IGUANA Manual

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Contents

| 1 | Intr | oduction | 3 |
|--------------|------|-------------------------------------|----|
| 2 | Frai | nework/API | 4 |
| | 2.1 | Setting up | 4 |
| | 2.2 | Interfaces | 4 |
| | | 2.2.1 Testcase | 4 |
| | | 2.2.1.1 ResultSet | 6 |
| | | 2.2.1.2 Connection | 6 |
| | | 2.2.2 Clusterer | 7 |
| | | 2.2.3 ConverterI with RDFVocabulary | 7 |
| 3 | Ben | chmark | 9 |
| | 3.1 | In general | 9 |
| | 3.2 | Starting the benchmark | 9 |
| | 3.3 | Implementations | 10 |
| | | 3.3.1 PGNConverter | 10 |
| | | 3.3.2 SortedStructureClusterer | 10 |
| | | 3.3.3 BorderFlowQueryClusterer | 10 |
| | | 3.3.4 BorderFlowStructureClusterer | 11 |
| | | 3.3.5 Testcases | 12 |
| | | 3.3.5.1 UploadTestcase | 12 |
| | | 3.3.5.2 QueryTestcase | 12 |
| | | 3.3.5.3 StressTestcase | 14 |
| | 3.4 | Configuration | 15 |
| | | 3.4.1 Root node | 15 |
| | | 3.4.2 Data description | 15 |
| | | 3.4.3 Databases | 16 |
| | | 3.4.4 Log-clustering | 19 |
| | | 3.4.5 Benchmark | 19 |
| | | 3.4.5.1 Testcases | 21 |
| | | 3.4.6 Email notification | 22 |
| 4 | Fur | her information | 24 |
| | 4.1 | Links | 24 |
| | 4.2 | License | 24 |
| \mathbf{A} | The | MIT License (MIT) | 26 |

1 Introduction

The IGUANA Benchmark Framework (IGUANA) is both a program and an API framework for benchmarking SPARQL Endpoints.

On one side it provides the ability to benchmark different SPARQL endpoints (e.g. triplestores) with given or generated datasets and predefined test cases and on the other side it provides a framework and API to implement either a complete different benchmark or just parts of the benchmark (e.g. own test cases).

In the following the manual will describe first how to use IGUANA as a framework then how to use it as a benchmark and how to configure it.

In the last part there will be a list of further information (such as where to find the source code)

2 Framework/API

2.1 Setting up

To use IGUANA as an API for your own benchmark, please consider the javadoc.

To use IGUANA as a framework you don't have to change the code itself, just write a library (jar file) and use IGUANA as a library for your code, which is necessary to include. If you deploy your code as a library for IGUANA you don't need IGUANA as a library in your deployed code.

IGUANA will load your classes on runtime if they are in the lib folder by the IGUANA jar file.

2.2 Interfaces

To replace parts of the Benchmark with your code IGUANA use Class.forName(className) to load your classes on runtime. To achieve this and also give you the possibility for properties, the classes must implement the provided interfaces. There are three parts you can change like this and therefore three interfaces.

Write a class which implements one of these interfaces and you can use it by writing the class name into the configuration file (see also 3.4).

2.2.1 Testcase

To use your testcases in IGUANA you have to implement the interface *Test-case.java*. Therefore you have to implement the following functions:

Please be aware that ResultSet is a class provided by IGUANA (see also 2.2.1.1) and Connection is a class provided by an adjusted version of wookieConnection (see also 2.2.1.2 and 4.1)

| Function | parameters | return | Description |
|--------------------------------|------------------------------------|-----------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------|
| start | none | void | This starts the whole test- case after setting the prop- erties, current connection, dbname and percentage |
| $\operatorname{getResults}$ | none | ${\it Collection}{<}{\it ResultSet}{>}$ | This will return the results after starting the testcase |
| ${ m addCurrentResults}$ | Collection <resultset></resultset> | void | This will add the previous results for the specific percentage. Which means you'll get the results for all tested connections for the current percentage. |
| $\operatorname{setProperties}$ | Properties | void | If you want to give users a little bit more control over your testcase, you can use properties for parameters etc. |
| $\operatorname{setConnection}$ | Connection | void | This will set the connection to test |
| ${\bf set Current DBN ame}$ | String | void | This will give you the current id of the tested connection |
| ${\bf set Current Percent}$ | String | void | This will set the current percentage on which the connection will be tested |

2.2.1.1 ResultSet The ResultSet is an Iterator which gives you the opportunity to rapidly go through the current results and easily add results to it. Also you can save the ResultSet as a CSV file (the ResultSet ist simply a table with a header) and as a bar chart png file.

Example:

| header: | Connection | Query1 | Query2 | | QueryN |
|---------|-----------------------|--------|--------|---|--------|
| Table: | con1 | 200 | 230 | | 176 |
| | con2 | 400 | 120 | | 300 |
| | : | : | : | : | : |
| | conM | 234 | 1233 | | 89 |

If this is the header and table in your ResultSet and you have one more connection to test, you can simply add the row

$$con M1 \quad 123 \quad 123 \quad \dots \quad 123$$

by the function addRow.

For more information on the ResultSet please consider the javadoc.

- **2.2.1.2 Connection** The connection has several functions to give you an interface for SPARQL endpoints which is easy to use. The connection can. . .
 - ... upload a file either in the default graph or in a given graph.
 - ... execute a SPARQL/UPDATE query (as String)
 - ... execute a SPARQL query (as String) and get a java.sql.ResultSet
 - ... execute an UPDATE query (as String) and get a boolean if it worked.
 - ...set the upload type to POST or PUT
 - ...drop a given graph

2.2.2 Clusterer

To use your own log clustering in IGUANA you have to implement the interface *Clusterer.java*. Therefore you have to implement the following functions:

| Function | parameters | return | Description |
|----------------|--------------------------------------|--------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| cluster | String logPath String queriesFile | void | This is the main method of your clusterer. You'll get the path of the log files and the name of the output file (queriesFile) where the queries should be written in. |
| set Properties | Properties | void | If you want to give users a little bit more control over your log clustering process, you can use prop- erties for parameters etc. |

2.2.3 ConverterI with RDFVocabulary

To convert your data from non RDF files into RDF Files (RDF/XML, TURTLE, N-TRIPLE,...) you can write your own Converter simply by implementing the interface *ConverterI* and write a class which extends the class *RDFVocabulary*. The first one will simply convert the files while the latter one just setup the namespace etc. the converter should use. The *RDFVocabulary* will be initialized before the converter starts. Be aware that even though the *ConverterI* only needs the function to convert one file, the rest will be handled by the benchmark class itself.

${\bf Converter I}$

| Function | parameters | return | Description |
|---------------------------|---------------------------------------|---------|-----------------------------------------------------------------------------------------------------------------------|
| processToStream | String inputFile String outputFile | Boolean | This is the main method of the converter and will convert the given input file to the specified output file. |
| ${\bf set Output Format}$ | String outputFormat | void | This will simply just set the outputFormat. It will be set before the convert- ing process |

${\bf RDFVocabulary}$

| Function | parameters | return | Description |
|----------|---------------------------------------------------------------------------------------------------------------------|--------|-----------------------------------------------------------------------|
| init | String namespace String anchor String prefix String resourceURI String propertyPrefixName String resourcePrefixName | void | This will just set the user wanted resourceURI etc. to the converter. |

3 Benchmark

3.1 In general

The IGUANA as a benchmark can ...

- \bullet ... convert non RDF data into a RDF dataset and use this as the 100% dataset
- ...cluster and analyze query logs with different strategies and writes out query patterns to test
- ... generate smaller and greater datasets out of an initial 100% dataset.
- ...upload the datasets before testing them and measure the time if the UploadTestcase is specified in the configuration
- ...test different test cases on different connections (SPARQL endpoints) with different sized datasets
- ... save all results as csv files and if possible save them as barchart pgn files.
- ... send you an email if the benchmark succeeded with the results attached (in a zip folder) or aborted with the exception that causes the abortion and the by then calculated results.

3.2 Starting the benchmark

Before you start you must install IGUANA.

- 1. Download the latest version in a choosen folder and all the dependent libraries in a folder lib in the choosen folder.
- 2. If you want to use the data generation part of the benchmark with the specific type "coherence" you'll need lp_solve 5.5.2.0. You can download it here: http://sourceforge.net/projects/lpsolve/files/lpsolve/5.5.2.0/ otherwise skip to 3.
 - (a) The files you need are platform specific (replace "platform" with your platform. Be aware that you can have a x64 processor but your JVM is a 32-bit installation. If this is the case you need the 32-bit files):
 - i. lp_solve_5.5.2.0_exe_"platform" .zip or .tar.gz
 - ii. lp_solve_5.5.2.0_dev_"platform" .zip or .tar.gz
 - (b) Extract the libraries (or if you're lazy all the files) into a new folder called lpsolve_lib in your folder with the iguana.jar file
- 3. configure your config.xml file as you wish

To start the benchmark open a terminal and write the following:

```
>>java -cp "lib/*"
    de.uni_leipzig.iguana.benchmark.Benchmark config.xml
```

if you're using the data generator with the "coherence" type you'll need to start the benchmark as following:

Where config.xml is the configuration xml file you use (see also 3.4)

3.3 Implementations

As IGUANA has several parts which are exchangeable and provides the interfaces for this case there are some implementations yet.

3.3.1 PGNConverter

This converter simply converts PGN-files into rdf data.

You can use it with the following two class-names:

 $converter-class: \ de. uni_leipzig. informatik. swp13_sc. converter. PGNToRDFC onverterRanged \ rdf-vocabulary-class: \ de. uni_leipzig. informatik. swp13_sc. datamodel. rdf. ChessRDFV ocabulary-class: \ de. uni_leipzig. informatik. swp13_sc. datamodel. rdf. ChessRDFV ocabulary-class: \ de. uni_leipzig. informatik. swp13_sc. datamodel. rdf. ChessRDFV ocabulary-class. \ de. uni_leipzig. \ de. \ de. uni_leipzig. \ de. \ de$

3.3.2 SortedStructureClusterer

The SortedStructureClusterer clusters the queries in the log files through their structures and returns the frequentest queries for the frequentest structures. The following properties can be set:

| name | values/type | description | required? |
|-------------------|-------------|----------------------------------------------------------------------------------------------|------------|
| threshold-queries | Integer | The minimum number a query must occur in the log files to be part of the clustering | default=10 |
| threshold-structs | Integer | The minimum number a structure must occur in the log files to be part of the clustering | default=10 |

 $class: \ de.uni_leipzig.iguana.clustering.clusterer.SortedStructureClusterer$

${\bf 3.3.3}\quad {\bf Border Flow Query Cluster er}$

The BorderFlowQueryClusterer is a log cluster and analyzing implementation [3] using the Border Flow library (see also 4.1)

Even though it works it is not fully implemented in the current version! The following properties can be set:

| name | values/type | description | required? |
|-------------------|----------------------------------------------------|-------------------------------------------------------------------------------------|--------------|
| threshold-queries | Integer | The minimum number a query must occur in the log files to be part of the clustering | default=10 |
| delta | Integer | Parameter for threshold calculation [3] | default=2 |
| min-nodes | Integer | The minimum nodes a cluster must have to be part of the final query selection | default=3 |
| conn-threshold | Double [0:1] | Threshold parameter for Border Flow algorithm (see also 4.1) | default=0.8 |
| test-one | Boolean | Parameter for Border Flow algorithm (see also 4.1) | default=true |
| heuristic | Boolean | Parameter for Border Flow algorithm (see also 4.1) | default=true |
| caching | Boolean | Parameter for Border Flow algorithm (see also 4.1) | default=true |
| harden | hardensuperset hardensharedshed | Parameter for Border Flow algorithm (see also 4.1) | required |
| quality | qualitymeasurerelativeflow qualitymeasuresilhoutte | Parameter for Border Flow algorithm (see also 4.1) | required |

 $class: \ de.uni_leipzig.iguana.clustering.clusterer.BorderFlowQueryClusterer$

3.3.4 BorderFlowStructureClusterer

Same as the BorderFlowQueryClusterer but instead clustering the Queries with their similarities the Structures will be clustered.

 $class: \ de.uni_leipzig.iguana.clustering.clusterer.BorderFlowStructureClusterer$

3.3.5 Testcases

3.3.5.1 UploadTestcase This testcase is needed if you need to upload datasets to your connections!

It simply uploads the files for the datasets and measure the time it needs to take to upload the current file.

class: de.uni_leipzig.iguana.testcases.UploadTestcase

3.3.5.2 QueryTestcase The query testcase will test the query patterns for a given time-limit against one Connection.

The query testcase can handle a query pattern file with SPARQL queries and/or UPDATE queries in it or a query pattern file with SPARQL queries and you can set a path where your changesets are located. Changesets must have the following regex form: [0-9]{6}(added—removed).nt

Added means they will be inserted, removed means the will be deleted through the testcase.

The query testcase needs a file where the query patterns are written in (one query per one line). These query patterns can contain variables like %%var%% or %%var2%% etc. These variables will be turned into real data results described in the following lines.

The first step the query testcase will do (if the benchmark hasn't done it yet) is to convert the query patterns into queries. To achieve this the patterns will be converted to select queries with the variables of the pattern as SELECT variables. For example let the following query be the third query of the query patterns file:

SELECT ?v1 ?v2 WHERE {\"\var1\"\ <URI> ?v1. ?v1 ?v2 "2"}

As you can see var1 is a pattern variable and to test the connection with queries which have results, the pattern will be converted into the following query

SELECT ?var1 WHERE {?var1 <URI> ?v1. ?v1 ?v2 "2"}

Now all results (maximum the given limit) will be saved in the file 3.txt, by replacing %%var1%% with the results, in the directory QueryTestcase.

This will happen for every query pattern in the query patterns file.

The second step is to test all queries in a given order against the connection.

The given order can be set with the updateStrategy (In the following the UP-DATE query can be also changesets if you're using changesets instead of UP-DATE queries).

if you set this to fixed you can either set a parameter x or it will be calculated. This states that after every SPARQL (UPDATE) query x UPDATE (SPARQL) queries be tested (It'll be calculated if there are more UPDATE or more SPARQL queries).

if you set this to variation you can either set the parameter x or it will be calculated. Herby the x sets an interval like the following [1:x] if set or it will

calculate an accurate interval.

After every run of SPARQL (UPDATE) and the following UPDATE (SPARQL) the new x will be calculated (a random number in the given interval) If you're using live data (changesets) instead of UPDATE queries you can set the ldLinking parameter.

This parameter states in what order to execute the changesets.

| strategy | resulting order |
|--------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| insertsFirst | This states the every 'added' changeset (orderd by their number) will be before the 'removed' changesets (orderd by their number). So you have 000010.added.nt before 000234.added.nt and 999999.added.nt before 000001.removed.nt and so on. |
| deletesFirst | Like insertsFirst but first 'removed' then 'added' changesets |
| ID | This states that the changesets will be first ordered by the number and then by 'added' before 'removed'. So you have 000011.added.nt before 000011.removed.nt but directly after 000010.removed.nt |
| DI | Like ID but 'removed' changesets before 'added' ones. |

The following properties can be set:

| name | values/type | description | required? |
|--------------------------|------------------------------------------|---------------------------------------------------------------------------------------------------|-----------------------|
| ${\it queryPatternFile}$ | String | The filename with the query patterns | required |
| updateStrategy | fixed variation | The described strategy to use if you're using SPARQL and either UPDATE queries or live data files | optional |
| x | Integer | If updateStrategy is set this is the parameter which specified the described parameter x | optional |
| time-limit | Integer | The time in milliseconds how long the testcase should run | required |
| limit | Integer | The number of how much queries should be generated for one query pattern | default=5000 |
| ldLinking | insertsFirst deletesFirst ID DI | The internal live data strategy to consider which is the next changeset to querying | (only with ldPath) |
| ldPath | String | If you're using live data you can set the path where your changesets are located | (only with ldLinking) |

class: de.uni_leipzig.iguana.testcases.QueryTestcase

3.3.5.3 StressTestcase The stress testcase provides an approach to test a real live situation with a given number of SPARQL users and UPDATE users. It will test the QueryTestcase with several threads (users) to achieve this real live situation.

It takes the same properties like QueryTest case (which are the properties the threads will have) plus the properties $update\text{-}users,\ user$ and their attribute value .

class: de.uni_leipzig.iguana.testcases.StressTestcase

3.4 Configuration

The IGUANA is configured through a XML file.

3.4.1 Root node

The root node is required and every section after the root node section is a tag inside this root node.

The root node looks like the following:

3.4.2 Data description

The data description tag contains the description how the converter will handle the data. For example which resource URI the converter should use. The data-description tag itself is only required if a converter will be used. The following table shows the tags the data-description needs

| tagname | attributes | description | required? |
|----------------------|------------|-------------------------------------------------------------------------------------------------------------------|-----------|
| namespace | name=URI | This provides the actual namespace | required |
| anchor | name=CDATA | This provides the property suffix so the property URI will be namespace+anchor | required |
| resource-uri | name=URI | This provides the resource-uri | required |
| property-prefix-name | name=CDATA | this is needed if you use outputFormat like TURTLE so the propertyURIs can be replaced by the prefix | required |
| resource-prefix-name | name=CDATA | this is needed if you use output- Format like TURTLE so the re- sourceURIs can be replaced by the prefix | required |

Example:

3.4.3 Databases

The databases tag contains several database tags.

The database tag has two attributes, one is the id tag and the other one is the type tag. The type can be either *impl* or *curl*.

The impl type converts files which will be uploaded into an INSERT tag. Also all UPDATE queries will be send directly over the SPARQL endpoint. This can be slow and the RAM can be very quickly out of memory.

The curl type sends all UPDATE queries through a specified script command (e.g. curl). The tags which will be only for the curl type have the prefix *curl*-. On these one you can use several variables like \$USER, \$PWD, \$CURL-URL, \$GRAPH-URI, \$UPLOAD-TYPE, \$FILE, \$UPDATE

Some of them can be replaced by the explicit string, but the variables FILE, UPDATE and GRAPH-URI must be set.

Both use the SPARQL endpoint directly for SPARQL queries.

| tagname | attributes | description | required? |
|--------------|------------|----------------------------------------------------|-------------|
| endpoint | uri | The SPARQL endpoint | required |
| user | value | The user for the endpoint | optional |
| pwd | value | The password for the user | optional |
| curl-url | url | The curl-url (as it can differs from the endpoint) | (only curl) |
| curl-command | command | The script command to upload a file | (only curl) |
| curl-drop | command | The script command to drop a specific graph | (only curl) |
| curl-update | command | The script command to send an UPDATE query | (only curl) |

Example:

```
<?xml version="1.0" encoding="utf-8"?>
<iguana xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">
       <databases>
              <database id="curl" type="curl">
                      <endpoint uri="localhost/sparql-auth" />
                      <user value="dba" />
                      <pwd value="dba" />
                      <curl-url url="http://localhost/sparql-graph-crud-auth" />
                      <curl-command command="curl --digest --user</pre>
                             $USER:$PWD --url '$CURL-URL?graph-uri=$GRAPH-URI'
                             $UPLOAD-TYPE -T $FILE"/>
                      <curl-drop command="curl -X DELETE</pre>
                             $GRAPH-URI --url '$CURL-URL'"/>
                      <curl-update command="curl -i -d '$UPDATE' -u '</pre>
                             $USER:$PWD' -H 'Content-Type:
                             application/sparql-query' --url 'http://$ENDPOINT'"/>
              </database>
              <database id="impl" type="impl">
                      <endpoint uri="localhost/sparql-auth" />
                      <user value="dba" />
                      <pwd value="dba" />
              </database>
              <database id="impl2" type="impl">
                      <endpoint uri="localhost/sparql" />
               </database>
       </databases>
</iguana>
```

3.4.4 Log-clustering

The log-clustering tag which is optional, contains three attributes which are all required and are the following:

attribute description

class The full classname. This states the class to use for the log clustering process. Remember this one must be a class which implements

the Clusterer.java interface

path The path of the logfiles

output-file The output file in which the final queries should be written in.

The output-file can be used in the testcases. So you can both test clustered queries and some queries you provide if you want to.

To set a property simply add a tag inside the log-clustering tag, called property, with the attributes name and value. Name declares the property name and value the value for the property. These properties will be set to the *Clusterer* as a *Properties* object.

Example:

3.4.5 Benchmark

To convert your files into RDF files (notice that these files will be string together to one file which will be used as the initial file and will be uploaded in the reference connection) before the benchmark starts you must add the following five tags.

| tag name | attributes | description |
|----------------------|------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| convert-processing | value | The tag to state if there is data which must be converted |
| convert-input-path | name | The path of the files which should be converted |
| convert-class | class | The actual Converter class name to use for converting the files |
| rdf-vocabulary-class | class | The class name to use for the RDFVocabulary. |
| output-path | name | This sets the path in which the converted files will be saved in (Be aware that you don't need the output-path or the file-names in it to test the dataset and/or generated new datasets out of it. Just let the initFile attribute of the random-function tag empty. The benchmark will handle it.) |

If you want to drop the connection (or if graph-uri is set the graph) every time the connection will be tested (for every percent) you can set the **drop-db** tag with the attribute *value* which can be true, to state that the dataset in the connection should be dropped, or false, to state that the dataset should not be dropped.

Be aware that if you specify a *graph-uri* tag it will only drop this specific graph. It is highly recommend to set this tag to true!

To set that the tests should only be tested on a specific graph you can add a **graph-uri** tag which has the attribute *name*. In this tag you simply write the graph URI which will be tested.

The **random-function** tag states which percentage/files will be tested and if they needs to be generated.

The tag itself has three attributes. The first one is called *generate* which states if the datasets (smaller and greater than 100%) must be generated.

To set the algorithm for generating smaller datasets there is the second attribute which is called *type*. This attribute can be either *RandomInstance*, *RandomTriple* or *coherence*. The first two are the generation algorithms out of the first DBpedia SPARQL benchmark paper[2]. In this paper they are called *seed* and *rand*. The latter one is an implementation based on coherence of the dataset [1]. The third attribute sets the initial file (the 100% file). This should be empty if the converter generates this file.

If you use the type coherence you can specify two more attributes by adding

roh and coherence as attributes. roh is the relaxation parameter specified in the referenced paper and coherence is the parameter to maintain by the generation which is also specified in the referenced paper.

To set which percentages should be tested you can add a **percent** tag in the random-function tag. The percent tag has two attributes. The first one called value which should be a double (1.0 stands for 100%) and states the percentage to test. The second attribute is called file-name which can set the name of the file/path if you don't need to generate it and you have the specific file or files in the specified path.

To warmup the connections before you test them (this will happen for every percentage you'll set) you can add a **warmup** tag which has three attributes. The first one is the *time* attribute. The value is the time in milliseconds how long the warmup will go on. The second one is the *file-name* attribute. This specify where to find the file with the queries (not query patterns!) to warmup the connection with. The last attribute is *update-path*. This specify the directory where to find rdf files to upload. This can be empty which means there are no updates.

The **test-db** tag sets which databases (must be existing ones in the databases tag) will be tested.

The test-db tag has two attributes, the first one is the *reference* attribute which is the id (the database tag id) which of the given connections should be used as a reference (for example to convert the query patterns into queries with real data in it).

The second attribute is the type attribute which can be either all or choose. If it's set to all every connection in the databases tag will be tested except for the reference connection. If it's set to choose only the connection specified in the test-db tag will be tested (even though you can specify the reference connection to be tested it will not be tested!).

To specify the connections you add a db tag in the test-db tag with the attribute id. The id attribute must match one of the database ids.

3.4.5.1 Testcases The testcases tag contains several testcase tags. The testcases tag is required even though it can be empty.

The testcase tag has one attribute called *class* in which the full class name of the testcase class should stand. Remember that the testcase class must implement the *Testcase.java* interface.

To set a property simply add a tag inside the testcase tag called property, with the attributes name and value. Name declares the property name and value the value for the property. These properties will be set to the *Clusterer* as a *Properties* object.

Example:

<?xml version="1.0" encoding="utf-8"?>

```
<iguana xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance">
   <benchmark log="Benchmark">
       <convert-processing value="true"/>
       <convert-input-path name="src/test/resources/" />
       <rdf-vocabulary-class
         class="de.uni_leipzig.informatik.swp13_sc.datamodel.rdf.ChessRDFVocabulary"/>
       <converter-class
         class="de.uni_leipzig.informatik.swp13_sc.converter.PGNToRDFConverterRanged"/>
       <output-path name="src/test/java/"/>
       <drop-db value="false" />
       <graph-uri name="http://example.com" />
       <query-diversity value="1" />
       <random-function type="coherence" generate="true" initFile="" coherence="0.75" roh="</pre>
              <percent value="0.5" file-name=""/>
              <percent value="0.25" file-name=""/>
       </random-function>
       <warmup time="20000" file-name="Q.txt"/>
       <test-db type="choose" reference="impl2">
              <db id="curl" />
              <db id="impl" />
       </test-db>
       <testcases>
              <testcase class="de.uni_leipzig.iguana.testcases.QueryTestcase">
                     cproperty name="queryPatternFile" value="Q.txt"/>
                     property name="limit" value="2"/>
                     cproperty name="time-limit" value="3600000"/>
                     cproperty name="updateStrategy" value="fixed"/>
              </testcase>
              <testcase class="de.uni_leipzig.iguana.testcases.StressTestcase">
                     cproperty name="users" value="3"/>
                     cproperty name="queryPatternFile" value="Q.txt"/>
              </testcase>
              <testcase class="de.uni_leipzig.iguana.testcases.UploadTestcase"/>
       </testcases>
   </benchmark>
</iguana>
```

3.4.6 Email notification

The optional email-notification tag has one attribute called attach-results and can be set either true or false. If this attribute is set true, the results (or temporary results) will be zipped and attached to the email.

| tagname | attributes | description | required? |
|------------|------------|------------------------------------------------------------------------------------------------------------------------------------------------|-----------|
| hostname | value | The hostname of the email server | required |
| port | value | The port of the host | required |
| user | value | The user from which the email will be send | required |
| password | value | The password for the user. For security reason you can simply leave this tag away and the benchmark will ask you over the command line. | optional |
| email-name | address | The adress which will be shown in the sended email | required |
| email-to | address | The adresses to send the email to. Be aware if you want to send it to the email-name too you must add this explicit. Can be set multiply times | required |

Example:

4 Further information

4.1 Links

Beware that these following information can be outdated

| website | https://aksw.github.io/IGUANA/ |
|------------------------------------|--------------------------------------------------------------------------------------------------------------------------------|
| distribution | https://github.com/AKSW/IGUANA/tree/master |
| javadoc | https://aksw.github.io/IGUANA/javadoc |
| source code | https://github.com/AKSW/IGUANA |
| issue tracker | https://github.com/AKSW/IGUANA/issues |
| contact information | mai12cpg@studserv.uni-leipzig.de |
| Border Flow (Clustering algorithm) | http://borderflow.sourceforge.net |
| DBpedia SPARQL Benchmark | http://aksw.org/Projects/DBPSB.html |
| adjusted wookieConnection | https://github.com/AKSW/IGUANA/blob/develop/de.uni_ leipzig.iguana/src/main/resources/lib/connection-0.0. 1-SNAPSHOT.jar |

4.2 License

IGUANA itself is licensed under the MIT License [A], but be aware that there are libraries we use which are licensed under different licenses.

References

- [1] S. Duan, A. Kementsietsidis, K. Srinivas, and O. Udrea. Apples and oranges: A comparison of rdf benchmarks and real rdf datasets. In *Proceedings of the 2011 ACM SIGMOD International Conference on Management of Data*, SIGMOD '11, pages 145–156, New York, NY, USA, 2011. ACM.
- [2] M. Morsey, J. Lehmann, S. Auer, and A.-C. Ngonga Ngomo. DBpedia SPARQL Benchmark – Performance Assessment with Real Queries on Real Data. In ISWC 2011.
- [3] M. Morsey, J. Lehmann, S. Auer, and A.-C. Ngonga Ngomo. Usage-Centric Benchmarking of RDF Triple Stores. In *Proceedings of the 26th AAAI Conference on Artificial Intelligence (AAAI 2012)*, 2012.

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