangeUri*. The *subjectUri* hold the value of the instance on whose profile the link is, the *predicateUri* is for the predicate, with which the new dataset is connected to the subject, and the *rangeUri* is an optional paramater.

Figure 4.4 shows the workflow of a data entry process. On the right profile page of a skeletal inventory can be seen, which lists the added skeletal elements. The profile page is configured that the *rangeUri* parameter holds in the links the URI of the classes of skull and vertebral column respectively. These parameters have to be considered by the form loading because they influences the options of the first selector which let the user add the skele-

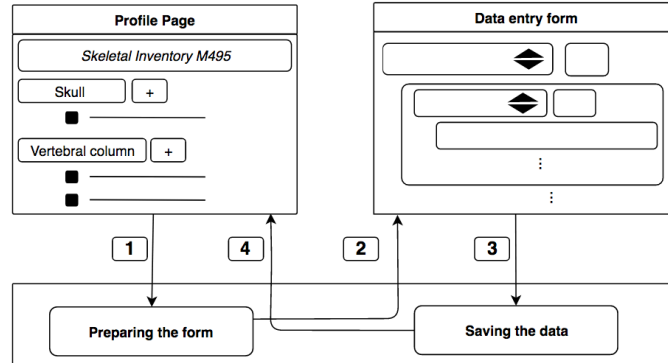


Figure 4.1: Complete workflow of data input process

tal sub sub division. So therefore it necessary to introduce a flag into the vocabulary that sign if a variable is coming with the HTTP request for the form loading, or with the JSON object after the submission.

Figure 4.3 shows all the nodes types and their attributes. Above the *mainInput* boolean flag, there is the variable name, which is required, and the constant value in case. There are three types of variable, the class, resource and literal. The literal variable itself denotes a string value and it has several subclasses for the other primitive types.

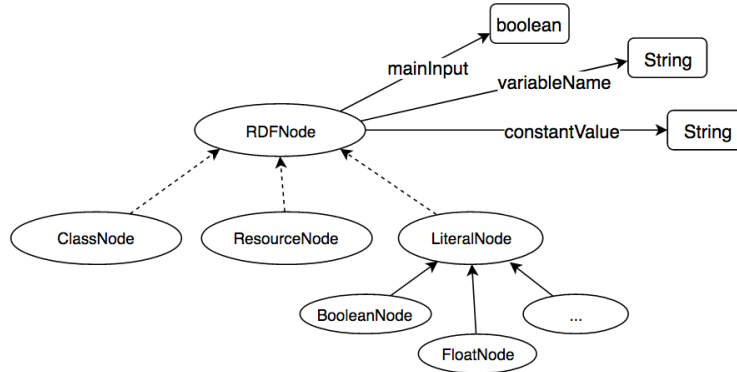


Figure 4.2: Variable types and their attributes

The other, more important is the modeling of the triples in the dataset. The vocabulary for triples has the purpose to express different constraint on the data scheme as well. Above the class *Triple* and *MultiTriple*, which were addressed in the end of the last chapter, there are two types of re-

striction triples. One for the classes and one for the instances. As we have seen in the description of the OWL ontologies, there are different types of restrictions can be defined. For this reason it should be possible to allow the definition of which restriction is used by the ontology, upon the entry form should operate. This can be expressed by the three boolean, types of the *classRestrictionTriple*. Moreover the greedy boolean flag means that the SPARQL query that queries the ontology has to return not only the result class but their superclasses too. Finally the instance restriction triples as the name indicates, expresses constraints between instances on the form. The examples in the next section will make the usage of the vocabulary more clear.

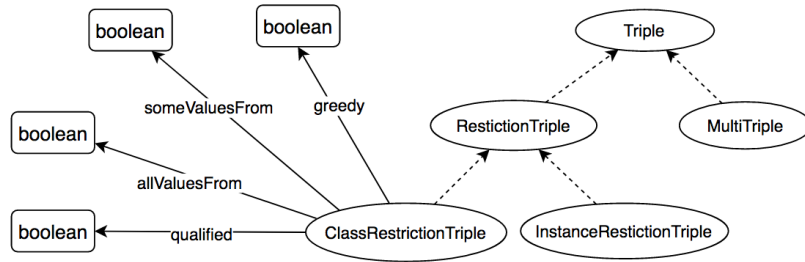


Figure 4.3: Triple types

4.1.2 Form definition

The class *Form* acts like a container for the form elements. There are two main types of form elements, the literal field and the selector. The literal field do not have further subclasses because its type, is defined through the type of the variable it represents. It would be a sort of over definition. This is the simplest case where the form adopts to the data model.

The selector can refer both instances and classes. If it represent an instance, so it let the selection of an existing instance, then it is possible to define an *InstanceViewer*. This feature allows to define a table with several columns. The utility is that the application can show more information about an instance than only the label in the selector field. Each column has a title and a number, and they refer to as well *RDFNodes*, whose values they show in the entries.

The class *SubformAdder* is the subclass of the selector, and has a relationship to the form with the predicate sub form. With this connection it is possible to define the sub form as new form instance. Moreover it has boolean flag, that allow to define a button add all should appear on, which adds all the possible subforms. This feature is useful be the skeletal subdivisions that contains much bones.

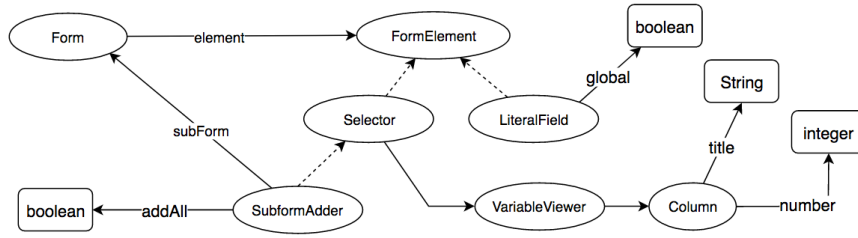


Figure 4.4: Form definition

4.2 Use-cases of the *RDFBones* project

The aim of this section is to show how it is possible to solve different problems with designed vocabulary. It covers the two main tasks of the project, the skeletal inventories and the study design execution. These two examples are sufficient to show the utility of the particular elements in the vocabulary and exemplify how their assembly can lead to a compact definition of complex web application problems.

4.2.1 Skeletal Inventories

Skeletal inventories were already addressed in section 3.3.1. Their goal is to define what kind of skeletal elements are present. In this part the creation of the primary skeletal inventories are discussed in more detail. This use-case addresses some additional challenges of the software, that were not mentioned yet. The explanation starts with the illustration of the implemented interface, to show what problem the high level logic has to define. As it was mentioned in figure 4.4, the entry forms can have inputs, which in this case the input is the class URI of the skeletal subdivision that has to be added with it sub subdivisions and bone organs. Therefore the first element of the

entry form is the selector of the sub subdivision.

Skull

Skeletal Regions **Viscerocranium**

Viscerocranium

BoneOrgans **Left temporal bone**

Left temporal bone **complete**

Neurocranium

BoneOrgans **Right parietal bone**

Right parietal bone **partly present**

Figure 4.5: Input form for skeletal inventories

Next to the selector the buttons *Add* and *Add all* can be seen, that let the user add the sub forms. Figure 4.5 shows the layout, when two sub sudvision were added, to each of them, a bone organ. The bone organ selector works exactly the same as the sub sudvision selector, but the selector for the completeness state is simple selector, without a sub form.

Figure 4.5 shows the triple scheme representing the skeletal inventories, where the nodes are representing variables. All the rectangle are representing instances. All of them will be newly created instead of the subjectUri, which comes as a main input, and the arrows with double line depicts the multi triples. Important to note that between the variable *boneOrgan* and *boneSegment* there is only a single triple (*fma:regional_part_of*) because in this use-case is simplified version and only entire bone segments will be added.

Above the instances to be created the classes have to be represented in the model too, because they can be the subjects of the class restriction triples. Figure Figure 4.7 is depicts the complete data model of the problem.

For the better readability the predicates are not denoted, but their value can be found in figure Figure 4.6. Each instance (rectangles) is connected to the class variable (ellipses). The red dotted arrows indicates restriction statements. The large three rectangles, that encompass set of triples are the graph, which are connecting to each other by means of multi triples. This is

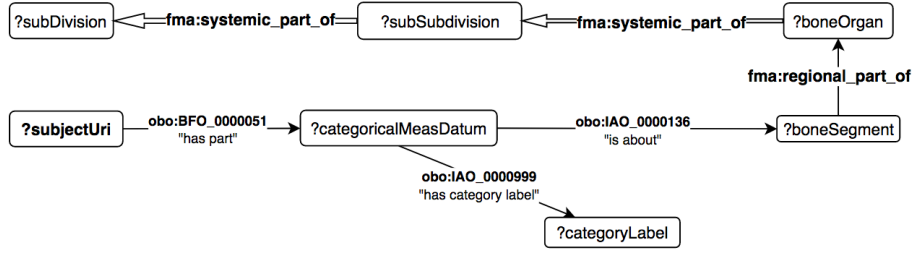


Figure 4.6: Skeletal inventory data triples

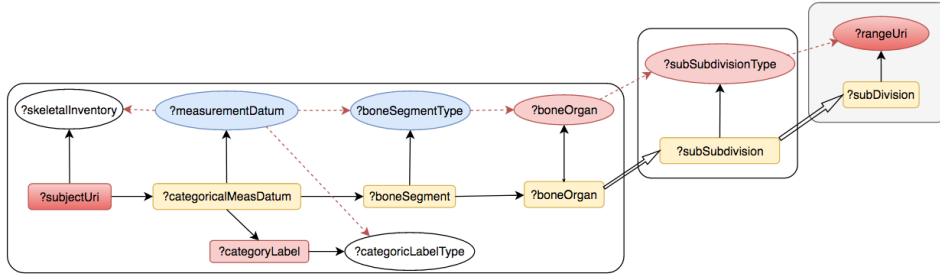


Figure 4.7: Complete data definition

the structure which is followed by the form as well.

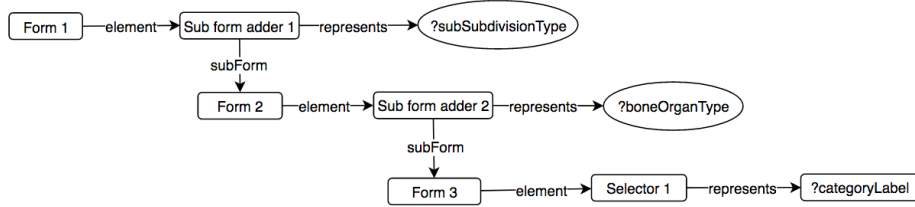


Figure 4.8: Form layout definition

Above the auxiliary rectangles for the graph, the nodes are colored to indicate their role in the process. The information that the colors hold is not defined in the vocabulary but it can be inferred from the whole data and form model. The first rule is that the main input nodes (*subjectUri* and *rangeUri*) are denoted with red, while the variables appear on the interface are light red. Based on that information it is already possible to determine to which instances it is required to assign an unused URI. Those instances are denoted with yellow. Furthermore, there are two classes in the data model that do not

appear on the interface, but their values can be evaluated through SPARQL queries. These are denoted with blue. And the classes with without color, do not appear on the final dataset, but they indicates constraint on the existing instances. Finally Figure 4.8 display the configuration data describing the form structure.

4.2.2 Study Design Execution

The entry form for study design execution has as well the hierarchical layout like the one for skeletal inventories, but there are additionally two elements on the main form. The first is the selector from skeletal inventories. It plays a role by the selection of the bone organs as input for the assays. To each assay a set bone segment types is defined in the extensions, that can be can be assigned to them as input. These bone on this form are not created newly but existing ones are selected, that were already added in the frame of the skeletal inventory data input. However there can be a large amount bone segments stored in the system, and thus the search is facilitated by showing only the ones that belong to the preselected skeletal inventory. The second is a global label field, whose value will be the label of all newly created instances.

Study Design Execution

Skeletal Inventory: Dry Bone Skeletal Inventory

Label: M-342

Assays: Assay.Glabella [Add] [Add all]

Assay.Glabella

Bone Segment: [Select]

Measurement Type: SexScore.Glabella [Add]

SexScore.Glabella: Indifferent [Close]

[Submit] [Cancel]

Figure 4.9: Input form for study design execution

Moreover the bone segment selector is not just a HTML selector input field, but a floating window implemented by JavaScript that allows the convenient browsing (Figure 4.10). It has two advantage with respect to

the conventional selector. Firstly it allows to display additional information about the instances above their labels, like their types or longer descriptions. Secondly it does not loads the form layout with additional subform for the selected instances, which by large amount assays and measurement datums is an important aspect.

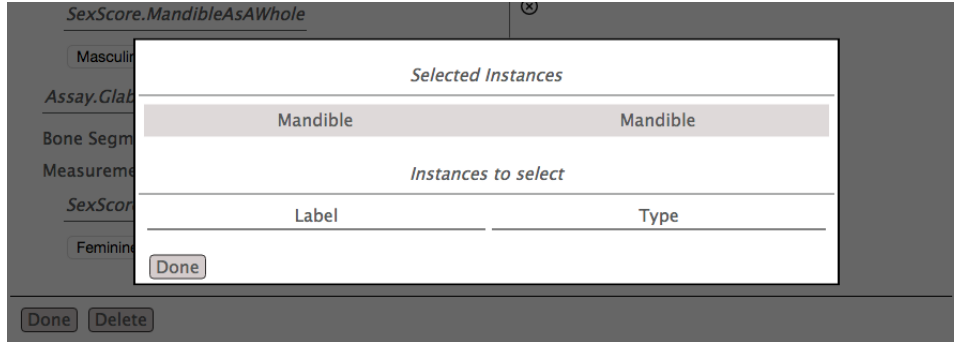


Figure 4.10: Instance selector for existing bone segment

On Figure 4.10 can be seen that there are two sections, one for the selected instances, and one for the instances to select.

The complete data structure of the form can be seen on the following image.

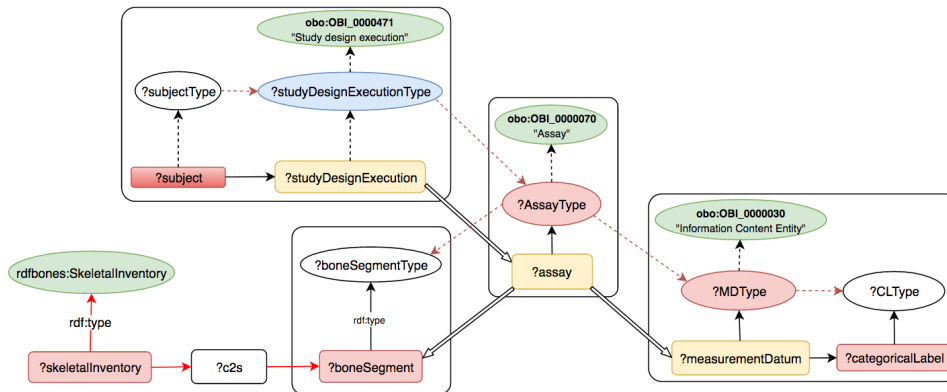


Figure 4.11: Complete data model

Chapter 5

Framework functionality

5.1 Main software modules and tasks

5.1.1 Descriptor data processor

5.1.2 Web interface generation and user action handling

5.1.3 Saving and querying the RDF data

5.2 Implementation

5.2.1 Server side

- This section introduces how application on the server is able to operate based on data queried from the configuration dataset

Overview

- The following image shows the most important steps of the data input process

Form representation in Java

- The literal fields asks for the type of the variable it represents and the type of the descriptor will be based on this literal field.

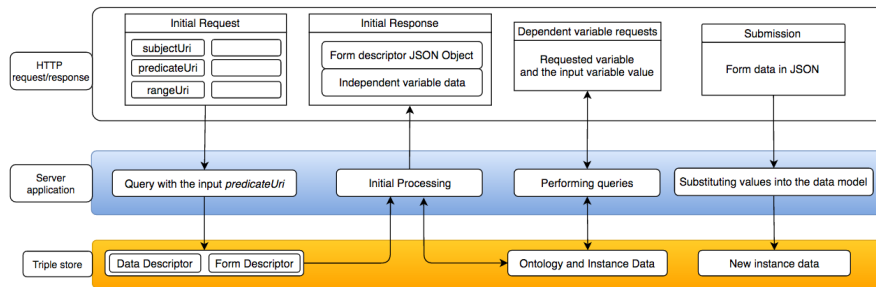


Figure 5.1: Overview of the mechanism on the server

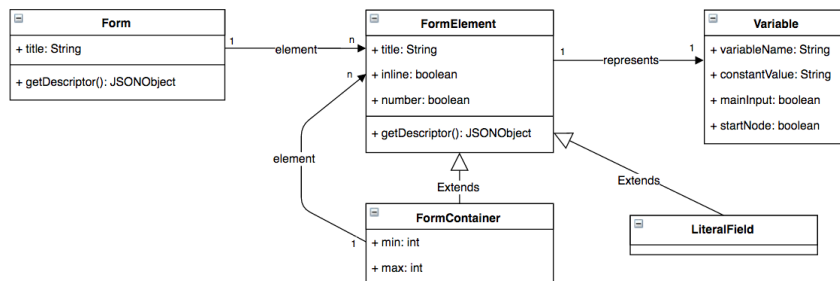


Figure 5.2: UML Diagram of the classes for the form

Data model in Java

- Querying the configuration triples regarding statements
- Processing into the tree structured graph model

Validation:

- Data dependencies can be over graphs
- But graphs can be connected only through multi statements
- The graph has to be a tree. There are no use cases right now where any loop would be required
- Data saving mechanism explanation
- Data retrieval mechanism explanation

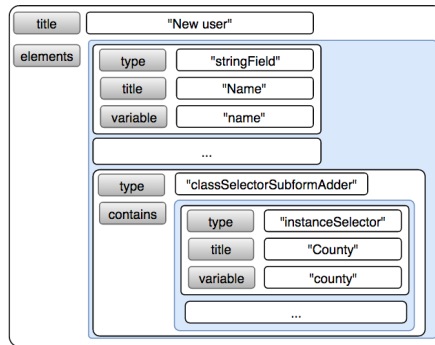


Figure 5.3: Form descriptor JSON object

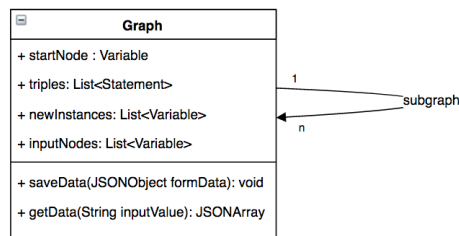


Figure 5.4: Graph UML

Data Dependencies

- The data dependencies are important only for the form.
- Everything starts with the form descriptor.
- There are cases where from the main form no element appears on the form.
- A have to bring examples to some problems that illustrate the problem.
- Here comes at first the ontology awareness into question....

where,

- Post processing for different restriction types

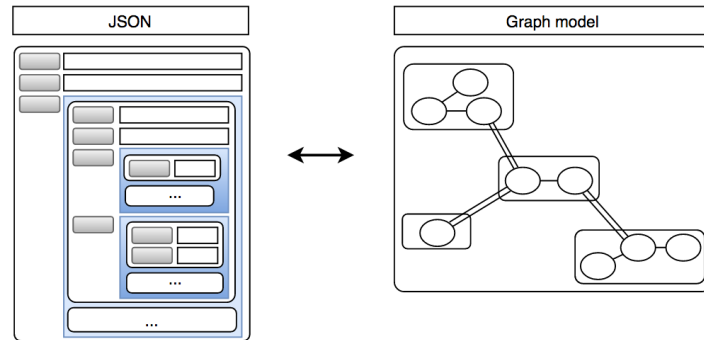


Figure 5.5: JSON vs graph model

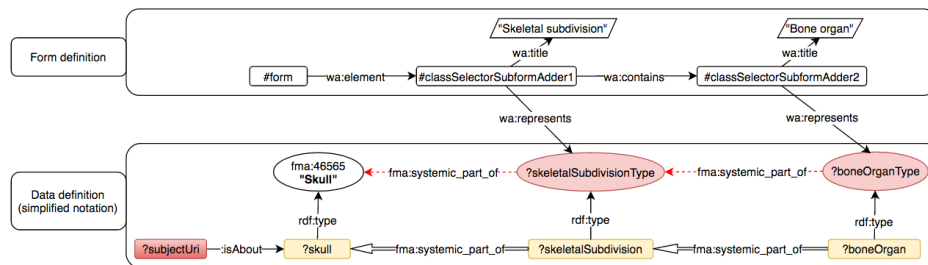


Figure 5.6: Example problem

Editing and deleting data

- This feature was not introduced on the overview image but it is really important.

5.2.2 Client side

- In this chapter it is discussed how the forms are implemented in JavaScript
- Each subsection contains code examples that gradually introduce the functionalities
- Codes are mostly simplified to facilitate the understanding

Object oriented JavaScript

- The task of code in on the interface is to operate based on a configuration data dynamically.

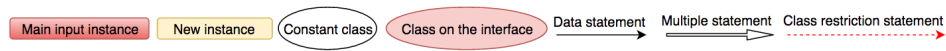


Figure 5.7: Elements of the simplified notation



Figure 5.8: Two restriction directions

- There two subtasks, the generation of the UI elements (i.e input fields, buttons, etc.) and manage the data input and display on the form
- To solve these problem, an object oriented approach is applied
- This means that there are classes that handles both the UI and the data related tasks
- See an example for the class definition in JS

```
class StringField {
  constructor (...) {
    this.container = $("<div/>")
    ...
  }
  someMethod () { ... }
}
```

Listing 5.1: JavaScript class

- Each form elements are represented by such objects, and the form loading based on the descriptor runs by the initialisation of these elements.

```
var formData = new Object ()
for (var i = 0, i < formElements.length; i++) {
  var descriptor = formElements[i]
  var element = null
  switch (descriptor.type) {
    case "stringField":
      element = new StringField (descriptor , formData)
      break;
    case "...":
```

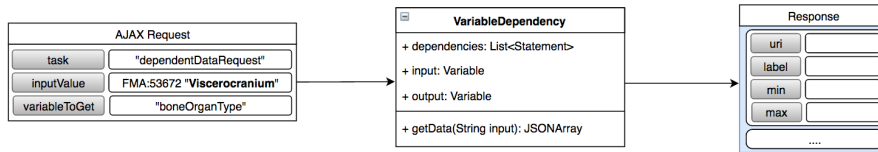


Figure 5.9: Getting dependent data

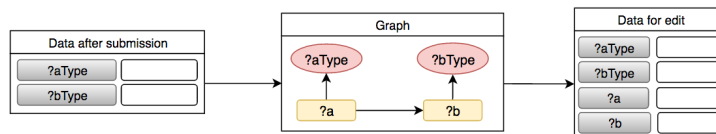


Figure 5.10: Difference between the submissions and edit data on the form

```

}
$( "#formContainer" ).append( element . container )
}

```

Listing 5.2: Form generation based on configuration data

- Explanation of the code ...
- This is the way how it is possible to generate interfaces

Handling data

- Above the element generation for the different forms it is necessary to handle of course the data based on the descriptor
- Each form element's descriptor contains a field called dataKey. The value of this field will be the key of data in the form data object.

```

class StringField {
  constructor(descriptor , formData){
    this.descriptor = descriptor
    this.formData = formData
    this.inputField = $( "<input/>" ).change( this . handler )
  }
}

```

```
handler() {  
  this.formData[this.descriptor.dataKey]=this.inputField.val()  
}  
}
```

Listing 5.3: Data saving

- The previous code illustrates how the form element object set the global form data field based on configuration data
- Further details about the code...
- Handling existing data
- It can be the case that form is loaded for editing. Then in this case the formData variable coming as input to the constructor contains the value for the dataKey?

```
class StringField {  
  constructor(descriptor , formData){  
    if(formData[descriptor.dataKey] != undefined){  
      this.editMode = true  
    }  
  }  
  
  handler() {  
    if(this.editMode){  
      var oldValue = this.formData[this.descriptor.dataKey]  
      var newValue = this.inputField.val()  
    }  
    AJAX.updateField(oldValue , newValue)  
  }  
}
```

Listing 5.4: Data saving

Sub form adders

- Descriptor of the sub form adder contains a field called subform
- Then the initialisation of the form happend through the Form class.

- This is used by the initial form loading as well

```
class SubformAdder {
  constructor(descriptor, formData){
    ...
    this.addButton = $("").text("Add").click(this.add)
    this.subFormDescriptor = this.descriptor.subForm
    this.formData[this.descriptor.dataKey] = []
  }

  add(){
    var subformDataObject = new Object()
    this.formData[this.descriptor.dataKey].
    push(subformDataObject)
    this.subFormContainer.append(
      new Form(this.subFormDescriptor, subformDataObject))
  }
}
```

Listing 5.5: Sub form adder routine

- Important for each subform a new JSON object is generated which is pushed to the array
- The same way the subform adder checks if the this.formData[this.descriptor.dataKey] contains already existing data and adds them if they are there

5.2.3 Data dependency

5.2.4 Form validation

Chapter 6

Evaluation

6.1 Expressivity of the framework

- Most of the problems of the project were solved by the higher level language
- Thus the code of the application could be significantly reduced
- This enables a more concise and readable documentation, which is a really important issue by such open-source projects.

6.2 Reduced amount of code

- As the configuration data is in the form of RDF triples as well, they could be added through web interfaces to accelerate the development.
- The most important advantage of the framework that the web application ontology can describe the input forms for itself.
- Another utility that in such case not only the data can be exchanged between institutes using RDF data, but the application descriptor as well.

6.3 Limitations of the system

Conclusion and future work

7.1 Summary of achievements

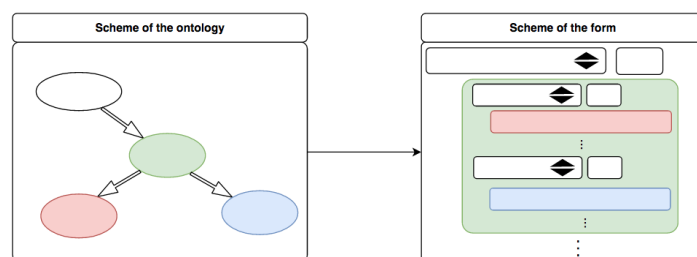


Figure 7.1: Scheme of scheme

7.2 Future work

Appendix A

Glossary

Just comment `\input{AppendixA-Glossary.tex}` in `Masterthesis.tex` if you don't need it!

Symbols

\$ US. dollars.

A

A Meaning of A.

B

C

D

E

F

G

H

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P

Q

R

S

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V

W

X

Appendix B

Appendix

B.1 Something you need in the appendix

Just comment `\input{AppendixB.tex}` in `Masterthesis.tex` if you don't need it!

Erklaerung

Hiermit erkläre ich, dass ich diese Abschlussarbeit selbständig verfasst habe, keine anderen als die angegebenen Quellen/Hilfsmittel verwendet habe und alle Stellen, die wörtlich oder sinngemäß aus veröffentlichten Schriften entnommen wurden, als solche kenntlich gemacht habe. Darüber hinaus erkläre ich, dass diese Abschlussarbeit nicht, auch nicht auszugsweise, bereits für eine andere Prüfung angefertigt wurde.

Ort, Datum

Unterschrift

Bibliography

- [1] OWL 2 web ontology language document overview. W3C recommendation, W3C, October 2009. <http://www.w3.org/TR/2009/REC-owl2-overview-20091027/>.
- [2] Marcelo Arenas, Claudio Gutierrez, and Jorge Pérez. On the semantics of sparql. In *Semantic Web Information Management*, pages 281–307. 2009.
- [3] Anita Bandrowski, Ryan Brinkman, Mathias Brochhausen, Matthew H. Brush, Bill Bug, Marcus C. Chibucos, Kevin Clancy, Mélanie Courtot, Dirk Derom, Michel Dumontier, Liju Fan, Jennifer Fostel, Gilberto Fragoso, Frank Gibson, Alejandra Gonzalez-Beltran, Melissa A. Haendel, Yongqun He, Mervi Heiskanen, Tina Hernandez-Boussard, Mark Jensen, Yu Lin, Allyson L. Lister, Phillip Lord, James Malone, Elisabetta Manduchi, Monnie McGee, Norman Morrison, James A. Overton, Helen Parkinson, Bjoern Peters, Philippe Rocca-Serra, Alan Ruttenberg, Susanna-Assunta Sansone, Richard H. Scheuermann, Daniel Schober, Barry Smith, Larisa N. Soldatova, Christian J. Stoeckert, Jr., Chris F. Taylor, Carlo Torniai, Jessica A. Turner, Randi Vita, Patricia L. Whetzel, and Jie Zheng. The ontology for biomedical investigations. *PLOS ONE*, 11(4):1–19, 04 2016.
- [4] Cornelius Rosse and José L.V. Mejino Jr. A reference ontology for biomedical informatics: the foundational model of anatomy. *Journal of Biomedical Informatics*, 36(6):478 – 500, 2003. Unified Medical Language System.