

# How do various plagiarism detection tools differ in their detection results when applied to java/spring coding exercises

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## **Abstract**

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# 1 Introduction

## 2 How Does Plagiarism Detection Work

\*Fand den Abschnitt bei moss ganz nett könnte man ja mit reinpacken\* Dennis G.

Moss and other plagiarism detection tools are not perfect, so a human should go over the results, and it should be checked if the claims are valid. "In particular, it is a misuse of Moss to rely solely on the similarity scores. These scores are useful for judging the relative amount of matching between different pairs of programs and for more easily seeing which pairs of programs stick out with unusual amounts of matching. But the scores are certainly not a proof of plagiarism. Someone must still look at the code." (Aiken, 2021)

## 3 Use Case and Software Experiment

### 3.1 Use Case

- Java/Spring exercises in ST2 - Automated pipeline for evaluation and creation of tasks - minor attempts to stop plagiarism -> random selection of exercise terms -> pipeline requires these terms - otherwise no differences between exercises - plagiarism attempts did occur -> high amount of submissions > 100(ask for number) makes it difficult to detect

### 3.2 Software Experiment

- attempt to find best tool to integrate into divekit - evaluate known plagiarisms - evaluate all submission for the ST2 module -> look for best results - systematically create plagiarisms to evaluate the tools' robustness

- Plagiarism s - rename variables and method -> mandatory for the tool due to the exercise design - change control structures -> for loop to while, if to switch etc. - duplicate code - extract methods from other methods - combine methods - move methods(create new classes)

### 3.3 Criteria for Evaluation

- must be able to evaluate Java 11 and spring exercises - Known plagiarisms must be detected  
- must be able to evaluate over 100(?) exercises in a reasonable speed - must be able to be integrated into the divekit pipeline or workflow **ask Stefan for his own criteria**

## 4 Evaluation of the different Tools

	C	C++	C#	Java	Kotlin	Python	PHP	VB.net	Javascript	Is expandable?
MOSS	Yes	Yes	Yes	Yes	No	Yes	No	Yes	Yes	No
JPlag	No	Yes	Yes	Yes	Yes	Yes	No	No	No	Yes
Plaggie	No	No	No	Yes	No	No	No	No	No	Yes <sup>1</sup>
AC2	Yes	Yes	No	Yes	No	Yes	Yes	No	No	Yes

Table 1 [The native supported programming languages for each plagiarism detection algorithm]

<sup>1</sup>Plaggie is open source, therefore a custom tokenizer for different languages can be implemented.

## 4.1 MOSS

"Moss (for a Measure Of Software Similarity) is an automatic system for determining the similarity of programs. To date, the main application of Moss has been in detecting plagiarism in programming classes. Since its development in 1994, Moss has been very effective in this role." (Aiken, 2021)

Moss claims to be one of the best cheating detection algorithms.<sup>2</sup> Moreover, moss supports the largest native programming languages variety<sup>3</sup>, but no other languages can be added. Moss supports the following languages: C, C++, Java, C#, Python, Visual Basic, Javascript, FORTRAN, ML, Haskell, Lisp, Scheme, Pascal, Modula2, Ada, Perl, TCL, Matlab, VHDL, Verilog, Spice, MIPS assembly, a8086 assembly, a8086 assembly and HCL2.

A unique feature of Moss is that it "is being provided as an Internet service. The service has been designed to be very easy to use[...]" (Aiken, 2021). In order to use this internet service, a Perl script must be executed which passes the files to the server. "In response to a query the Moss server produces HTML pages listing pairs of programs with similar code. Moss also highlights individual passages in programs that appear the same, making it easy to quickly compare the files. Finally, Moss can automatically eliminate matches to code that one expects to be shared (e.g., libraries or instructor-supplied code), thereby eliminating false positives that arise from legitimate sharing of code." (Aiken, 2021)

A disadvantage of Moss is that since Nov 13, 2022 there is a transmission limit of 100 submissions per day per user. This could cause problems if one wants to integrate Moss into an automated process.

### 4.1.1 Moss comparison algorithm

"Moss currently uses robust winnowing, which is more efficient and scalable (in the sense that it selects fewer fingerprints for the same quality of results) than previous algorithms we have tried. There are a few issues involved in making such a system work well in practice. For this application, positional information (document and line number) is stored with each selected fingerprint. The first step builds an index mapping fingerprints to locations for all documents, much like the inverted index built by search engines mapping words to positions in documents. In the second step, each document is fingerprinted a second time and the selected fingerprints are looked up in the index; this gives the list of all matching fingerprints for each document.

Now the list of matching fingerprints for a document  $d$  may contain fingerprints from many different documents  $d_1, d_2, \dots$ . In the next step, the list of matching fingerprints for each document  $d$  is sorted by document and the matches for each pair of documents  $(d, d_1), (d, d_2), \dots$  is formed. Matches between documents are rank-ordered by size (number of fingerprints) and the largest matches are reported to the user." (Schleimer et al., 2003)

### 4.1.2 Preliminary results

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<sup>2</sup>"The algorithm behind moss is a significant improvement over other cheating detection algorithms (at least, over those known to us)." (Aiken, 2021)

<sup>3</sup>Of the four plagiarism detection programs listed in this paper

File 1	File 2	Lines Matched
ST2M0-Geber1(81%)	ST2M0-Nehmer1(73%)	311
ST2M0-Geber2(62%)	ST2M0-Nehmer2(66%)	276
ST2M0-Geber1(14%)	ST2M0-Geber2(12%)	81
ST2M0-Geber2(11%)	ST2M0-Nehmer1(12%)	76
ST2M0-Nehmer1(9%)	ST2M0-Nehmer2(9%)	68
ST2M0-Geber1(9%)	ST2M0-Nehmer2(8%)	62

Table 2 [Table of Milestone [0] distance between projects]

File 1	File 2	Lines Matched
ST2M4-Geber1(54%)	ST2M4-Nehmer2(57%)	938
ST2M4-Geber1(52%)	ST2M4-Nehmer1(52%)	908
ST2M4-Nehmer1(51%)	ST2M4-Nehmer2(54%)	884
ST2M4-Geber2(13%)	ST2M4-Nehmer2(17%)	372
ST2M4-Geber2(12%)	ST2M4-Nehmer1(15%)	363
ST2M4-Geber1(14%)	ST2M4-Geber2(11%)	333
ST2M4-Geber2(8%)	ST2M4-Geber3(6%)	254
ST2M4-Geber3(6%)	ST2M4-Nehmer1(9%)	253
ST2M4-Geber1(9%)	ST2M4-Geber3(6%)	238
ST2M4-Geber3(5%)	ST2M4-Nehmer2(9%)	239

Table 3 [Table of Milestone [4] distance between projects]

## 4.2 JPlag

JPlag is a plagiarism detection tool, which was developed by in 2000 at the University of Karlsruhe to help with the detection of plagiarized coding exercises. At that the tool was only available as a WWW service and could analyze C, C++. Scheme and Java programs (Perchelt et al., 2000, p. 4). Currently, the tool is available as an open source application which can be run locally and can be used with a CLI or a Java API. JPlag also currently supports 12 programming languages including the original four languages and more modern languages such as Kotlin ("JPlag - Detecting Software Plagiarism", 2022, Supported Languages ) while also allowing to add new languages (Sağlam, 2022).

### 4.2.1 JPlag's comparison algorithm

JPlag's algorithm is split into two parts the tokenizing and the comparison of the token strings. During the tokenization process all programs are parsed and converted into token strings. In the second phase all token strings are compared in pairs to determine their similarity. This comparison uses a specially optimized version of the Greedy String Tiling algorithm. This algorithm tries to cover one token string with substrings of the other string. The percentage of the covered token strings is the similarity between the two programs (Perchelt et al., 2000, p. 10).

**Tokenization** The tokenization process is the only language dependent process of the JPlag algorithm (Perchelt et al., 2000, p. 10). The extracted tokens represent syntactic elements of the language like statements or control structures (Sağlam, 2022, How are submissions represented? — Notion of Token). During the parsing process each file is parsed with the result being a set of abstract syntax trees(AST) for each submission. Each AST is traversed depth first with nodes representing grammatical units of the language. During the traversal when entering and exiting a node a token can be created that match the type of the node and will then be added to the current list of tokens. There are block type nodes which can represent classes, if expressions or other elements of the language which have a corresponding beginning and end. When creating tokens from those nodes each have a corresponding "BEGIN" and "END" token. The token list should always have a pair of matching "BEGIN" and "END" tokens (Sağlam, 2022, How does the transformation work?).

**Comparing token strings** JPlag's comparison algorithm is essentially just the greedy string tiling algorithm for the comparison of two strings, however it is differently optimized to improve its runtime(Perchelt et al., 2000, p. 5). The goal of the algorithm is to find a set of substrings which are not only the same but also satisfy these three rules:

1. "Any token of A may only be matched with exactly one token from B."
2. "Substrings are to be found independent of their position in the string."
3. "Long substring matches are preferred over short ones[...]"

(Perchelt et al., 2000, p. 11)

These rule have some consequences. The first rule doesn't allow the matching of code that have been duplicated while the second rule makes the reordering of the source code not viable. Furthermore, the third rule is introduced since short matches are more likely to be spurious (Perchelt et al., 2000, p. 11).

When these rule 3 is applied sequentially then a greedy algorithm consisting of two phases will be created.

In phase 1 the longest contiguous match in the two strings is searched. This is done by 3 nested loops. The outer loop iterates over all extracted tokens in string A, the second loop compares token T from string A with every token in string B. If two tokens are identical then the innermost loop extends this match as far as possible (Perchelt et al., 2000, p. 11). These matches are called tiles which are permanent and unique (Wise, 1993, p. 3).

In the second phase all matches of the maximal length are marked, when this happens all matched tokens are also marked and can't be used for any further matches. This satisfies the first rule. These two phases will be repeated until not further matches are found (Perchelt et al., 2000, p. 11). Also, of importance is the MinimumMatchLength, which defines the length at which maximal matches and any tile below the maximum matches are ignored. The smallest possible value for MinimumMatchLength is 1. However, in general this value should be higher than 1 since at that length it is unlikely that a match will be significant (Wise, 1993, p. 3)

JPlag only considers two strings to match 100% when the shorter string is completely matched by the longer string. The logic being that the longer string represents a program which has been entirely copied and then extended. The similarity between two programs can be calculated with the following formula.

$$sim(A, B) = \frac{2 + coverage(tiles)}{|A| + |B|}$$

$$coverage(tiles) = \sum_{match(a,b,length) \in tiles} length$$

(Perchelt et al., 2000, p. 13).

#### -Detail the specific optimizations

GST has a worst-case complexity of  $O(n^3)$  (Wise, 1993, p. 5) and a best-case complexity of  $O(n^2)$  (Perchelt et al., 2000, p. 14). The worst-case complexity can't be improved the average-case complexity is improved to  $O(n)$  by implementing an idea from the Karp-Rabin algorithm (add foot note) (2000, p. 14). The modified algorithm computes the hash values for all substrings of length MML in string A and B. Then each value from string A is compared with each value from string B. String A is always chosen to be shorter than B because string A will always be traversed completely while string B will be accessed through a hash table, which will further decrease the complexity. If two hash values are the same then a potential possible match has been found beginning at that token where the values match. The possible match is then verified by comparing the substrings token by token. The modified algorithm doesn't start

the comparison from the beginning but rather from the back after the maxmatch tokens since all matches before that are not interesting for the algorithm. The algorithms also attempts to simultaneously expand the match further (Perchelt et al., 2000, p. 14).

#### 4.2.2 Preliminary results

JPlag displays its result in a list with all possible combinations of all submissions while also showing the amount of pairs which are in a certain similarity range. The different pairs can be further investigated allowing the user to see the specific matched code.

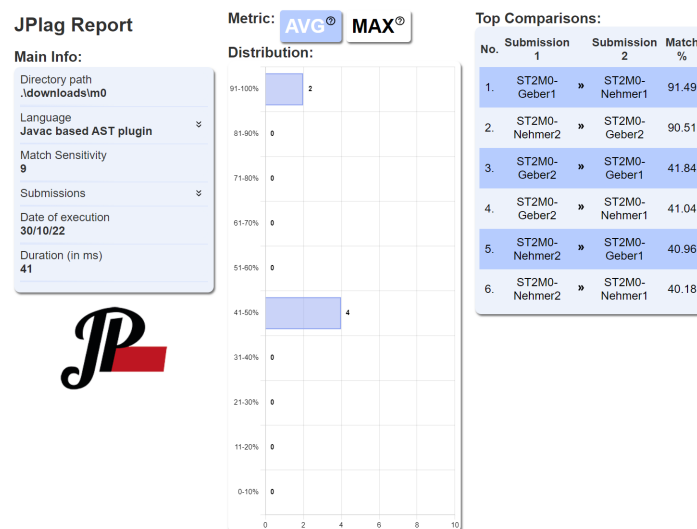


Figure 1 The JPlag overview showing the similarities of m0 code

When using JPlag to check the similarity of known plagiarized code it had similarity value for the plagiarized m0 code of 91.49% and 90.51% while the none plagiarized code had similarity of around 40%. The results for m4 were somewhat less clear since they had a similarity value of 84.77% and 76.2%. Which should still lead to an investigation of the submitted code.

Submission 1	Submission 2	Similarity	Similarity with base code
ST2M0-Geber1	ST2M0-Nehmer1	91.49%	88.11%
ST2M0-Nehmer2	ST2M0-Geber2	90.51%	86.11%
ST2M0-Geber2	ST2M0-Geber1	41.84%	16.34%
ST2M0-Geber2	ST2M0-Nehmer1	41.04%	16.21%
ST2M0-Nehmer2	ST2M0-Geber1	40.96%	14.61%
ST2M0-Nehmer2	ST2M0-Nehmer1	40.18%	14.56%

Table 4 [Table of Milestone [0] showing the similarity between submissions]

It should also be noted that no files were excluded and that there was no base code passed to JPlag, which explains the high similarity between unrelated programs as all submissions for a milestone use the same base code. When excluding common base code then the results for m0 become significantly more clear with the known plagiarism having a similarity value between 86 and 89 percent while the none plagiarized submissions have a value between 14 and 17 percent.

Looking at m4 with the base code excluded still shows a less clear image. With the known



plagiarized have a similarity value which is lower by 9 to 7 percentage points and with the none plagiarized submissions having a similarity value which lower by 18 to 16 percentage points. All of which are considerably smaller percentage point drops than those observed in m0. It is not clear why exactly the similarity values of m4 are closer to each other in m4 and why the difference between them doesn't grow that much when excluding the base code in comparison to m0.

Possible theories why yes no? Ask Stefan

Submission 1	Submission 2	Similarity	Similarity with base code
ST2M4-Nehmer2	ST2M4-Geber	84.47%	77.97%
ST2M4-Nehmer2	ST2M4-Nehmer1	77.48%	68.30%
ST2M4-Geber	ST2M4-Nehmer1	76.20%	67.42%
ST2M4-Nehmer2	ST2M4-NoPlag2	49.00%	32.70%
ST2M4-Geber	ST2M4-NoPlag2	47.20%	30.97%
ST2M4-Nehmer1	ST2M4-NoPlag2	46.16%	29.90%
ST2M4-Nehmer1	ST2M4-NoPlag1	41.82%	25.33%
ST2M4-Nehmer2	ST2M4-NoPlag1	40.55%	22.78%
ST2M4-Geber	ST2M4-NoPlag1	39.99%	22.69%
ST2M4-NoPlag2	ST2M4-NoPlag1	36.40%	20.15%

Table 5 [Table of Milestone [4] showing the similarity between submissions]

#### 4.2.3 Integration into an automated evaluation pipeline

JPlag probably show the most potential to be used in an automated pipeline as it not only provides a CLI but also a Java API, while also being able to be run locally("JPlag - Detecting Software Plagiarism", 2022) which means that there are no submission limits like in MOSS and that JPlag can be easily integrated into the existing pipeline. However, results are always saved as a Zip file and are view in a web application which can also be run locally. This poses a problem as???

-How does the pipeline work, are files permanently saved or deleted after it finishes evaluation, can they be accessed

### 4.3 Plaggie

-Was the only open source tool when release(Ahtianien et al., 2006), JPlag now also open source

#### 4.3.1 The Algorithm

-Uses CUP as a Parser -Uses standard LALR(1) parser generation("CUP", n.d.) -GST with no special optimization attempts -Algorithm was extended to support exclusion of common code(Ahtiainen, 2006, 4. Algorithm used)

-known unsuccessful attacks - Changing the comments - Changing the indentation - Method and variable name changes(Ahtiainen, 2006, Known successful attacks)

-know successful attacks - Moving inline code to separate methods and vice versa - Inclusion of redundant program code - Changing the order of if-else blocks and case-blocks(Ahtiainen, 2006, Known unsuccessful attacks)

#### **4.3.2 The results**

-TODO still have to run it.

#### **4.3.3 Integration into an automated evaluation pipeline**

-only Java 1.5(Ahtiainen, 2006; Ahtianien et al., 2006) -no further development since 2006 -has a cli -> can be automated -results as html files

## 4.4 AC2

"AC is a source code plagiarism detection tool. It aids instructors and graders to detect plagiarism within a group of assignments written in languages such as C, C++, Java, PHP, XML, Python, ECMAScript, Pascal or VHDL (plaintext works too, but is less precise). AC incorporates multiple similarity detection algorithms found in the scientific literature, and allows their results to be visualized graphically." (Freire, 2022a) The first version of "AC was born in the Escuela Politécnica Superior of the Universidad Autónoma de Madrid to deter and detect source-code plagiarism in programming assignments". (Freire, 2022a)

2016 AC version 2 was released. In the version 2.0 the developers switched to Maven and Antr4<sup>4</sup>. The developers of AC also moved their git repository to GitHub an internet hosting service for software development and version control.

AC2 lists some advantages over the other plagiarism detection tools on their GitHub page. In this regard, it is

- "local, and does not require sending data to remote servers (not the case of Moss)local, and does not require sending data to remote servers (not the case of Moss)"
- "robust, using Normalized Compression Distance as its main measure-of-similarity, but with the possibility of integrating other, additional measures to gain better pictures of what is going on (JPlag, Moss and Plaggie have hard-coded analyses, mostly based on sub-string matching after tokenization)."
- "heavy on information visualization. AC will not provide "percentage of copy"; instead, it will create graphical representations of the degree of similarity between student submissions within a group, so instructors can build their own explanations regarding what really happened (see here and here for papers on AC's visualizations)."

(Freire, 2022a)

### 4.4.1 AC2 comparison algorithm

In order for the AC algorithm to check files for equality, it allows adding folders or zip-files into it. Two filters can be applied to these files. The first filter selects the submissions that will be examined, and the second filter is determining which files are compared to each other.

For example student s1 has in his subfolder 'a.c' and student s2 has in his folder 'aa.c'. So the first filter should determine that each submission starts with the letter 's'. The second filter would look for files that ended with the string '.c'.

When all submission and files are correctly grouped together the analysis part can start. "The default analysis (Zip NCD), uses normalized compression distance (NCD) to detect redundancy (= similarity) between submissions. If AC2 knows the programming language of its source files,

<sup>4</sup>"In computer-based language recognition, ANTLR (pronounced antler), or ANOther Tool for Language Recognition, is a parser generator that uses LL(\*) for parsing. ANTLR is the successor to the Purdue Compiler Construction Tool Set (PCCTS), first developed in 1989, and is under active development."(Wikipedia contributors, 2022)

they will be tokenized prior to comparison. This will allow AC2 to ignore comments, whitespace and exact identifiers, so that simply renaming variables or altering formatting will not affect similarity calculations." (Freire, 2022b)

**NCD (normalized compression distance)** NCD (normalized compression distance) is a method to determine the similarity between two objects. These two objects can be of any type e.g. text, email, code, ... . To determine how equal two objects are, NCD checks how easy it is to transform the two objects into each other.

**Tokenization** If AC2 can recognize the programming language it will use a lexer and a parser to tokenize the text file. "AC2 uses Antlr4<sup>4</sup> grammars to generate lexers (= tokenizers) and parsers for languages." (Freire, 2022b) Therefore it is relatively easy to add support for a new programming language.

#### 4.4.2 Preliminary results

In order to present the advantages of AC2, several projects were compared using the AC2 algorithm. As explained in section 4.4.1 AC2 comparison algorithm, the filters were used to select and group the submissions. After passing the project files, the algorithm analyzes them and creates an output window. The output that AC2 generates is mainly visual based. So AC "will create graphical representations of the degree of similarity between student submissions within a group". (Freire, 2022a). See figure 2 and figure 3.

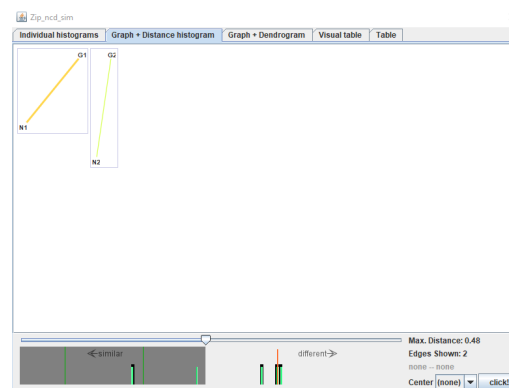


Figure 2 The graphical output of AC2 from milestone [0]

Figure 2 shows milestone 0 and how the different projects have copied from each other. This suggests that project N1 has maybe copied its code form G2 and N2 from G2.

Figure 3 illustrates that G1, N1 and N2 have probably copied from each other, since a correlation exists between the 3 projects. This again illustrates the advantage of AC2 over the other plagiarism detection programs, since it is easier to detect multiple correlations between projects.

Table 6 and table 7 show how far apart the individual projects are. This distinguishes AC2 from other plagiarism detection programs, as they often reflect the similarity in percent to the user.

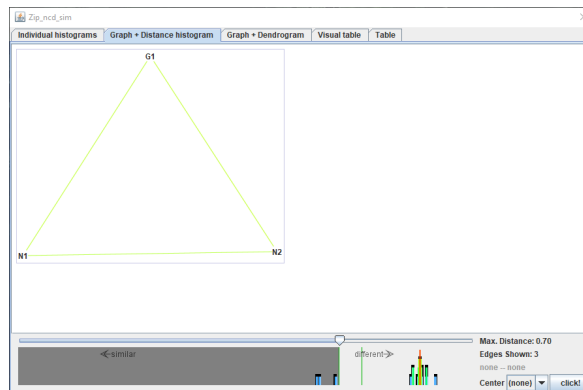


Figure 3 The graphical output of AC2 from milestone [4]

One	The other	Distance
ST2M0-Geber1	ST2M0-Nehmer1	0.2926
ST2M0-Geber2	ST2M0-Nehmer2	0.4609
ST2M0-Geber1	ST2M0-Nehmer2	0.6302
ST2M0-Nehmer2	ST2M0-Nehmer1	0.6705
ST2M0-Geber2	ST2M0-Geber1	0.6718
ST2M0-Geber2	ST2M0-Nehmer1	0.6811

Table 6 [Table of Milestone [0] distance between projects]

One	The other	Distance
ST2M4-Geber1	ST2M4-Nehmer2	0.6538
ST2M4-Geber1	ST2M4-Nehmer1	0.6603
ST2M4-Nehmer2	ST2M4-Nehmer1	0.6957
ST2M4-NoPlag2	ST2M4-Nehmer1	0.6809
ST2M4-Geber1	ST2M4-NoPlag1	0.8647
ST2M4-NoPlag1	ST2M4-Nehmer1	0.8657
ST2M4-NoPlag2	ST2M4-NoPlag1	0.8772
ST2M4-NoPlag1	ST2M4-Nehmer2	0.8792
ST2M4-NoPlag2	ST2M4-Nehmer2	0.8809
ST2M4-NoPlag2	ST2M4-Geber1	0.8823

Table 7 [Table of Milestone [4] distance between projects]

## 5 Conclusion

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