

Clone Overflow:

An empirical study of online code clone

RESEARCH QUESTIONS

- **RQ1 (online code clone):** *can we find any reused source code fragments between Q&A sites and open source projects?* We would like to observe whether this phenomenon has happened or has been happening and at what scale.
- **RQ2 (Direction of online code clone flow):** *in which direction source code have been reused?* If the code reuse between the two locations exist, we would like to discover the which direction the code that have been copied. Is it from Q&A site to open source projects, or the other way around, or both?
- **RQ3 (Online code clone classification):** *what are categories of online code clone?* How and why did these online code clone happen?
- **RQ4 (Effects of online code clone):** *Is this phenomenon of online code clone harmful and how?* is there any evidence of problems caused by reusing code between Q&A sites and open source projects?

DATASET

In our study, we selected Qualitas corpus containing 64 Java open source projects [5]. However, we found that *eclipse* project does not contain source code so we removed it from the dataset resulting in totally 63 projects being analysed. The details of the 63 Qualitas projects are listed in Table IV.

CLONE DETECTORS

We selected two clone detectors for this study: Simian [1] and NiCad [2], [3]. **FIXME: Add more info**

EXPERIMENTAL RESULTS

The results of running 2 clone detectors: Simian and NiCad, to detect clones between 144,075 Stackoverflow fragments (Java accepted answers) and 63 open-source projects in Qualitas dataset is presented below. There are 2 tools selected: Simian and NiCad. They are configured using two different settings: default settings, and settings from another study [6]. Full Simian's parameter names can be found from the footnote¹.

Manual investigation of Simian's clone report showed that there were problematic 11 fragments. These fragments generate false clone containing array initialisation. Hence, they were removed from the result set before analysis.

Table I: Qualitas-*O* (2013-09-01r): Simian and NiCad clone results

Statistics	Simian Default settings				Simian FSE'13 settings				NiCad Default settings				NiCad FSE'13 settings			
	threshold=6				threshold=5				threshold=6, UPI=0.30				threshold=5, UPI=0.2			
	iCharC,iCurlB,iIdC,iMod,iStrC				iCurlB,iId,iIdC,iStr,iChar,iSbtNm,bSqBrck				rename=none,abstract=none				rename=blind,abstract=literal			
	Frag.	C_{pairs}	C_{SLOC}	$C_{\%}$	Frag.	C_{pairs}	C_{SLOC}	$C_{\%}$	Frag.	C_{pairs}	C_{SLOC}	$C_{\%}$	Frag.	C_{pairs}	C_{SLOC}	$C_{\%}$
Total	654	24,929	–	–	1,175	16,957,362	–	–	985	105,118	–	–	12,084	113,557,298	–	–
Mean	–	38	7.54	0.27	–	14,444	4.80	0.28	–	107	9.52	0.25	–	9,397	5.21	0.20
Std Dev.	–	87	3.21	0.22	–	281,747	1.22	0.18	–	198	3.07	0.18	–	12,098	1.73	0.16
Max	–	551	49.00	0.94	–	9,599,676	18.00	0.89	–	1,792	39.00	0.80	–	227,077	44.00	0.86
Min	–	1	5.00	0.01	–	1	4.00	0.02	–	1	7.00	0.02	–	1	3.00	0.01
Median	–	3	7.00	0.23	–	22	5.00	0.24	–	15	8.00	0.19	–	6,105	5.00	0.15
Mode	–	1	7.00	0.25	–	1	4.00	0.50	–	1	8.00	0.53	–	1	4.00	0.33

Table II: Qualitas-*N* (2016-08-05): Simian and NiCad clone results (44 new + 19 old Qualitas projects)

Statistics	Simian Default settings				Simian FSE'13 settings				NiCad Default settings				NiCad FSE'13 settings			
	threshold=6				threshold=5				threshold=6, UPI=0.30				threshold=5, UPI=0.2			
	iCharC,iCurlB,iIdC,iMod,iStrC				iCurlB,iId,iIdC,iStr,iChar,iSbtNm,bSqBrck				rename=none,abstract=none				rename=blind,abstract=literal			
	Frag.	C_{pairs}	C_{SLOC}	$C_{\%}$	Frag.	C_{pairs}	C_{SLOC}	$C_{\%}$	Frag.	C_{pairs}	C_{SLOC}	$C_{\%}$	Frag.	C_{pairs}	C_{SLOC}	$C_{\%}$
Total	792	61,268	–	–	1,229	25,725,857	–	–	1,141	666,140	–	–	12,400	615,263,639	–	–
Mean	–	78	7.31	0.26	–	20,932	4.86	0.28	–	584	9.49	0.25	–	49,618	5.25	0.20
Std Dev.	–	208	2.70	0.21	–	382,015	1.30	0.19	–	1,488	3.11	0.18	–	62,659	2.80	0.16
Max	–	1,300	49.00	0.94	–	13,202,807	21.00	0.92	–	10,980	39.00	0.84	–	1,319,242	250.00	0.96
Min	–	1	5.00	0.01	–	1	4.00	0.12	–	1	7.00	0.02	–	1	2.00	0.01
Median	–	2	7.00	0.20	–	21	5.00	0.25	–	25	8.00	0.19	–	52,329	5.00	0.16
Mode	–	1	7.00	0.33	–	1	4.00	0.50	–	1	8.00	0.67	–	1	4.00	0.33

¹Simian's parameters: iChar = ignoreCharacters, iCurlB = ignoreCurlyBraces, iId = ignoreIdentifiers, iIdC = ignoreIdentifierCase, iMod = ignoreModifiers, iStrC = ignoreStringCase, iStr = ignoreStrings, iSbtNm = ignoreSubtypeNames, bSqBrck=balanceSquareBrackets

Table III: Qualitas-*N* (2016-08-05): Simian and NiCad clone results (only 44 new Qualitas projects)

Statistics	Simian Default settings threshold=6 iCharC,iCurlB,ildC,iMod,iStrC				Simian FSE'13 settings threshold=5 iCurlB,ild,ildC,iStr,iChar,iSbtNm,bSqBrck				NiCad Default settings threshold=6, UPI=0.30 rename=none,abstract=none				NiCad FSE'13 settings threshold=5, UPI=0.2 rename=blind,abstract=literal			
	Frag.	C _{pairs}	C _{SLOC}	C _%	Frag.	C _{pairs}	C _{SLOC}	C _%	Frag.	C _{pairs}	C _{SLOC}	C _%	Frag.	C _{pairs}	C _{SLOC}	C _%
	707	60,536	—	—	1,205	22,797,190	—	—	1,068	648,165	—	—	10,231	573,438,528	—	—
Total	—	86	7.37	0.26	—	18,919	4.85	0.28	—	607	9.39	0.25	—	47663	5.22	0.20
Mean	—	217	2.22	0.21	—	326,588	1.31	0.19	—	1,499	2.98	1.81	—	58,833	2.77	0.15
Std Dev.	—	1,298	32.00	0.94	—	11,127,286	21.00	0.92	—	10,550	36.00	0.84	—	1,246,598	250	0.96
Max	—	1	5.00	0.01	—	1	4.00	0.02	—	1	7	0.02	—	1	2.00	0.01
Min	—	2	7.00	0.21	—	20	5.00	0.25	—	33	8	0.19	—	52,077	5.00	0.15
Median	—	1	7.00	0.25	—	1	4.00	0.50	—	1	8	0.67	—	1	4.00	0.33
Mode	—	1	7.00	0.25	—	1	4.00	0.50	—	1	8	0.67	—	1	4.00	0.33

AGREED CLONE PAIRS

The agreed clone pairs using Bellon's *good-match*(0.7) and *ok-match*(0.7) criteria are listed in Table VI. We use the threshold of 0.7 for both *good-match* and *ok-match*. Visualisation of *good-match* common clone pairs between sets of parameter settings can be seen from Figure 1. There are 1,406 unique *good-match* pairs. The *ok-match* subsumes the *good-match* resulting in totally 10,879 common clone pairs (Figure 2).

However, for NiCad's settings, we found a few renaming and clustering errors which resulted in fewer projects with clone reports. For NiCad default settings (NiCad_{df}), 6 projects had clustering failed errors. For NiCad FSE13 settings (NiCad_{fse13}), 4 projects had renaming failed errors and 13 projects had clustering failed errors as depicted in Table V. So these projects are also missing from agreed clone pairs. **FIXME: Report the errors.**

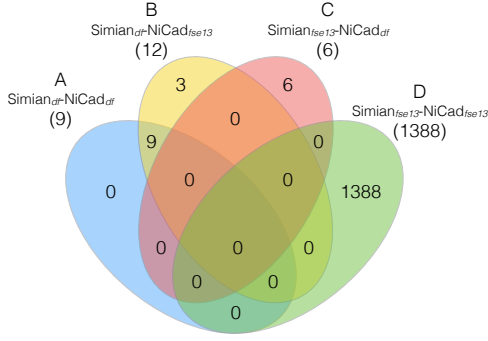
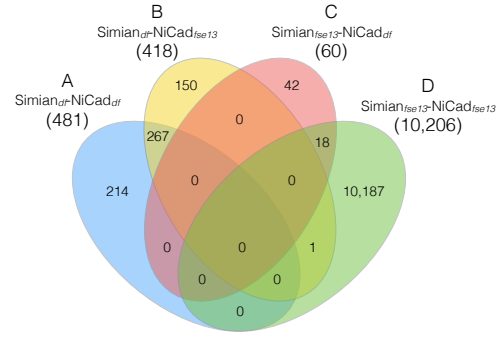
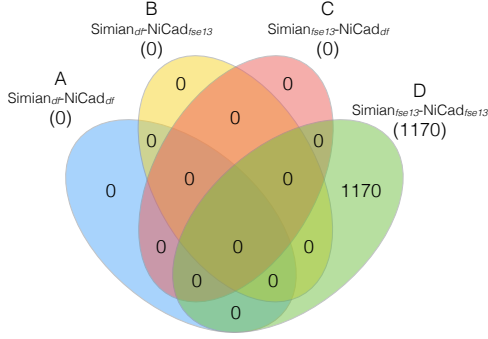
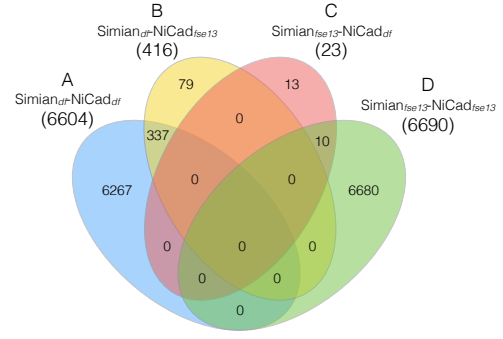
We are interested in discovering reused code in the latest versions of Qualitas projects. So, we downloaded the newest release of each project and found 44 of them having newer updates. Then, we reran the experiment again on these 44 projects. Several projects triggered NiCad problem of clustering and renaming again as listed in Table V. The agreed clone pairs using Bellon's *good-match*(0.7) and *ok-match*(0.7) criteria of this new dataset are also listed in Table VI.

Table V: No. of projects in Qualitas Original (Qualitas-*O*) and New (Qualitas-*N*) successfully analysed by Simian and NiCad

Qualitas- <i>O</i>			Qualitas- <i>N</i>		
Simian _{df} /Simian _{fse13}	NiCad _{df}	NiCad _{fse13}	Simian _{df} /Simian _{fse13}	NiCad _{df}	NiCad _{fse13}
63	57	46	44	40	34
	6 clustering failed cayenne checkstyle db-derby geotools iReport hibernate	13 clustering failed cayenne checkstyle db-derby geotools iReport ArgoUML castor drjava ganttpjproject ivatagroupware jasperreports jboss jchempaint		4 clustering failed jboss (wildfly) hadoop db-derby hibernate	6 clustering failed ArgoUML checkstyle db-derby cayenne jena geotools
		4 renaming failed Vuze aspectj eXoPortal hibernate			4 renaming failed Vuze hadoop jboss (wildfly) hibernate

Table VI: Distribution of clone pairs reported based on *good-match*(0.7) and *ok-match*(0.7) criteria

Tool		Qualitas- <i>O</i>		Qualitas- <i>N</i>	
Simian	NiCad	<i>good-match</i>	<i>ok-match</i>	<i>good-match</i>	<i>ok-match</i>
default	default	9	481	0	6,604
default	fse13	12	418	0	79
fse13	default	6	60	0	23
fse13	fse13	1,388	10,206	1,170	6,690
Total		1,415	11,165	1,170	13,396
Total (unique)		1,406	10,879	1,170	13,386

Fig. 1: Qualitas-O *good-match*(0.7) pairsFig. 2: Qualitas-O *ok-match*(0.7) pairsFig. 3: Qualitas-N *good-match*(0.7) pairsFig. 4: Qualitas-N *ok-match*(0.7) pairs

MANUAL INVESTIGATION

A. Investigation of agreed good and ok-match clone pairs

Table VII: Classifications of clone creation

Category	Descriptions
A	Code in Stackoverflow is copied from Qualitas ($Q \rightarrow S$).
A'	Code in Qualitas is copied from Stackoverflow ($S \rightarrow Q$).
B	Code is copied either from each other or a third source (unknown) ($S \leftrightarrow Q \vee (T \rightarrow S \wedge T \rightarrow Q)$).
C	Code in both places are copied from a third source T (known) ($T \rightarrow S \wedge T \rightarrow Q$).
D	Code is a boiler-plate or IDE auto-generated.
E	Code in both places initialise a similar/the same object; extend the same class/its subclass; implement the same interface.
F	Accidental similarity, false clone

1) *Qualitas-O* (v. 2013-09-01r): The classification scheme is described in Table VII and the classification results are shown in Table VIII. We have manually investigated all of the 1,406 *good-match* ones reported by agreement of four different Simian and NiCad settings. However, for the *ok-match*, we could not investigate all of the 10,879 pairs manually. According to the distribution of category from *good-match* results, we can see that $\text{Simian}_{fse13}\text{-NiCad}_{fse13}$ produces a large number, 1,387, of false positive results (D, E, and F). Thus, we decided to leave them out of the manual investigation of *ok-match* pairs. There are totally 664 *ok-match* pairs that were investigated. The 57 true positive pairs found are combinations of 10 unique Stackoverflow fragments, and 13 unique Qualitas Java files.

Since we are not certain about the direction of copying in the B-classified pairs, we checked the modification time of each Java file in Qualitas project and compare it to the timestamp of Stackoverflow answers. We found that all Stackoverflow code fragments were posted after their respectively similar Java files in Qualitas project. This means that the copying can only be either (1) $Q \rightarrow S$ or (2) from a third source to both S and Q independently.

Qualitas-N: We decided to filter out the agreed clone pairs reported by agreement of $\text{Simian}_{fse13}\text{-NiCad}_{fse13}$ altogether due to its large number of false positive. With the remaining 3 agreements, we did manual investigation of agreed *ok-match* clone pairs from agreements of $\text{Simian}_{df}\text{-NiCad}_{df}$, $\text{Simian}_{df}\text{-NiCad}_{fse13}$, and $\text{Simian}_{fse13}\text{-NiCad}_{df}$. We had not done any investigation of *good-match* since there were no *good-match* pairs for these 3 agreements. The number of pairs that have been looked at is 6,706 (excluding duplicates). The results are shown in Table.

Table VIII: Qualitas-*O*: Classification results of *good*- and *ok*-match pairs which excludes the subsumed *good*-match and $\text{Simian}_{fse13}\text{-NiCad}_{fse13}$ pairs.

Classificaion	A	A'	B	C	Sum	S_u	Q_u	Q_{up}	D	E	F	Sum	S_u	Q_u	Q_{up}	Total	S_u	Q_u	Q_{up}
<i>good-match</i> (0.7)	1	0	1	3	5	5	4	4	26	6	1369	1401	56	416	31	1406	61	420	32
<i>ok-match</i> (0.7)	8	0	36	8	52	8	10	6	474	57	81	612	77	62	17	664	84	70	20

Table IX: Qualitas-*O*: Distribution of classification category A–F according to *good*-match pairs

Category	A	A'	B	C	D	E	F	Total
$\text{Simian}_{df}\text{-NiCad}_{df}$	1	0	1	0	0	4	0	6
$\text{Simian}_{df}\text{-NiCad}_{fse13}$	0	0	0	3	1	1	1	6
$\text{Simian}_{fse13}\text{-NiCad}_{df}$	0	0	0	0	7	0	0	7
$\text{Simian}_{fse13}\text{-NiCad}_{fse13}$	0	0	0	0	18	1	1,368	1,387
Total	1	0	1	3	26	6	1,369	1,406

Table X: Qualitas-*O*: Distribution of classification category A–F according to the *ok*-match pairs

Category	A	A'	B	C	D	E	F	Total
$\text{Simian}_{df}\text{-NiCad}_{df}$	3	0	17	1	428	6	7	462
$\text{Simian}_{df}\text{-NiCad}_{fse13}$	5	0	18	7	17	51	50	148
$\text{Simian}_{fse13}\text{-NiCad}_{df}$	0	0	1	0	29	0	24	54
Total	8	0	36	8	474	57	81	664

2) *Qualitas-N* (v. 2016-08-05): We decided to filter out the agreed clone pairs reported by agreement of $\text{Simian}_{fse13}\text{-NiCad}_{fse13}$ altogether due to its large number of false positive. With the remaining 3 agreements, we did manual investigation of agreed *ok*-match clone pairs from agreements of $\text{Simian}_{df}\text{-NiCad}_{df}$, $\text{Simian}_{df}\text{-NiCad}_{fse13}$, and $\text{Simian}_{fse13}\text{-NiCad}_{df}$. We had not done any investigation of *good*-match since there were no *good*-match pairs for these 3 agreements. The number of pairs that have been looked at is 6,706 (excluding duplicates). The results are shown in Table XI.

Table XI: Qualitas-*N*: Classification results of *good*- and *ok*-matches (excluding the subsumed *good*-match pairs).

Classificaion	A	A'	B	C	Sum	S_u	Q_u	Q_{up}	D	E	F	Sum	S_u	Q_u	Q_{up}	Total	S_u	Q_u	Q_{up}
<i>good-match</i> (0.7)	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>ok-match</i> (0.7)	2	0	0	0	-	-	-	-	6676	1	27	-	-	-	-	6706	-	-	-

Table XII: Qualitas-*N*: Distribution of classification category A–F according to the *ok*-match pairs

Category	A	A'	B	C	D	E	F	Total
$\text{Simian}_{df}\text{-NiCad}_{df}$	0	0	0	0	6603	1	0	6604
$\text{Simian}_{df}\text{-NiCad}_{fse13}$	2	0	0	0	73	0	4	79
$\text{Simian}_{fse13}\text{-NiCad}_{df}$	0	0	0	0	0	0	23	23
Total	2	0	0	0	6676	1	27	6706

B. Investigation of Simian and NiCad clone pairs

1) *Qualitas-O* (v. 2013-09-01r): In the preliminary stage of our experiment, we found that there are 41 Stackoverflow fragments reported by Simian with default configurations. However, only 10 of them appear in the new results using tool's agreement. Thus, we further investigated the clone pairs reported by Simian and NiCad but *without* an agreement.

With our 4 settings, we decided to investigate only 2 settings, Simian_{df} and NiCad_{df} , and drop Simian_{fse13} and NiCad_{fse13} due to their large number of false positives as shown in Table IX and X. With the 2 selected settings, we investigated clone pairs having the minimum clone size of 10 SLOC as they are meaningful and tend to be real clone in modern clone detection [4].

For Simian_{df} , there were 9,393 clone pairs reported by the tool. Out of 9,393 pairs, 143 of them are the ones found in *ok*-pairs using agreement-based detection. We filtered the results further by removing false positives such as similar equals(), hashCode() methods, getters and setters out by using regular expression. We managed to remove 8,958 pairs using this method. Eventually, there were 292 clone pairs remaining for manual investigation. For NiCad_{df} , we obtained 7,062 clone pairs to look at which is infeasible for manual investigation. Hence, result filtering was also needed. However, regular expressions could not be used effectively as in Simian's case since NiCad allowed clones that are different at keywords/variable names or even added/deleted lines. So we decided to filter the results by selecting pairs that pass stricter clone criteria with $\text{UPI} = 0.2$. By reducing the UPI to 0.2, there were totally 171 pairs left. Out of 171, 57 are *ok*-pairs and 114 are remaining pairs for manual

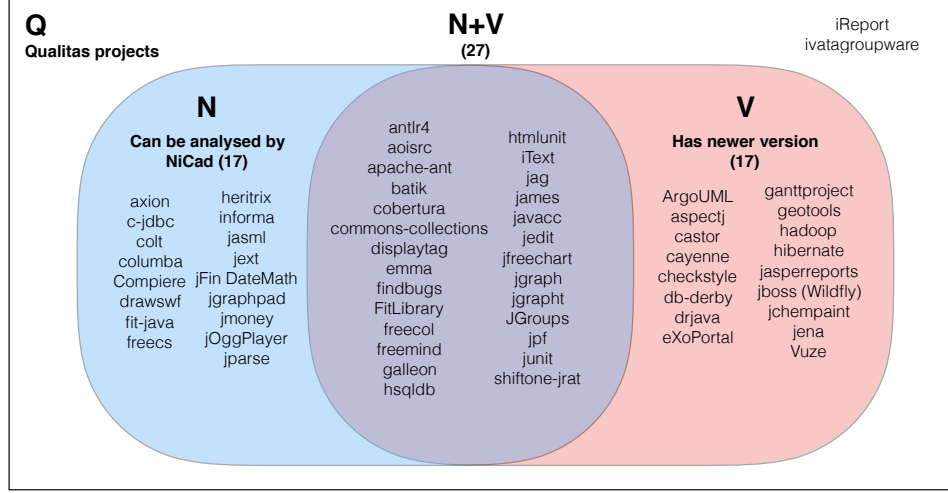


Fig. 5: Qualitas projects categorised by NiCad's results and their versions

check (18 pairs are from *cayenne* and *iReport* that could not be analysed using $UPI = 0.3$). The statistics of the clones and classification results are reported in Table XIII and XIV.

Table XIII: Statistics of $Simian_{df}$ and $NiCad_{df}$ clone pairs.

Tool	Clone pairs	ok-pairs	filtered pairs	remaining pairs
$Simian_{df}$	9393	143	8958	292
$NiCad_{df}$	7062	231	6717	114

Table XIV: Classification results of 292 $Simian_{df}$ and 114 $NiCad_{df}$ individual unique pairs. $NiCad_{df}^*$ shows the classification without *cayenne* and *iReport*.

Tool/Classification	A	A'	B	C	Sum	S_u	Q_u	Q_{up}	D	E	F	Sum	S_u	Q_u	Q_{up}	Total	S_u	Q_u	Q_{up}
$Simian_{df}$	35	0	90	8	133	68	59	23	13	12	134	159	39	72	23	292	103	125	31
$NiCad_{df}$	4	0	5	0	9	9	5	4	24	3	78	105	41	39	12	114	48	44	14
$NiCad_{df}^*$	4	0	1	0	5	4	5	3	23	1	67	91	28	31	10	96	33	37	13

Table XV: Numbers of true online clone pairs ($A+A'+B+C$) found by manual investigation

good-pairs	ok-pairs	$Simian_{df}$ pairs	$NiCad_{df}$ pairs	Total
5	52	133	9	199

C. Evolution of clone pairs from Qualitas-O to Qualitas-N

After located and classified all the online clone pairs, we are now interested in analysing the evolution of true online clone pairs. This is to answer RQ4. In this setting, we only focus on 44 projects in the region in V in Figure 5. They are projects which can be analysed Simian and also have newer versions than the versions in Qualitas-O. After filtering, there are 45 pairs removed because they do not have new versions (7 from *c-jdbc*, 21 from *Compiere*, 7 from *heritrix*, and 10 from *iReport*) and 154 pairs remaining.

Table XVI: Projects that do not have results in Qualitas- N but do have results in Qualitas- O

Project	Qualitas- O				Reason of missing results
	<i>good-match</i>	TP	<i>ok-match</i>	TP	
jboss	2	0	63	1	NiCad's failure
db-derby	0	0	0	0	NiCad's failure
hadoop	191	1	2432	5	NiCad's failure
hibernate	19	0	11	0	NiCad's failure
ArgoUML	0	0	0	0	NiCad's failure
checkstyle	0	0	0	0	NiCad's failure
cayenne	0	0	0	0	NiCad's failure
geotools	0	0	0	0	NiCad's failure
jena	0	0	0	0	NiCad's failure
Vuze	0	0	0	0	NiCad's failure
Compiere	127	2	178	8	No update
axion	7	0	12	0	No update
c-jdbc	66	0	768	0	No update
colt	0	0	5	0	No update
columba	14	0	177	0	No update
drawswf	0	0	0	0	No update
fit-java	0	0	2	0	No update
freecs	5	0	22	0	No update
heritrix	16	0	190	0	No update
iReport	0	0	0	0	No update
informa	31	0	22	0	No update
ivatagroupware	0	0	0	0	No update
jFin	15	0	72	0	No update
jOggPlayer	12	0	52	0	No update
jasml	23	0	47	0	No update
jext	80	0	85	0	No update
jgraphpad	1	1	55	0	No update
jmonev	0	0	0	0	No update
jpase	0	0	0	0	No update
freecol	1	1	0	0	<i>FreeColMenuTest.java</i> exists with the <i>createImageIcon()</i> method. Still can't find the reason why it is not reported.
aoisrc281	0	0	33	33	<i>MovieEncoder.java</i> (in 2 locations) are missing from the latest version.
findbugs	0	0	2	2	<i>UnionBugs.java</i> has been changed.
jgraph	0	0	3	3	<i>HelloWorld.java</i> has been changed.
Total	610	5	4231	52	

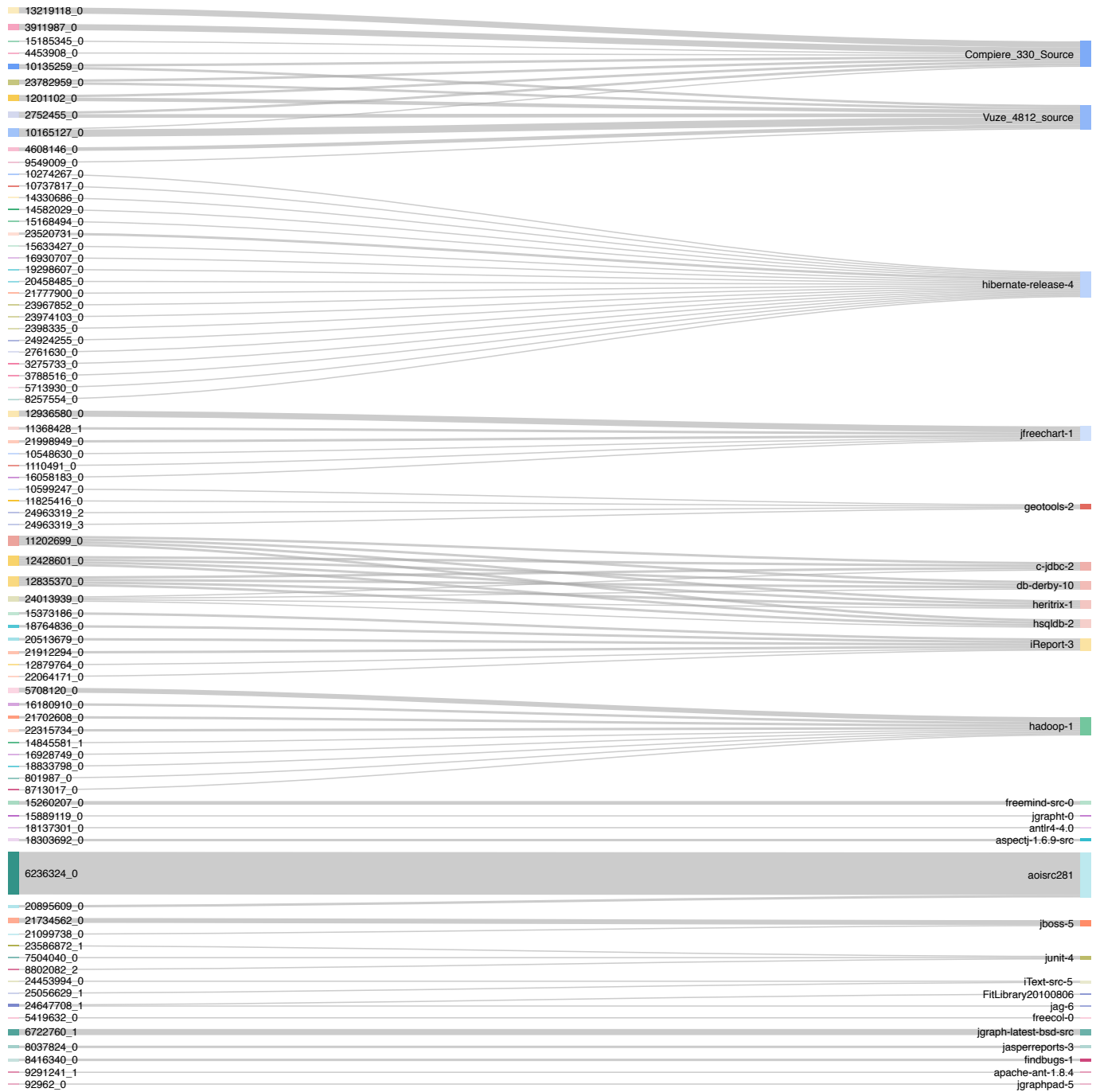


Fig. 6: Relationships between 199 true online code clone found between Stackoverflow and Qualitas projects

INVESTIGATION OF MISSING A/B CLONE PAIRS REPORTED BY SIMIAN (DF)

We investigated the 41 clone pairs previously reported by Simian with default configurations and manually investigated. The 41 pairs were searched for in 4 new results sets: Simian_{df} , Simian_{fse13} , NiCad_{df} , NiCad_{fse13} . The investigation results are shown in Table XVII.

Table XVII: Results of matching the original 41 Simian(default) pairs in the pretty-printed result sets

Settings	Found	Not found
Simian_{df}	40	1*
Simian_{fse13}	0	41
NiCad_{df}	17	24
NiCad_{fse13}	24	17

The single missing Stackoverflow fragment (19051537_0.java) (denoted by *) is one of the 11 false clones generated by Simian. It is removed from the results of the pretty-printed version because it is an outlier. The rest are missing because of different parameter settings.

SIMIAN'S PARAMETERS

We have carefully investigated the effects of the Simian's parameter `-balanceSquareBrackets+`. I found that it works in the expected way of handling a pair of brackets ([,]) that span over multiple lines. For example, the two code fragments in Figure 7 would match by having `-balanceSquareBrackets+` enabled.

<pre> 1 public class MagicSquare { 2 private int[][] square; 3 private boolean[] possible; 4 private int totalSqs; 5 private int sum; 6 private int numsquares; 7 public static void main (String[] args) { 8 MagicSquare m = new MagicSquare (3); 9 } 10 } </pre>	<pre> public class MagicSquare2 { private int[][] square; private boolean[] possible; private int totalSqs; private int sum; private int numsquares; public static void main (String[] args) { MagicSquare m = new MagicSquare (3); } } </pre>
-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------	------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------

Fig. 7: Two identical fragments with only differences in locations of the square brackets. All 7 lines are reported by Simian if `-balanceSquareBrackets+` is enabled. If not, the clone pairs is reported as (MagicSquare.java [3,8], MagicSquare2.java [5,10]).

However, the `-balanceSquareBrackets+` parameter only works on a small testing environment having toy programs or only small pairs from the full datasets. It does not work with the full complete set of 144,075 Stackoverflow fragments and Qualitas projects. Please find the summary of all the testing scenarios in Table XVIII.

Table XVIII: Simian's `-balanceSquareBrackets+` (`-bsb+`) is observed to have unpredictable behaviours when running against big datasets. CP_2 means the reported clone(s) do not contain lines having dislocated brackets (L_b) (i.e. $CP_2 = CP_1 - L_b$).

Project 1	Project 2	Dislocated brackets?	-bsb+	Clones pair reported
<i>Only run Simian against the pair</i>				
MagicSquare.java	MagicSquare_exact_copy.java	no	0,1	CP_1
MagicSquare.java	MagicSquare2.java	yes	0	CP_2
MagicSquare.java	MagicSquare2.java	yes	1	CP_1
stackoverflow/4298836_0.java	Qualitas/aoisrc281/./ExprModule.java	no	0	CP_3
stackoverflow/4298836_0.java	Qualitas/aoisrc281/./ExprModule.java	no	1	CP_3
stackoverflow/4533682_1.java	Qualitas/cobertura-1/./TouchCollector.java	no	0	CP_4
stackoverflow/4533682_1.java	Qualitas/cobertura-1/./TouchCollector.java	no	1	CP_4
<i>Run Simian against the complete stackoverflow data and the project</i>				
stackoverflow/4298836_0.java	Qualitas/aoisrc281/./ExprModule.java	no	0	CP_3
stackoverflow/4298836_0.java	Qualitas/aoisrc281/./ExprModule.java	no	1	—
stackoverflow/4533682_1.java	Qualitas/cobertura-1/./TouchCollector.java	no	0	CP_4
stackoverflow/4533682_1.java	Qualitas/cobertura-1/./TouchCollector.java	no	1	—

REFERENCES

- [1] Simian. <http://www.harukizaemon.com/simian>. Accessed: 07.04.2016.
- [2] James R. Cordy and Chanchal K. Roy. The NiCad Clone Detector. In *ICPC '11 Proceedings of the 2011 IEEE 19th International Conference on Program Comprehension*, pages 3–4, 2008.
- [3] Chanchal K. Roy and James R. Cordy. NICAD: Accurate Detection of Near-Miss Intentional Clones Using Flexible Pretty-Printing and Code Normalization. In *2008 16th IEEE International Conference on Program Comprehension*, pages 172–181, 2008.
- [4] Hitesh Sajnani, Vaibhav Saini, Jeffrey Svajlenko, Chanchal K Roy, and Cristina V Lopes. SourcererCC: Scaling Code Clone Detection to Big-Code. In *Proceedings of the 38th International Conference on Software Engineering - ICSE '16*, pages 1157–1168, 2016.
- [5] Ewan Tempero, Craig Anslow, Jens Dietrich, Ted Han, Jing Li, Markus Lumpe, Hayden Melton, and James Noble. Qualitas corpus: A curated collection of java code for empirical studies. In *2010 Asia Pacific Software Engineering Conference (APSEC2010)*, pages 336–345, December 2010.
- [6] Tiantian Wang, Mark Harman, Yue Jia, and Jens Krinke. Searching for Better Configurations: A Rigorous Approach to Clone Evaluation. In *Proceedings of the 2013 9th Joint Meeting on Foundations of Software Engineering*, pages 455–465, 2013.

Table IV: 63 Qualitas projects (new versions checked on 2016-08-05)

Projects	Old version	New versions	Latest change	Repo.	License	Notes
ArgoUML	0.34	0.35.1	15/01/2014		Eclipse Public License v.1.0	
Compiere	330	–	01/06/2010	git	GNU General Public License v.2.0	No longer OSS
FitLibrary	20100806	20111026	12/07/2014		GNU General Public License v.2.0	
JGroups	2.10.0.GA	3.6.10	27/06/2016		Apache License, v.2.0	
Vuze	4812	5721-20	12/05/2016		GNU General Public License v.2.0	
antlr4	4.0	4.2.1	15/01/2014		BSD License	
aoisrc	281	302	12/01/2015		GNU General Public License v.2.0	
apache-ant	1.8.4	1.10.0	09/04/2016	git	Apache License, v.2.0	
aspectj	1.6.9	1.8.9	14/03/2015		Eclipse Public License 1.0	
axion	1.0-M2	–	08/03/2013		Proprietary (BSD/Apache-style)	
batik	1.7	1.8	16/03/2015		Apache License, v.2.0	
c-jdbc	2.0.2	–	16/09/2005		GNU Lesser General Public License, v.2.1	
castor	1.3.1	1.4.2	20/06/2016	git	Apache License, v.2.0	
cayenne	3.0.1	4.0.M4	27/07/2016	git	Apache License, v.2.0	
checkstyle	5.1	7.2	04/08/2016	git	GNU Lesser General Public License, v.2.1 & Apache License, v.2.0	<i>Cli, Logging and Beanutils</i> packages are from the Apache Commons project.
cobertura	1.9.4.1	2.1.2	01/06/2015	git	GNU General Public License v.2.0	
colt	1.2.0	–	09/09/2014		Proprietary (CERN)	
columba	1.4	–	20/04/2007		Mozilla Public License 1.1	
commons-collections	3.2.1	4.1	25/11/2015		Apache License, v.2.0	
db-derby	10.6.1.0	10.12.1.1	13/09/2015		Apache License, v.2.0	
displaytag	1.2	2.0	17/08/2014	svn	MIT License	
drawswf	1.2.9	–	02/04/2013		GNU General Public License v.2.0	
drjava	20100913-r5387	20140826-r5761	26/08/2014		BSD License	
eXoPortal	???	4.4	24/06/2016	git	GNU Lesser General Public License v.3.0 & proprietary	
emma	2.0.5312	2.0.5312	09/05/2013		Common Public License 1.0	
findbugs	1.3.9	3.0.1	06/03/2015		GNU Lesser General Public License, v.2.0	
fit-java	1.1	–	04/06/2013		GNU General Public License v.2.0	
freecol	0.10.7	0.11.6	04/08/2016	git	GNU General Public License v.2.0	
freecs	1.3.20100406	–	22/04/2013		GNU General Public License v.3.0	
freemind	0.9.0	1.0.0	26/06/2016		GNU General Public License v.2.0+	
galleon	2.3.0	2.5.6	29/04/2013		GNU General Public License v.2.0	
gantproject	2.0.9	2.8.1	05/08/2016		GNU General Public License v.3.0	
geotools	2.7-M3	16	02/08/2016	git	GNU Lesser General Public License, v.2.0	
hadoop	1.0.0	2.6.4	11/02/2016		Apache License, v.2.0	
heritrix	1.14.4	–	05/06/2013		GNU Lesser General Public License, v.2.1	
hibernate	4.2.2	5.2.2	04/08/2016	git	GNU Lesser General Public License, v.2.1+	
hsqldb	2.0.0	2.3.4	16/05/2016	svn	BSD License	
htmlunit	2.8	2.24	27/07/2016	svn	Apache License, v.2.0	
iReport	3.7.5	–	28/05/2014		Affero GNU Public License, v.3.0	
iText	5.0.3	7.0.0	02/05/2016		Affero GNU Public License, v.3.0	
informa	0.7.0-alpha2	–	07/11/2008		GNU Lesser General Public License, v.2.1 & Apache Software License, v.1.1	
ivatagroupware	0.11.3	–	27/02/2013		GNU General Public License v.2.0	
jFin DateMath	RI_0_1	–	25/04/2013		GNU General Public License v.2.0	
jOggPlayer	114s	–	15/04/2013		GNU General Public License v.2.0	
jag	6.1	6.2	08/04/2013		GNU General Public License v.2.0 & BSD License	BSD License is for libraries.
james	2.2.0	2.3.2.1	14/08/2015		Apache License, v.2.0	
jasml	0.10	–	08/03/2013		Apache Software License	
jasperreports	3.7.4	6.3.1	20/06/2016	git	GNU Lesser General Public License, v.3.0	
javacc	5.0.0	7.0.0	07/07/2016	svn	Proprietary (Sun)	
jboss (Wildfly)	5.1.0.GA	10.0.0	29/01/2016		GNU Lesser General Public License, v.2.1	Renamed to Wildfly.
jchempaint	3.0.1	3.1.3	18/04/2016	git	GNU Lesser General Public License, v.2.1+	
jedit	4.3.2	5.3.0	20/10/2015		GNU General Public License v.2.0	
jena	2.6.3	3.1.0	10/05/2016		Apache License, v.2.0	
jext	5.0	–	18/08/2004		GNU General Public License v.2.0	
jfreechart	1.0.13	1.2.0-ea1	08/05/2016	svn	GNU Lesser General Public License, v.2.0	
jgraph	5.13.0.0	3.5.1.5	20/07/2016	git	Proprietary (mxGraph License)	
jgraphpad	5.10.0.2	–	10/11/2006		GNU General Public License & GNU Lesser General Public License (derivatives)	
jgraphpt	0.8.1	0.9.2	03/04/2016		GNU Lesser General Public License, v.2.1 & Eclipse Public License, v.1.0	
jmoney	0.4.4	–	17/04/2016		GNU General Public License v.2.0	
jpase	0.96	–	29/07/2004		GNU Lesser General Public License, v.2.1	
jpf	1.5.1	???	13/01/2012		Apache License, v.2.0	
junit	4.11	4.12	04/12/2014		Eclipse Public License, v.1.0	
shiftone-jrat	0.6	1-beta-1	25/04/2013	svn	GNU Lesser General Public License, v.2.0	