

#### SOFTWARE EVOLUTION SERIES 2

# Report

December 14, 2017

Students: Cornelius Ries 11884827

Piotr Kosytorz 11876964 Tutor: Riemer van Rozen Course: Software Evolution

#### 1 Introduction

This documents contains our notes and answers to the questions about software metrics (practical lab Series 2).

Rewrite the introduction

#### 2 Definitions

- Clone (reference to kamiya2002ccfinder)
- Clone class (reference to kamiya2002ccfinder)
- The biggest clone
- The biggest clone class

#### 2.1 Clone type-1

Per definition from [2] type 1 clones are strict textual clones, excluding comments and whitespaces.

#### 2.2 Clone type-2

Per definition from [2] type 2 clones are "a syntactically identical copy; only variable, type, or function identifiers have been changed".

#### 2.3 Clone type-3

Per definition from [2] type 3 clones are copies with further changes like adding/removing statements.

### 3 Algorithm design

- General algorithm design (sketch: text description)
- Type-3 clean method (pseudocode) / We dont have it yet
- Formalised algorithm (pseudocode with references to type-1, type-2 clean methods)
- Describe how the algorithm is covered in the literature (and provide references).

#### 3.1 General algorithm design

The idea and algorithm of our duplication detection is based on the information from [3] and [1]. The main idea behind this approach is to use the abstract syntax tree for clone detection. After creating the ast from the source files, the nodes are hashed into different buckets. If a bucket has more than one element, it follows, that there are multiple locations in the source code of this node. This counts as one clone class. There are multiple types of clones that our tool can detect. These are type 1 and type 2 clones.

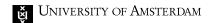
#### 3.2 Type-1 cleaning

If using the AST approach, the bucket key is in general just the node itself. With rascal though, there was a change in the implementation that shifted the loc and other informations of a node from the annotations on the node, to information contained in the node. This messed up the matching because every location was unique, which resulted in every node (because of subnodes) being unique. To circumvante the problem we remove the unnessecary information from all the relevant nodes. The actual cleaning cleanNode is done by visiting all the subnodes of a relevant node and replacing the node with a cleaned version of the original node using the insert statement.

#### 3.3 Type-2 cleaning

For type 2 detection we also use the AST. To streamline all the information that is contained in the node we visit all the subnodes of a node and replace specific variables/identifiers/types with a general one. The replacement is done using the rascal visit - case => notation.

```
1 case Type _ => lang::java::m3::AST::float()
 2 case Modifier => lang::java::m3::AST::\public()
  3 \ \mathbf{case} \ \backslash \mathtt{method}(\_, \_, \mathtt{a}, \mathtt{b}) \ \Rightarrow \ \backslash \mathtt{method}(\mathtt{lang} :: \mathtt{java} :: \mathtt{m3} :: \mathtt{AST} :: \mathbf{float}() \ , \ \mathtt{"method"}, \mathtt{a}, \mathtt{b} ) 
 4 case \method(_, _, a, b, c) => \method(lang::java::m3::AST::float(), "method",
           a, b, c)
 5 case \parameter(a, _, b) => \parameter(a, "parameter", b)
 6 case \vararg(a, __) => \vararg(a, "vararg")
 7 case \annotationTypeMember(a, _) \Rightarrow \annotationTypeMember(a, "
         annotationTypeMember")
 8 case \lambda \text{annotationTypeMember(a, -, b)} \Rightarrow \lambda \text{annotationTypeMember(a, "
         annotationTypeMember", b)
 9 case \typeParameter(_, a) => \typeParameter("typeParameter", a)
10 case \constructor(_, a, b, c) => \constructor("constructor", a, b, c)
11 case \setminus interface(_, a, b, c) \Rightarrow \setminus interface("interface", a, b, c)
            \begin{array}{l} \langle {\bf class}\,(\_,\ a,\ b,\ c) \Rightarrow \langle {\bf class}\,("\,{\bf class}",\ a,\ b,\ c) \\ \langle {\bf enumConstant}\,(\_,\ a) \Rightarrow \langle {\bf enumConstant}\,("\,{\bf enumConstant}",\ a) \end{array} 
12 case
13 case
           \operatorname{Constant}(a, a, b) \Rightarrow \operatorname{Constant}(a, a, b)
14 case
           \methodCall(a, _, b) => \methodCall(a, "methodCall", b) \methodCall(a, b, _, c) => \methodCall(a, b, "methodCall", c)
15 case
16 case
           \simpleName(_) => \simpleName("simpleName")
17 case
18 case \number(_) => \number("125681651")
```



```
19 case \variable(_, a) => \variable("variable", a)
20 case \variable(_, a, b) => \variable("variable", a, b)
21 case \booleanLiteral(_) => \booleanLiteral(true)
22 case \stringLiteral(_) => \stringLiteral("stringLiteral")
23 case \characterLiteral(_) => \characterLiteral("a")
```

#### 3.4 The main algorithm

Here we provide a textual representation of our main algorithm. For further information please have a look into the source file DuplicationsAnalyzer.rsc.

Build the AST of the project.

For all nodes in AST

- If size of node > threshold
- Clean node for type 1 detection.
- Clean node for type 2 detection.
- Collect node in map with cleaned node as key,
   relation of original node and location as value

For all keys in Map build a set of clone classes

- Collect all values from the map for current key
- If size of collected values > 1 Then add values as a new clone class to set

Filter subclone classes

- For all previously detected clone classes
- If another clone class exists for which all locations include the locations of the current one Then

Else

Add to new Set

For all filtered clone classes

- Collect them in output format
- Collect locations
- Determine line statistics
- Determine clone type

# 4 Main program validity

Write about automatic tests - describe what the tests do (be convincing)

All tests are in separate files that extend their original rascal module:

- DuplicationsAnalyzerTests
- RaterTests
- UtilsTests
- VolumeAnalyzerTests

To run the tests, import all the modules above and execute :test in the rascal console. The projectLocation in Configuration.rsc has to be set to the projects location in your eclipse!

#### 5 The tool

- General description and purpose, used solutions (webserver, REST api, ReactJS app, d3 graphs, etc)
- The 3 main requirements (your tool satis es from the perspective of a maintainer) [ref to storey1999cognitive]:
  - 1. Get a comprehensible overview (of code quality and duplications)
    - Readable table
    - SIG maintanability index
  - 2. Get a deep insight into the clones
    - navigate easily through them
    - provide and extended search function (easy to use = we reduce eort)
  - 3. See how the clones spread over my project (=Improve comprehension)
    - provide insightful visualization = Provide eective presentation styles (graph)
    - show how the les containing clones relate to each other (=indicate options for further exploration)
- Implementation choices
- The visualisation (constellation) how does it help a maintainer or a developer?

#### 6 Tool manual

- How to run
- How to use
- Where is the Rascal project
- Where is the original visualisation project

To use the tool we provide the source code as a eclipse project

#### 6.1 How to run

In order to run the program, please follow the steps:

- 1. Please import the project into your eclipse with a working rascal installation.
- 2. Open Configuration.rsc and adjust the location of the projectLocation to match the path of the project to your eclise
- 3. Do the same for the smallSqlProject and hqSqlProject
- 4. Start a rascal console and import the Main module
- 5. run startServe();
- 6. open a browser and point it towards http://localhost:5433 or to the location of serveAddress in case you changed it

### 7 Summary

- Sum things up
- Write how the tool performs on hsqlsb,
- Write how it can be improved

## References

- [1] Ira D Baxter et al. "Clone detection using abstract syntax trees". In: Software Maintenance, 1998. Proceedings., International Conference on. IEEE. 1998, pp. 368–377.
- [2] Rainer Koschke. Identifying and Removing Software Clones. 2008.
- [3] Flavius-Mihai Lazar and Ovidiu Banias. "Clone detection algorithm based on the Abstract Syntax Tree approach". In: Applied Computational Intelligence and Informatics (SACI), 2014 IEEE 9th International Symposium on. IEEE. 2014, pp. 73–78.