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ANALYZING MUTANT DENSITY AS A COMPLEMENTARY METRIC  
TO CYCLOMATIC COMPLEXITY FOR FINDING REFACTORING  
OPPORTUNITIES

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## Abstract

The complexity of code is a research field that has existed for multiple decades and has a large amount of existing work. Despite the large amount of existing work, it still remains a very active field due to its lasting importance in the current DevOps era. Within the context of DevOps, maintainability is an important term that is supported by testability for example. Certain complexity metrics have the ability to be related to testing effort needed which is why it remains relevant. One of the most researched metrics for estimating this effort is cyclomatic complexity. It has a long history of usage, even in the current DevOps tooling, despite there being critique on the metric on different aspects. This work tries to contribute to finding refactoring opportunities related to testing effort by working with a complementary metric for cyclomatic complexity to provide additional opportunities and a failsafe for false positives. This is done by positioning mutant density as a complementary metric, which measures code complexity based on the principles of mutation testing. Mutant density utilizes a flexible set of mutation operators to measure the amount of mutants which is used to define complexity. This different approach, which is still related to testing effort due to the principles it is based upon, provides a potential complementary metric which can expand on the need for finding refactoring opportunities. The contributions could be divided in these aspects: 1) Defining the correlation between cyclomatic complexity and mutant density, 2) Analyzing different metrics utilizing mutant density and cyclomatic complexity that provide deeper insight in code complexity and refactoring opportunities , and 3) Investigating the influence of the modular aspect of mutant density by selecting different mutation operators.

## Nederlandstalige Samenvatting

De complexiteit van code is een onderzoeksveld dat al meerdere decennia bestaat en dat ook een grote hoeveelheid gepubliceerde werken rond het onderwerp heeft. Ondanks de lange geschiedenis van dit onderzoeksveld, is er nog steeds zeer actief onderzoek naar dit onderwerp vanwege het blijvende belang in het huidige DevOps tijdperk. Binnen de context van DevOps is maintainability een zeer belangrijke term that onder andere ondersteund wordt door testbaarheid. Bepaalde complexiteit metrieken hebben de capaciteiten om gerelateerd te zijn aan de moeite die in testen gestoken moet worden, dit is een van de redenen voor de blijvende relevantie. Een van de meest onderzochte metrieken voor deze moeite te meten is cyclomatische complexiteit. Het heeft een lange geschiedenis van gebruik, zelfs in de huidige DevOps tooling ondanks het feit dat er op sommige vlakken kritiek is tegen deze metriek. Dit werk probeert een bijdrage te leveren aan het vinden refactoring opportuniteiten die gerelateerd zijn aan de moeite voor het testen door te werken met een complementaire metriek. Deze complementaire metriek gaat cyclomatische complexiteit ondersteunen in het vinden van nieuwe refactoring opportuniteiten en het vermijden van false positives. Dit gebeurt door mutatie densiteit te positioneren als een complementair metriek, deze meet code complexiteit aan de hand van principes die origineel uit mutatie testen komen. Mutatie densiteit gebruikt een flexibele verzameling van mutatie operatoren om te meten hoeveel mogelijke mutaties er aanwezig zijn in code, dit aantal wordt gebruikt om de complexiteit te meten. Deze aanpak verschilt zeer hard van cyclomatische complexiteit maar is nog steeds gerelateerd aan de moeite om te testen vanwege de oorsprong uit mutatie testen. Dit geeft een potentieel complementair metriek die voor uitbreiding kan zorgen aan de nood voor refactoring opportuniteiten. De bijdragen van dit werk kunnen onderverdeeld worden in deze aspecten: 1) Definieren van de correlatie tussen cyclomatische complexiteit en mutatie densiteit, 2) Analyse van verschillende metrieken die gebruik maken van mutatie densiteit en cyclomatische complexiteit om dieper inzicht te krijgen in refactoring opportuniteiten, en 3) Onderzoek naar de invloed van het modulaire aspect van mutatie densiteit door de selectie van verschillende mutatie operatoren.

## **Acknowledgments**

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# 1 Introduction

Software development tries to create software that is not only functional but also includes a factor of maintainability and adaptability. The landscape of software development on its own is ever-evolving, causing applications to have a constant growth in complexity. This increases the difficulty surrounding maintenance in the face of changing requirements, technological advancements and the constant pursuit of improvement. Software maintainability therefore emerges as a critical factor in achieving long-term success. It encompasses the ease with which a software system can be changed, updated and expanded upon without causing unwanted ripples through the existing code or influencing the stability.

At the core of maintaining software lies the practice of refactoring, a specific approach to improve the internal structure and design of code without changing the external behavior. This is a key factor to supporting the concept of maintainable code [1], enabling developers to correct flaws, reduce technical debt and optimizing performance. Being able to identify refactoring opportunities is a fundamental functionality needed to support this.

Finding refactoring opportunities is widely supported in the current DevOps era where there is much more emphasis on automation of quality assurance [2]. Popular industrial tooling such as CodeScene [3] and Sonarqube [4] all support different aspects of automated quality assurance. The given examples each utilize code complexity metrics to determine refactoring opportunities, providing simpler ways of increasing and managing maintainability.

The actionability of these tools lies within their own specific recommendations for code improvements. SonarQube, for example, identifies code smells, potential bugs, refactoring targets and many others. Codescene similarly offers insights by analyzing code complexity, coupling, change patterns and also refactoring targets. Despite these tools being adopted in many different industries, acknowledging the potential false positives that can occur in such tooling is very important [5, 6]. This could occur by putting too much weight on a singular complexity metric to define refactoring targets, causing suggested improvements that may not be relevant. The relevance of suggestions is however also dependent on the context of the project itself. Complete avoidance of false positives is difficult but combining complementary metrics can reduce the probability.

Within the DevOps context, testing is a critical component in ensuring the quality, reliability and the evolvability of a software product [7]. There are many rapid and frequent deployments with DevOps which makes testing even more crucial. It functions as a safety net that aids in capturing bugs and issues as they emerge during the development process. By integrating testing into the DevOps pipelines, it is possible to automate various testing processes such as unit testing, integration testing, performance testing and security testing. This can also be seen by the code coverage inclusion in tools such as the previously mentioned Sonarqube. On top of that, testing also provides a certain sense of confidence in the stability and robustness of the software. Based on these factors and within the context of DevOps, it is very interesting to study refactoring opportunities from the perspective of complexity metrics which estimate testing efforts. Using complexity metrics to refactor methods with the goal of simplifying the testing process, while the testing process has so much importance, provides a very interesting field of study.

One of the more known metrics which is able to estimate testing effort is cyclomatic complexity [8]. This metric is closely linked to the testing effort since the cyclomatic complexity can be seen as the minimum amount of tests needed to create full coverage, therefore directly representing the difficulty of testing [9]. Cyclomatic complexity does have a relatively large history of critique on a few different aspects, however most emphasis for this work is placed upon how well cyclomatic complexity measures maintainability and not the direct critique. Current measures for calculating maintainability still frequently incorporate cyclomatic complexity [10, 11].

Placing too much emphasis on cyclomatic complexity alone could lead back to the principle of false positives or obtaining incomplete information. A possible method to avoid this is utilizing a complementary metric.

This work will investigate how well mutant density [12] can be used as the complementary metric to provide additional insight into refactoring opportunities.

Mutant density is a complexity metric which also estimates testing effort based on an approach derived from mutation testing. It counts the amount of potential mutations in a method as the complexity. Doing this also allows some modularity by choosing the mutation operators used to create mutants. Mutant density will not only be positioned as a failsafe for false positives but also as a metric capable of finding refactoring opportunities which are unavailable when only using cyclomatic complexity.

The corresponding research questions behind this general claim can be formulated as:

- RQ1. Correlation - what is the correlation between cyclomatic complexity and mutant density?

Investigating the correlation has its importance on a few different levels. An extremely strong correlation between the two metrics would cause redundancy for one of them. Since mutant density has a higher computational complexity, this metric would be more easily seen as redundant. Claiming that these metrics can be used in a complementary manner also needs a certain amount of correlation. A very low correlation would make this impossible since they would not be able to verify each other while also providing new refactoring opportunities.

- RQ2. Complementary - how can we utilize mutant density to provide additional insight into refactoring opportunities? Is there anecdotal evidence supporting this?

This research question essentially has two main reasons why it's necessary. Firstly, it needs to be defined how these two metrics are to be used to obtain new insights. There are different ways possible of combining these metrics so finding the difference between these combinations is important. Secondly, it needs to be defined how the complementary claim can be investigated and how the results can be obtained surrounding this.

- RQ3. Fine Tuning - what is the impact of fine-tuning mutant density on correlation and complementary?

Since mutant density has an aspect of modularity based on choosing different mutation operators, creating variations based on this can influence the results. It is therefore interesting to investigate whether this could positively impact the ability of finding refactoring opportunities or support the complementary aspect.

The rest of this work will be structured as follows. In Section 2, there will be some background information on the most relevant metrics used in the analyses and about the tooling for finding refactoring opportunities. The main methods used in this work will be described in Section 4. This is followed by a showcase of the results combined with extensive discussion in Section 5. After this the threats to validity are discussed in Section 6. An overview of the related work is shown in Section 3. Finally the conclusions and potential work is given in Section 7.

## 2 Background

Due to the emphasis mostly being placed on code complexity metrics which also relate to the testing effort, this section will dedicate itself to covering the most important metrics in question for this work. There will also be some coverage of Sonarqube and Codescene on the aspects that are utilized within this work. Since this work emphasises the detection of refactoring opportunities, some fixed code examples will be used. This also allows for clear visualisation and contextualization of the issues this work addresses. These two Java functions will be used for the examples, the exact reasoning behind choosing these will become clear in further subsections. Both of these examples are obtained from the Spring library through methods used in this work.

---

```
private static int nextPowerOf2(int val) {
    val--;
    val = (val >> 1) | val;
    val = (val >> 2) | val;
    val = (val >> 4) | val;
    val = (val >> 8) | val;
    val = (val >> 16) | val;
    val++;
    return val;
}
```

---

Listing 1: nextPowerOf2 function from Spring dataset

---

```
private boolean isSockJsSpecialChar(char ch) {
    return (ch <= '\u001F') || (ch >= '\u200C' && ch <= '\u200F') ||
        (ch >= '\u2028' && ch <= '\u202F') ||
        (ch >= '\u2060' && ch <= '\u206F') ||
        (ch >= '\uFFFO') || (ch >= '\uD800' && ch <= '\uDFFF');
}
```

---

Listing 2: isSockJsSpecialChar function from Spring dataset

### 2.1 Source lines of code

Source lines of code is potentially one of the oldest complexity metrics. Originating from the time when FORTRAN and assembly where the most commonly used languages. At that time the development was also done through the usage of punched cards, where a single card often represented a single line of code. Therefore, measuring the complexity based on this was an obvious conclusion.

This metric can be divided in two general types: physical SLOC (LOC) and logical SLOC (LLOC) [13]. The exact definitions might vary but the core concept remains the same. Physical SLOC counts all the lines in the source code without including the commented lines. Logical SLOC tries to measure based on the amount of executable statements, their specific definitions are however coupled to the programming languages.

In essence, this metric is very simple. The counting can be easily automated and the metric is very intuitive because it can be seen directly. Being one of the oldest metrics, it also has a lot of data available. Within this work this metric will be used alongside cyclomatic complexity and mutant density to capture the influence of function sizing. The influence of this metric on testing effort is not directly the most emphasised aspect. Still, functions with larger amount of lines tend to be more difficult to test. This aspect will be discussed further when discussing correlation between the metrics. One of the downsides to source lines of codes is effect of different programming languages on the same functionality.

The difference over languages can be clearly seen in Figure 1, where the same functionality over different languages changes the lines of code. Although it will be argued later on that the effect of this is limited for the proposed research

BASIC	C	COBOL
<pre>PRINT "hello, world"</pre>	<pre>#include &lt;stdio.h&gt;  int main() {     printf("hello, world\n"); }</pre>	<pre>identification division. program-id. hello . procedure division. display "hello, world" goback . end program hello .</pre>
Lines of code: 1 (no whitespace)	Lines of code: 3 (excluding whitespace)	Lines of code: 6 (excluding whitespace)

Figure 1: The effect of different languages on source lines of code computation when implementing the same functionality

questions, it remains relevant to keep in mind. In depth analysis of this particular issue will not be covered extensively within the scope of this work to not take away from the main research questions.

## 2.2 Cyclomatic complexity

Cyclomatic complexity is one of the most known complexity metrics and it will have a very important role throughout this whole work. Since its introduction in 1976 by Thomas J. McCabe [8], it has since seen quite some usage. Even industrial tooling frequently relies on this metric for complexity measurement [3,4,14].

The way cyclomatic complexity works is based on the measurement of the control flow of a program. This means the metric counts the amount of unique and independent paths through a program's source code. The formula for the calculation is as follows:

$$V(G) = E - N + 2 \quad (1)$$

where  $V(G)$  is the cyclomatic complexity,  $E$  the amount of edges in the control flow graph, and  $N$  the number of nodes in the graph. Since each program has a so called "entry" and "exit" points, the formula includes a "+2" factor.

---

```
private boolean example1(int b, int c) {
    int a = 10;
    if ( b > c ) {
        a = b;
    } else {
        a = c
    }
    System.out.println(a);
    System.out.println(b);
    System.out.println(c);
}
```

---

Listing 3: Example function to showcase cyclomatic complexity calculation

If we visualize this *example1* Function 3 on p. 12 as a flow graph we obtain this:

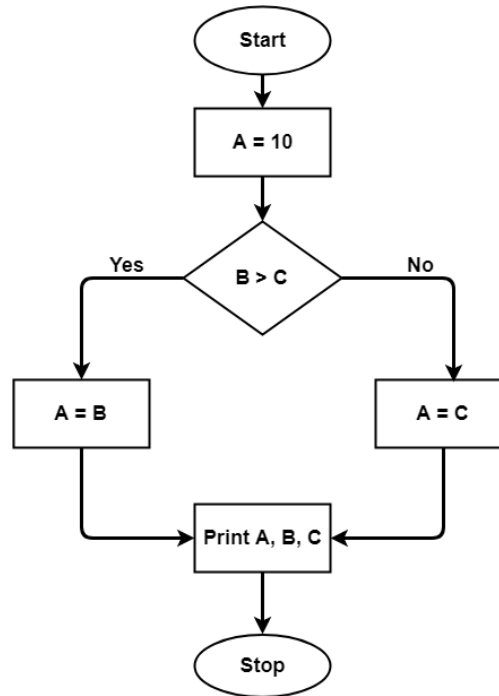


Figure 2: Flow graph representing the independent paths through function1

If we apply the previous formula 1 we would obtain a cyclomatic complexity of two for that function.

In general there are multiple different statements that can increase the cyclomatic complexity.

- Conditional statements: **if**, **else if**, **switch**
- Loops: **for**, **while**, **do-while**
- Boolean operators: **&&**, **||**, **!**
- Exception handling: **try**, **catch**, **except**

From this list, there are a few key takeaways. Not every single one of these statements or operators are available for every single programming language. This might cause small variation in cyclomatic complexity for the same functionality in different languages. Still, compared to how this influences source lines of code, the difference in this case is a lot smaller.

One of the advantages that something like cyclomatic complexity has over source lines of codes is the closer relation with the exact functionality. Since it does so by using statements that provide clear division in methods, it looks and feels very intuitive. This is also one of the reasons it is still used so frequently.

When looking back at the relationship between complexity and testing, the premise is that you need a single test for each independent path through a method. Therefore, cyclomatic complexity would dictate the amount of tests needed. This is not wrong, but it ignores one part of testing. The difficulty of writing tests is not always the amount of tests needed, but also how difficult it is to test all of the functionality. This is something that cyclomatic complexity does not directly cover.

There is also categorisation or threshold as to how cyclomatic complexity should be interpreted [15].

- 1 - 10 Simple procedure, little risk
- 11 - 20 More complex, moderate risk
- 21 - 50 Complex, high risk
- > 50 Untestable code, very high risk

This categorisation is not fixed however. In NIST [9] it is mentioned that there is sufficient support for this categorisation for general application but that variation might still be possible and is not necessarily bad depending on the provided context. They do however, never specify any real indication on why this categorisation works and which studies back these findings. Despite this, the threshold of 10 is often used as point where refactoring or splitting the method is appropriate. Since thresholding on its own is a core aspect of defining refactoring opportunities, this remains highly relevant.

## 2.3 Mutant Density

Originally complexity metrics are not tied to different defect types. Mutant density is introduced as a complexity metric that pinpoints statements more likely to be at risk of a fault. As such, creating a metric for increasing the maintainability of the software. The base concept of this already exists within the field of mutation testing.

Mutation testing injects intentional faults in the code and runs these mistakes through the tests to verify if the tests detect the introduced mistakes. Every injected mistake creates another mutant. The exact mutations created depends on the types of faults that are injected, so there is a certain amount of variation within this. These types of faults are called the mutation operators and allow for fine tuning the usage of mutant density.

The general workflow of mutation testing can be seen in Figure 3.

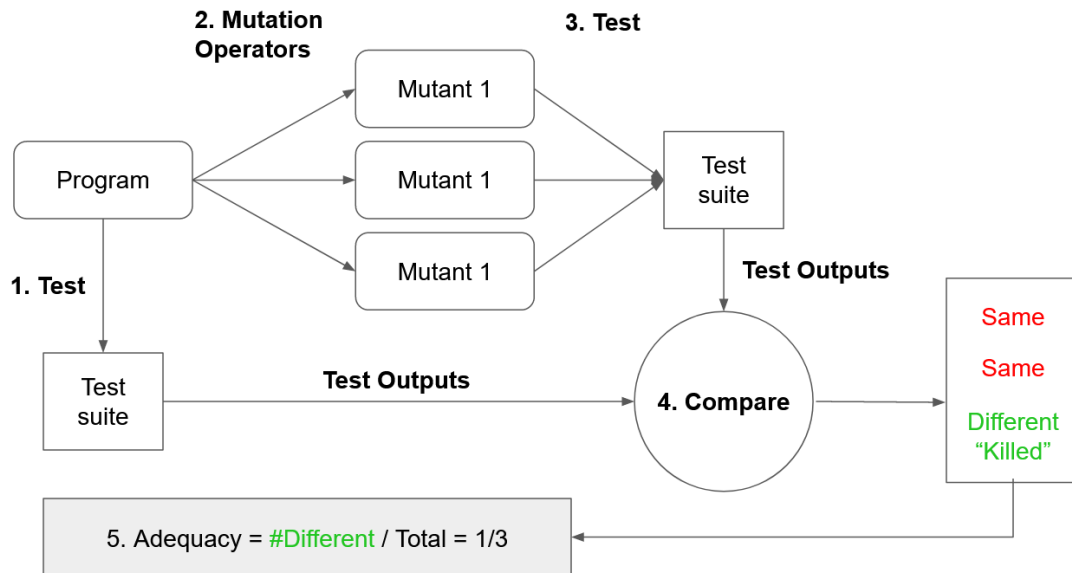


Figure 3: General workflow of mutation testing

Mutation testing was originally introduced in 1975 in the work "The design of a prototype mutation system for program testing" [16]. It is a technique provided for the evaluation of the effectiveness of a software testing suite. This is, as previously explained, done by measuring the ability to detect bugs and faults in the program. The work argues that mutation testing could provide a more accurate measurement of the effectiveness of tests. Especially in comparison to code coverage analysis because this metric can easily provide wrong insights. Mutation testing functions very differently and also directly measured the ability of the test suite to detect faults.

This is the concept that is used as a foundation for mutant density, which is the amount of mutants that can be generated for each line of code. The average mutant density for a file is thus defined by the amount of mutants divided by the number of relevant lines. Comments and whitespaces are ignored, just like source lines of code.

Lets take the previously described *nextPowerOf2* Function 1 on p. 11, this function has a larger mutant density of 12 due to its many operations. Based on cyclomatic complexity alone, it would be placed below the threshold of 10 and not detected as something to worry about. The discrepancy between those two and simply looking at the functions show clear signs of potential improvement.

### 2.3.1 Mutation operators

Before wrapping up this section, a recap will be given on potential mutant operators. This is necessary since it is a part of the modularity of mutant density. It should be noted that there is also variation possible between the types of mutations for different languages. Currently the focus will not be on addressing this but more on the available types of mutation and what they represent.

- Relational Operator Replacement (ROR): replaces a single operand with another operand, these include  $<$ ,  $<=$ ,  $>$ ,  $>=$ ,  $=$ ,  $!=$ ,  $true$ ,  $false$ . Depending on the implementation this could mean  $x < y$  can be mutated to:  $x <= y$ ,  $x! = y$ ,  $false$ . Although more mutations are possible, [17] proved that these three cover all other possibilities.
- Arithmetic Operator Replacement (AOR): replaces a single arithmetic operator with another operand. these include  $+$ ,  $-$ ,  $*$ ,  $/$ ,  $\%$
- Logical Connector Replacement (LCR): replaces logical operand with the inverse, these include  $||$ ,  $\&\&$
- Constant Replacement (CR): replaces literal values with other constants.

This only showcases the most basic examples, it should also be clear that not every mutation can be applied to every language in the same way. Take python for example, when looking at LCR, python also has the option to replace this *and/or*. The differences can also be larger than this, depending on the software, mutations may also be applied to more specific statements such as *try, except* or *switchcases*. Not every language includes this thus the mutations varies quite a lot based on this. By using different mutation operators, mutant density can go in depth in its modularity. Some portions will be covered in later sections but a full analysis on this is outside the scope of this work.

## 2.4 DevOps tools

This section will provide an overview of the tooling used such as: CodeScene and Sonarqube. This tooling will mostly be used for verification and strengthening of the anecdotal evidence at the end of the results. Both of these tools have their own specific ways of finding refactoring targets which aligns with the goal of this thesis. There are more tools available with similar functionality but the scope is limited to these for their easy usage and the inclusion of cyclomatic complexity.

### 2.4.1 CodeScene

Within the interface of CodeScene, the first step is connecting through the chosen git platform where your project is located. A project can be selected and there will be automatic analysis done after the selection. The functionality provided by CodeScene is broad so the scope is limited to functionality relevant to this specific work. On the main

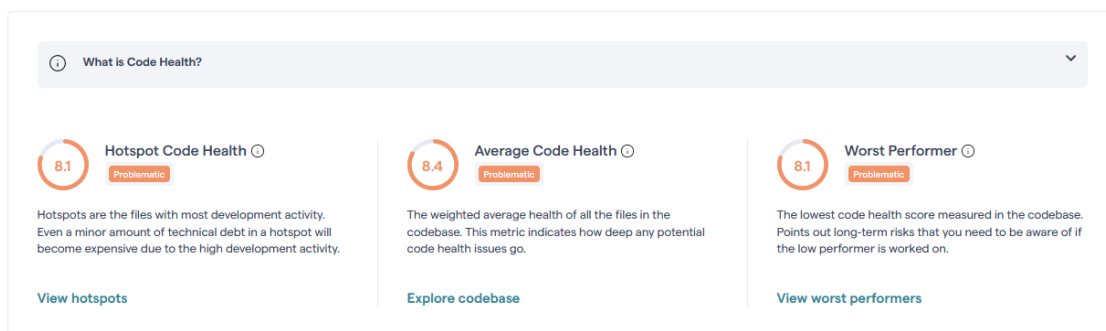


Figure 4: CodeScene main screen

screen, the first important part of information is given in the form of Figure 4. Of this code health is the most important aspect since it directly correlates to the maintenance costs. This cost is determined by a combination of both properties of the code as well as organizational factors. These include low cohesion, presence of god classes, lines of code complex methods, nested complexity and many others. One of the complexity metrics used in the case of code complexity is cyclomatic complexity.

When further inspecting code health, code scene provides an overview of code health hotspots as seen in

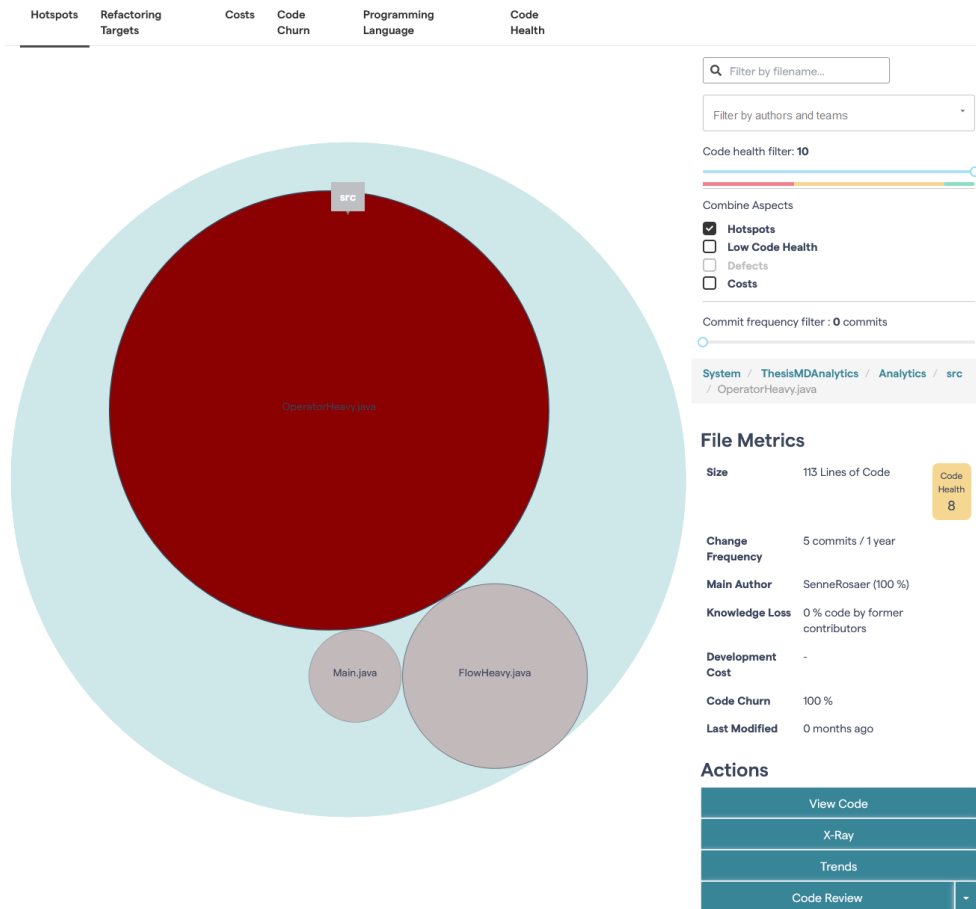


Figure 5: CodeScene hotspot overview

Figure 5. This allows finer details of the different files which cause the most impact on the current code health. It also provides a few actions that can be executed on the most important files, for the RG2. the emphasis will be on the X-Ray action. The results on a very small example project are given in Figure 6. It clearly highlights specific functions in different colors based on a few different factors. Certain functions such as `f1` in Figure 6, are also flagged as refactoring targets. It chooses this based on a complex conditional present instead of only on relying on cyclomatic complexity. Still, the *areBoxingCompatible* is indicated as the most complex function which is based on cyclomatic complexity, meaning this indication makes less usage of complementary metrics.

## 2.4.2 SonarQube

Just like CodeScene, Sonarqube provides a large amount of functionality so the scope will again be limited to the necessary parts. The main setup of SonarQube is a bit more involved and differs between programming languages, pipeline usage and other factors so it will not be discussed. The primary overview given by Sonarqube is shown in Figure 7. It mostly provides information about technical dept, bugs, code smells, code coverage and duplication. More specific information can however be found when moving over to the measures tab. In Figure 8 this is shown and the most important parts can be seen. Within SonarQube there is a clear separation between maintainability and complexity. Testing effort is more closely linked to the complexity metrics for this work which is a part of the maintainability. Sonarqube calculates maintainability only based the time needed to fix the code smells. As for complexity, this is separated into cyclomatic complexity and cognitive complexity. The definitions of cyclomatic complexity remains the same and it is mostly used to indicate the minimum amount of tests needed to have full test coverage. The cognitive complexity indicates how hard it is to understand the code's control flow. These metrics are however not calculated for each method but for each file, therefore refactoring targets are not directly defined but rather indicated through the code smells.



Hotspots

Internal Change Coupling

Structural Recommendations

Change Frequency Distribution

Function name	<div><div></div><div></div></div>	<div><div></div><div></div></div> Change Frequency	<div><div></div><div></div></div>	<div><div></div><div></div></div> Lines of Code	<div><div></div><div></div></div>	<div><div></div><div></div></div> Code Review	<div><div></div><div></div></div>	<div><div></div><div></div></div> Function Complexity	<div><div></div><div></div></div>	Actions
f1		1		16		• Complex Conditional		4		<div><div></div><div></div><div></div></div>
areBoxingCompatible		0		40		• Complex Method • Bumpy Road Ahead		14		<div><div></div><div></div><div></div></div>
f1_refactored		0		15		• Complex Conditional		4		<div><div></div><div></div><div></div></div>
nextPowerOf2_refactored2		0		10				2		<div><div></div><div></div><div></div></div>
nextPowerOf2_refactored		0		10				1		<div><div></div><div></div><div></div></div>
nextPowerOf2		0		10				1		<div><div></div><div></div><div></div></div>
f2		0		5		• Complex Method		10		<div><div></div><div></div><div></div></div>
val_operation		0		3				1		<div><div></div><div></div><div></div></div>
f1_operation		0		3				1		<div><div></div><div></div><div></div></div>

Figure 6: CodeScene precise file information

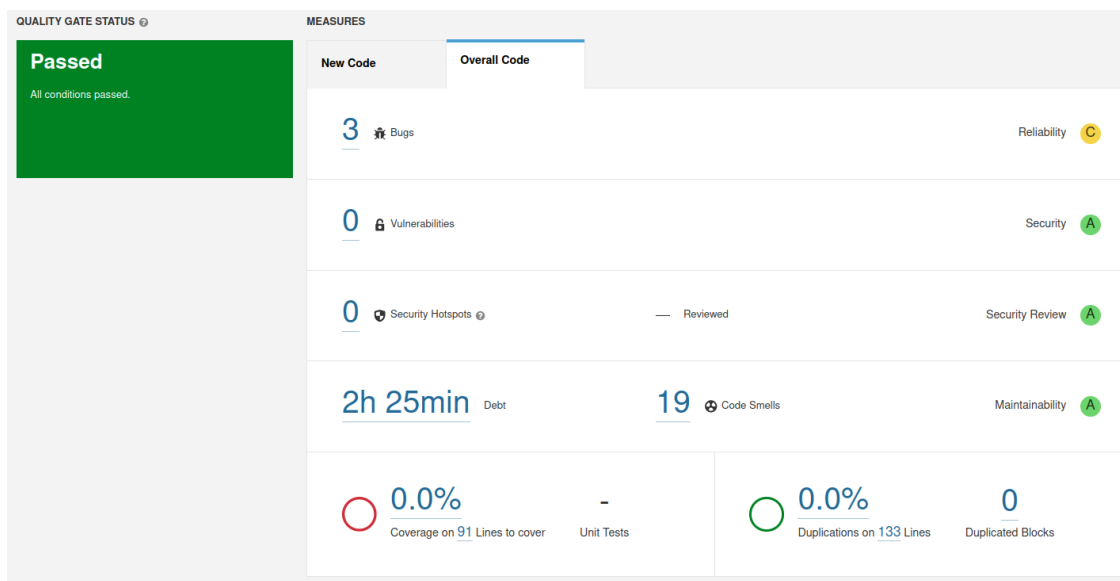


Figure 7: Sonarqube overview

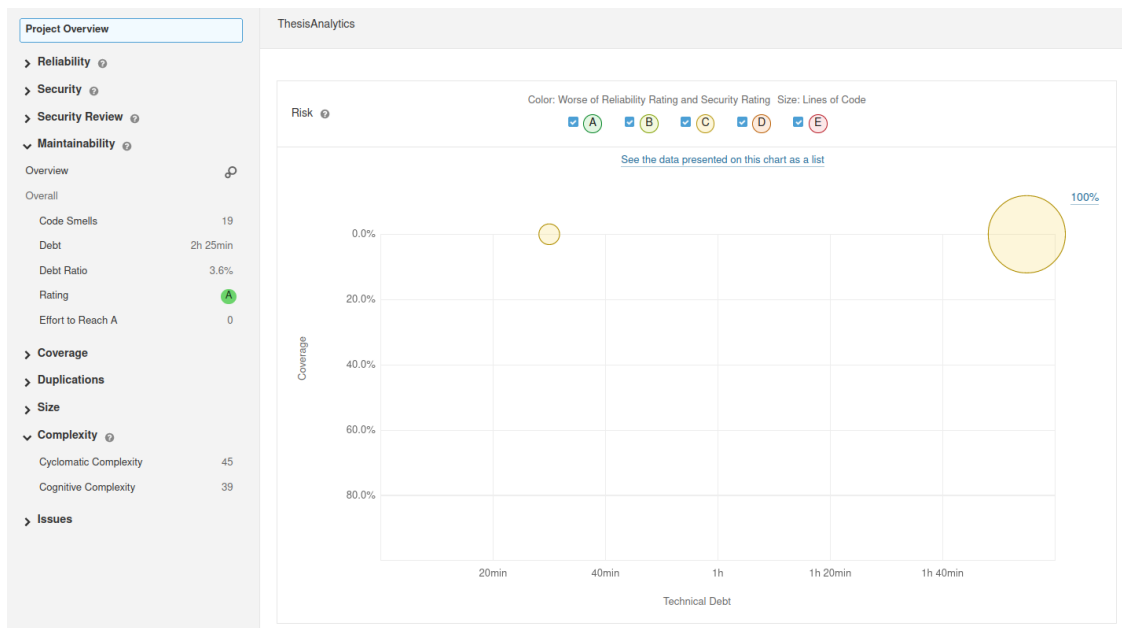


Figure 8: Sonarqube precise information

### 3 Related Work

Within the field of software engineering, various studies have focused on identifying refactoring opportunities and analysing the correlation between different software metrics. This section will provide an overview of literature on these subjects which contain relevant information surrounding the methods used in this work and the reasoning behind these.

#### 3.1 Correlation Analysis of Software Metrics

Understanding the correlation between software metrics is crucial for efficient utilisation of these metrics for measuring different software aspects. Metrics that are correlated on such a high level can cause redundancy. Finding this redundancy allows for more careful selection of software metrics to efficiently provide insights. This is one end of the spectrum of correlation which has its relevance within this work. Too high correlation with mutant density would cause mutant density to be redundant because of the higher computational complexity. It would also directly impact the complementary factor of RQ2. There is also the possible case of too low or almost no existing correlation. Lack of correlation would indicate that the metrics measure different aspects of the software. The goal of this work is complementary metrics which find refactoring opportunities based on testing effort. If these metrics are too independent, complementary usage is also out of question. Although correlation mostly relates to RQ1., complementary usage of mutant density and cyclomatic complexity in this work also includes usage of source lines of code. This causes correlation to remain relevant to avoid combining metrics which do not combine properly.

The work of [18] is an extensive work on the correlation between source lines of code and cyclomatic complexity to analyse redundancy. Other works covering the same goal exist but it remains important since the methods used correspond to the methods within this work. They propose correlation analysis through visualisation methods including boxplots, scatterplots and distribution plots. To quantify the strength and direction of the visual exploration, the Pearson correlation coefficient was employed. These methods are employed in a similar way for analysis surrounding RQ1.. Their final conclusions reject redundancy based on lower correlation of a larger general dataset compare to other works and the higher variation of cyclomatic complexity on higher source lines of code. Using the Pearson correlation coefficient is also applied in other works such as [19, 20] while visualising through boxplots and scatterplots is also applied in works such as [21].

#### 3.2 Refactoring Opportunities

Due to the importance of finding refactoring opportunities to improve different aspects of maintainability in a DevOps context, various researchers have covered different possibilities for locating those refactoring opportunities. Variation is possible within refactoring opportunities depending on the context of refactoring. Refactoring opportunities are not defined through singular methods but often through a combination of code smells [22–24]. These include duplicate code, god classes, complex conditionals, long parameter lists, complex code and a multitude of other possibilities. Every single one of these code smells could indicate potential refactoring even though it is not always coupled directly to a complexity metric or a certain threshold.

Complex code is however listed as a code smell which is based on cyclomatic complexity. Due to cyclomatic complexity being a complexity metric accumulated over a complete method, methods to decide refactoring opportunities need to be defined. Other code smells such as complex conditionals rely on the presence of the complex conditional and not a certain amount of occurrences to provide refactoring opportunities. For cyclomatic complexity, this mostly falls back to refactoring when it exceeds a threshold of 10 [25]. This threshold is even incorporated within NIST [9] although there is no provided method on how this exact threshold was derived. Mutant density within this concept lies close to cyclomatic complexity, in which a threshold would be needed to define a clear refactoring target. Even though 10 is chosen for cyclomatic complexity, within NIST [9], it is mentioned how variation might be possible within an organization or project depending on the argumentation.

This indicates that despite the existing empirical evidence for this threshold, the context surrounding the usage could indicate variation. Finding thresholds is outside of the scope of this work due to its complexity and needed scale. The chosen approach is using z-score to define outliers for various complexity metrics which are used to define refactoring opportunities. Using this outlier detection method is not conventional within the context of finding refactoring opportunities due to its limited usage on small projects. Despite this, it allows for a flexible threshold to be defined for each project without the direct necessity of a larger scale empirical analysis.

A reoccurring pattern within similar work is the limitation on programming languages covered and the potential effects on the results. Works such as [22, 25] are mostly limited to Java due to existing integration of refactoring

patterns and metric analysis in Eclipse. Although most visual examples within this work are based on Java as well, there will be analysis across other languages such as C, C++ and Python. This covers more potential variation possible when generalizing specific metrics. Due to the broader incorporation of different languages, the sizing of the datasets within each language is more limited.

## 4 Method of Analysis

The method of analysis takes a very step-wise approach. As most of the analysis is done with a combination of source lines of code, cyclomatic complexity and mutant density there is an initial need for this data. The gathering of this is not as one-sided due to difficult tooling and dataset generation. With the data in place there are different types of analysis that are done with each their own importance for the different research questions. Correlation analysis between the metrics is the first step in this process. Further analysis relies on finding patterns in how outliers for specific metrics correspond to testing effort and potential refactoring opportunities. This is done on a few different levels which are explained within the following sections. Outlier detection is at the core of finding refactoring opportunities and support for mutant density as a complementary metric. DevOps tooling will mostly be used alongside other information to highlight the applicability and provide anecdotal information.

### 4.1 Data collection

Table 1: Tools used for the metrics for each programming languages

	<b>Java</b>	<b>Python</b>	<b>C/C++</b>
<b>Cyclomatic complexity</b>	Chaosmeter	Mccabe	Metrix++
<b>Mutant density</b>	Chaosmeter	Mutpy	Dextool mutate
<b>Source lines of code</b>	Chaosmeter	Custom	Metrix++

The process of gathering data contained quite some hurdles due to difficulty with tooling. Since the goal is to obtain source lines of code, cyclomatic complexity and mutant density there is the need for a tool which supports all of these. For java this is not an issue since Chaosmeter [26] covers all of these metrics. Other languages do not have this luxury, since some analysis compare between languages this cannot be avoided.

For C and C++ the same tooling can be used. The calculation of the amount of mutations is done through Dextool mutate [27]. This tool was not created with the intention of just calculating mutant density but has more emphasis on mutation generation and execution. Due to this emphasis, Dextool is also not an out of the box solution. To gather the amount of mutations, an additional parser was written to obtain it from the output it gives. Dextool provides an html output with the mutations visualized, it would also be possible to alter the source code to provide this functionality, but writing the parser was an easier solution. The source lines of code and cyclomatic complexity are generated by Metrix++ [28]. Metrix++ keeps functions by the start and end line, therefor the parser needs to be able to read this information from the html files.

Python also has a few different tools to extract the data. Cyclomatic complexity is calculated by McCabe [29], the source lines of code are calculated by a custom tool and the mutations are calculated by mutpy [30]. Existing tooling for source lines of code mostly focuses on calculating it for a single method. This work does analysis for each method separately, therefor something additional had to be provided.

Again, the tooling used for mutations have a pure emphasis on the testing which is the same with mutpy. Therefore, mutpy had to be altered since the base version was not able to just generate and count mutants for each function. The mutations were originally coupled to the written tests, meaning any functionality which wasn't tested did not get mutated. This functionality was overwritten such that the presence of tests was no longer a necessity. The code was also altered in a way to provide the mutant, their corresponding function and file in structured manner.

Although this provided us with all of the data, there are still a few flaws that should be discussed. Since the tooling is not obtained from a single provider, there might be variations in implementation which could deliver small deviations in results. As was mentioned in the background information on mutation operators, there are differences between languages which is reflected in differences in the tooling. How source lines of code are counted could also vary across the tooling due to the lack of standardization.

Across the languages there are also issues regarding the projects used. Although dataset generation will be covered in a later section, there are some issues that are coupled with the metric extraction. For Java, the metrics are obtained in a static manner, this implies that there is no building of any sort included. Python allows for this to happen but the implementation for Mutpy does not completely support this. For example, loading a project that includes usage

of a GPU, forces you to have one working on your machine. This forces certain workarounds which makes the tooling more flawed than what would be optimal.

C and C++ have similar issues, Dextool needs to build the project before the mutant generation can be applied. This not only consumes quite some time, it also forces you to have a system that can build the project. With older C or C++ projects this is not always trivial.

## 4.2 Dataset composition

Table 2 gives an overview of all the projects used, their methods and the size. Note that for the method count and the nature of this paper, only methods which include at least a single possible mutation are included. The goal was to have a certain amount of variation within the projects sizes and also have some difference in functionality. Python for example, contains some projects for API's and frameworks while also having some projects focusing more on mathematical problems. Java does have Spring as it's largest project which is much larger than the largest projects for other languages. This is not ideal since it might provide certain imbalance. Equally large projects for other languages are also difficult to obtain due to more errors occurring during the generation of this data. Dextool often outputs that it skips certain methods or files due to internal issues so this also impacts the size of the resulting data.

Table 2: Project used for each language and their basic information

Language	Project Name	Method Count	SLOC	Github repo
Java	EventBus	157	1813	<a href="https://github.com/greenrobot/EventBus">https://github.com/greenrobot/EventBus</a>
Java	FastJson	2030	48479	<a href="https://github.com/alibaba/fastjson">https://github.com/alibaba/fastjson</a>
Java	Mockito	741	9805	<a href="https://github.com/mockito/mockito">https://github.com/mockito/mockito</a>
Java	Rebound	148	2228	<a href="https://github.com/facebookarchive/rebound">https://github.com/facebookarchive/rebound</a>
Java	Socket	67	977	<a href="https://github.com/socketio/socket.io-client-java">https://github.com/socketio/socket.io-client-java</a>
Java	Spring	8701	110078	<a href="https://github.com/spring-projects/spring-framework">https://github.com/spring-projects/spring-framework</a>
Python	BioPython	687	12984	<a href="https://github.com/biopython/biopython">https://github.com/biopython/biopython</a>
Python	Ansible	1536	34892	<a href="https://github.com/ansible/ansible">https://github.com/ansible/ansible</a>
Python	FastApi	28	752	<a href="https://github.com/tiangolo/fastapi">https://github.com/tiangolo/fastapi</a>
Python	Google Images	20	703	<a href="https://github.com/hardikvasa/google-images-download">https://github.com/hardikvasa/google-images-download</a>
Python	Photon	44	453	<a href="https://github.com/s0md3v/Photon">https://github.com/s0md3v/Photon</a>
Python	Scrapy	700	7993	<a href="https://github.com/scrapy/scrapy">https://github.com/scrapy/scrapy</a>
Python	Qbittorrent	65	356	<a href="https://github.com/v1k45/python-qbittorrent">https://github.com/v1k45/python-qbittorrent</a>
Python	Flask	131	2248	<a href="https://github.com/pallets/flask">https://github.com/pallets/flask</a>
C++	Duckdb	1420	35366	<a href="https://github.com/duckdb/duckdb">https://github.com/duckdb/duckdb</a>
C++	OpenCv	1678	40100	<a href="https://github.com/opencv/opencv">https://github.com/opencv/opencv</a>
C++	PrusaSlicer	226	7310	<a href="https://github.com/prusa3d/PrusaSlicer">https://github.com/prusa3d/PrusaSlicer</a>
C++	Transmission	521	11399	<a href="https://github.com/transmission/transmission">https://github.com/transmission/transmission</a>
C	Curl	438	19464	<a href="https://github.com/curl/curl">https://github.com/curl/curl</a>
C	LibrdKafka	283	7977	<a href="https://github.com/confluentinc/librdkafka">https://github.com/confluentinc/librdkafka</a>
C	Redis	1501	38523	<a href="https://github.com/redis/redis">https://github.com/redis/redis</a>
C	Mbedtls	842	34625	<a href="https://github.com/Mbed-TLS/mbedtls">https://github.com/Mbed-TLS/mbedtls</a>
C	SoftEthervpn	604	15084	<a href="https://github.com/SoftEtherVPN/SoftEtherVPN">https://github.com/SoftEtherVPN/SoftEtherVPN</a>

### 4.3 Correlation analysis

The analysis of correlation includes a visualisation part and a numerical verification part. The visualisation contains boxplots and scatterplots which are created with Seaborn using Python. Scatterplots are in place for every possible pair between cyclomatic complexity, source lines of code and mutant density. Boxplots are only used for pairs including source lines of code. The visualisation is supported by utilizing the Pearson correlation coefficient which is also computed in the same Python scripts through Scipy.

### 4.4 Outlier detection

Outlier detection is based on the calculation of a z-score over the distribution of different metrics and selection a cutoff value. The z-score can be given by  $Z = \frac{X-\mu}{\sigma}$  where X is the value we want to standardise,  $\mu$  the mean and  $\sigma$  the standard deviation. This is done through Scipy which for a set of values is able to calculate the z-score for each value compared to the mean and standard deviation of that set. The set of values in question contains the results of a metric for each method in a dataset. For example, this could be a set of the cyclomatic complexity or mutant density for each method. This set of z-score is also visualised through Seaborn to determine potential cutoff values. The exact values and reasoning behind the chosen values are discussed in Section 5.

Initial analysis is done on outliers calculated on the average cyclomatic complexity and average mutant density.

$$AverageMutantDensity = (MD/SLOC)$$

$$AverageCyclomaticComplexity = (MD/SLOC)$$

These metrics are calculated for each method within a dataset, out of these a mean and standard deviation is derived. The average mutant density and cyclomatic complexity are simply the previously described metrics divided by the source lines of code of the method.

#### 4.4.1 Scaled outliers

While looking for outliers and refactoring targets, using something such as average mutant density or average cyclomatic complexity leaves some potential issues.

Two methods might have the exact same average mutant density due to the relation of mutants to the source lines of code. However, if one of these functions is much longer this indicates a larger issue than the smaller function. A small function with a bad composition might be a choice to encapsulate the problematic code. Although it does not excuse the design of the function, the same bad composition for a larger method would have more direct influence on the maintainability of the code. Therefore, they should be more easily filtered or have a score that represents this better.

$$ScaledAverageMutantDensity = (MD/SLOC) * (MD + SLOC)/100$$

$$ScaledAverageCyclomaticComplexity = (CC/SLOC) * (CC + SLOC)/100$$

This applies a multiplication to the previous formula that takes both the size of the source lines of code and mutant density into account. This divided by a factor of 100 to limit the influence of this multiplication. The chosen factor of 100 is not based on specific empirical evidence and would need more verification, for the current use cases it is sufficient however. The same is done to obtain a scaled version of the average cyclomatic complexity. These formulas are not based on existing work but are derived from the intermediate results and issues that needed to be resolved.

#### 4.4.2 Cross metric

The previous ways of handling outlier detection, only focused on cyclomatic complexity and mutant density separately. They provide some initial insight but the main focus of RQ2. is investigating how cyclomatic complexity and mutant density can complement each other.

There are a few ways to look at relations between cyclomatic complexity and mutant density. The first one is using the previous outliers and applying set theory to look at the intersection and dissections. This can be visualized as Figure 9. The similarities between metrics has been a recurring topic until now, this will also remain so for the rest of this work. The dissections contain useful information due to the fact that only a single metric captures it as an outlier. Depending on the code contained in those outliers, it could show methods that are not noticed through a singular metric but still contain refactoring opportunities. This part is of utmost importance in terms of showing the

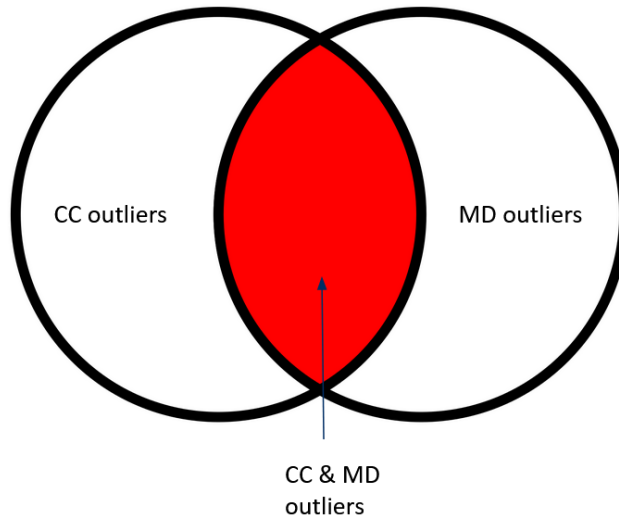


Figure 9: Visualisation of set theory application for RQ2.

additional functionality of mutant density. There needs to be sufficient functions to showcase that the additional outliers are also complex but that will be covered in the results.

The second way is by using a metric that uses the distance between the cyclomatic complexity and mutant density of a method.

$$AverageCC\&MDdistance = (CC - MD)/SLOC$$

If the outliers are calculated based on this metric it shows which functions have an abnormal difference between the cyclomatic complexity and the mutant density. Just like the previous metrics, there could be the introduction of some sort of ratio or scaling to take the sizing into account.

$$ScaledAverageCC\&MDdistance = (CC - MD)/SLOC * (MD + CC + SLOC)/100$$

Using this metric only functions to a certain extent if both cyclomatic complexity and mutant density have similarities. If there is completely no correlation than the difference between these two metrics would provide very random data. Therefore, the correlation analysis has its importance on supporting this analysis.

#### 4.5 Mutation operator fine-tuning

The computation of mutant density allows for a selection of mutation operators to be used to allow fine-tuning of the metric. Using all possible combinations for analysis is outside of scope.

However, if outliers are calculated for every single type of mutation operator separately there are still feasible analyses. This means that that instead of calculating mutant density with a set of mutation operators, there is a separate run for each mutation operator.

There are two ways of looking at this new data. The first one is looking at outliers that occur for each mutation operator separately. This implies that there are methods which have a very large variety of possible mutations spread out over different mutation operators, on top of that they are outliers in each case. The second possibility are outliers which are only outliers for a single mutation operator. They might not contain enough mutations to be a general outlier, but they would have enough when tweaking the mutation operators.

#### 4.6 Dataset comprised of a single project or multiple projects

Comparing the influence of calculating outliers over a single project or over a set of projects is an important aspect of potential generalisation. Verification is done by calculating outliers for the complete Java dataset, meaning it is based on a more general mean and standard deviation. Followed by outlier calculation on a singular project, Spring in this case. It can then be visualised if the larger dataset finds more or less outliers for Spring through a histogram.



#### 4.7 Selection of anecdotal evidence

Selecting anecdotal evidence as results for the analysis is done through selecting the highest z-scores, the biggest outliers in other words. The biggest outliers do not always show the patterns which cause outliers the best. Therefore, there is some personal influence in the selection to better show the potential and which patterns this show. Despite this, every given function is an outlier that has higher z-score than most other outliers.

#### 4.8 Overview of formulas

Recap of formulas and adding a name by which they will be referred to from now on.

<b>Metric</b>	<b>Formula</b>
Average Mutant Density	$(MD/SLOC)$
Average Cyclomatic Complexity	$(CC/SLOC)$
Scaled Average Mutant Density	$(MD/SLOC) * (MD+SLOC)/100$
Scaled Average Cyclomatic Complexity	$(CC/SLOC) * (CC+SLOC)/100$
Average CC & MD distance	$(CC-MD)/SLOC$
Scaled Average CC & MD distance	$(CC-MD)/SLOC * (MD+CC+SLOC)/100$

Table 3: Recap of the most important formulas which are used for outlier detection

## 5 Results

The following section will discuss the results of the analysis done to obtain answers for the provided research questions. The sections corresponding to RQ1. and RQ2. mostly cover the results based on the Java dataset. Java is chosen as a main focus since there is only a single tool used for calculating all the metrics instead of a combination of tools. This provides more certainty for consistent results. The Java dataset is also the largest dataset across the different languages and compared to languages such as C and C++, the code examples are less technical. The analysis was still executed over all of the different languages but these results are only covered whenever they deviate from the obtained Java results.

### 5.1 RQ1. Correlation

#### RQ1. Correlation - what is the correlation between cyclomatic complexity and mutant density?

Correlation can be calculated between all possible pairs of source lines of code, cyclomatic complexity and mutant density. Although the data on all of these pairs is generated, for RQ1. the only relevance is between cyclomatic complexity and mutant density. Pairs including source lines of code contain relevance for RQ2. and will therefore only be covered in the corresponding section, the methods used will remain the same.

Correlation between cyclomatic complexity and mutant density is necessary for avoiding redundancy and supporting the complementary claims of RQ2.. The results will be divided in two general steps, visualisation and pearson correlation. Although both directly apply to the same aspects of the correlation, pearson correlation is provided to avoid potential confusion surrounding the results of specific visualisations.

#### 5.1.1 Step 1.1 - visualisation of correlation

The correlation can be visualized in different ways but this work will focus itself to boxplots and scatterplots. The data that the graphs are created upon will be specified in the captions of the images since the following parts will contain some variation. For RQ1., only scatterplots are used since the boxplots are based on an inclusion of source lines of code.

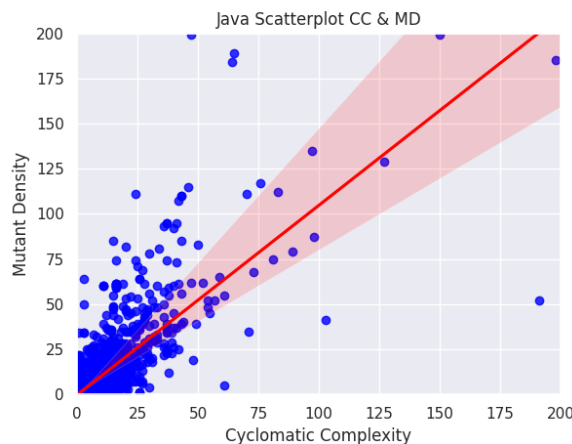


Figure 10: Scatterplot with a regression line on the correlation between cyclomatic complexity (y-axis) and mutant density (x-axis)

In Figure 10 the scatterplot is shown for the correlation between cyclomatic complexity and mutant density. This visualisation shows that specifically for the Java dataset, there is almost linear increase between the two metrics. In essence, this result is not unexpected. Both mutant density and cyclomatic complexity contain the inclusion of some form of functionality when viewing at complexity. Either through for statements or possible operators present in the code. Therefore, an even increase in both of these metrics could be expected. Another potential reason for this even increase is the fact that most statements that increase cyclomatic complexity, include operators that can be mutated. Despite this, for the same amount of lines, it is more likely to expect more possible mutations than statements that increase cyclomatic complexity. With the limited data available it is hard to determine if this is specifically due to the

chosen Java projects or due to the programming language itself. To verify if it is caused by the selection of projects, a larger dataset should be formed. Verification if it is based on the programming language is done in a following part.

Despite the low variation at the start of the regression line, there is a significant increase in variation visible on the regression line. At higher cyclomatic complexity and mutant density this is expected. In functions with a larger source lines of code there is more potential variation in how the code was written and in what it contains. When only using these two metrics which have high possible variation at the tail end of the regression line, this effect is amplified. It indirectly also means that those cases with higher variation have a higher source lines of code so this metric is slightly incorporated without direct usage.

The same scatterplot for C++ can be found in Figure 11. In the case of C++, the mutant density increases at a higher rate than the cyclomatic complexity.

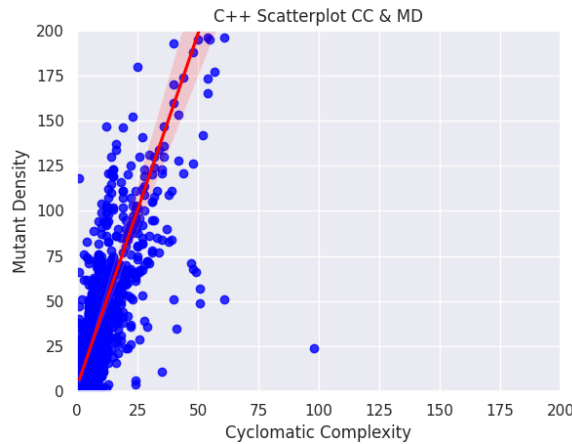


Figure 11: Scatterplot with a regression line on the correlation between cyclomatic complexity (y-axis) and mutant density (x-axis)

Languages such as C and C++ have a vastly different complexity in comparison to other languages such as Python, this also results differences in implementation time for the same functionality [31]. Languages with this higher complexity also contain more potential targets for mutations based on the built-in operators. An example of this is writing a for loop to iterate over a list. C++, Python and Java all contain simplified ways of creating this for loop. C does not have this luxury, the same for loop to iterate over list will contain an initialization, a condition and a update statement. Each of these parts can be mutated while a simplified version from Python has less possible mutations. There are other such difference between programming languages which could imply the presence of generally more targets for mutation.

The composition of the dataset is the biggest potential influence. If the selected projects contain more mathematical focused solutions, it's obvious this would create more influence for mutant density. Reducing the influence of the selection of projects would need a much larger dataset for each of the languages or a more thorough process for selecting the projects. The scatterplot shows some support for this statement due to the proportionally higher mutant density compared to cyclomatic complexity. Although the risk of the small dataset size remains, this indicates the general higher amount of mutations due to applications or programming language. Another potential cause could be the difference in tooling and computation method as mentioned earlier.

### 5.1.2 Step 1.2 - pearson correlation

The previous graphs are purely a visualisation and they need to be supplemented by additional numerical data to support it and avoid possible uncertainties due to chosen visualisation methods. This is done by utilizing the Pearson correlation. As mentioned in the methods, this is due to the continuation of the approach of [18].

Due to the differences visible between Java and C++, Table 4 covers all of the results from those languages. This table also includes correlation with source lines of code even though its relevance and visualisation is kept for the

following section. Even with their relevance for RQ2. instead of RQ1., they provide a bit more context to the exact value of the Pearson correlation coefficient.

Table 4: Results of pearson correlation for different metrics

	<b>Pearson MD CC</b>	<b>Pearson CC SLOC</b>	<b>Pearson MD SLOC</b>
<b>Java</b>	0.7427374837402358	0.8888096205091884	0.7606250350314062
<b>C++</b>	0.7142645801978144	0.9032900606076069	0.6993161787635052

For Java, the correlation between mutant density and cyclomatic complexity is positioned between the two other correlation coefficients. For C++, this correlation is in between the other metrics but still a lot closer to the lower correlations. This implies that the correlation is not stronger despite both of them having some incorporation of functionality. The more similar rate of increase does not say anything about the correlation based on these results. Although it might have been expected, when looking more closely at the scatterplots, it is clear that for Java the points are still a big clutter. The same spread of points for C++ does look a bit less cluttered which results in higher correlation but not a noticeable difference. A coefficient of 1 would imply perfect positive correlation while a coefficient of 0 implies no correlation. Despite the difference between the languages and between the different pairs, all of these correlations remain positive.

These results answer RQ1.. There is a positive correlation cyclomatic complexity and mutant density which is not high enough to cause redundancy and not low enough to discredit complementary usage. There are some issues surrounding consistency over different programming languages but despite the differences, the correlation does not change sufficiently to alter the conclusion. These results all provide a solid foundation to look further into the placement of mutant density as a complementary metric to cyclomatic complexity.

## 5.2 RQ2. Complementary

This section will focus mostly on the results surrounding analysis for RQ2.

### **Complementary - how can we utilize mutant density to provide additional insight into refactoring opportunities? Is there anecdotal evidence supporting this?**

This section will follow the following step-wise approach to gradually build upon previous findings.

- Step 2.1 - visualising contextual information to better grasp where outliers might be found
- Step 2.2 - outliers for average cyclomatic complexity and mutant density
- Step 2.3 - analysing the influence of using a dataset comprised of a single project or multiple projects
- Step 2.4 - anecdotal evidence for outliers of average cyclomatic complexity and mutant density
- Step 2.5 - outliers for scaled average cyclomatic complexity and mutant density anecdotal evidence
- Step 2.6 - investigating the complementary claim through set theory and altered metrics

It starts by providing contextual information which is in place to better grasp where outliers might be found. This allows for easier understanding of which patterns cause outliers within the context of this work. After the contextualisation, this is followed by an initial detection of outliers and refactoring opportunities using the z-score on the average cyclomatic complexity and average mutant density distributions. The metrics in question average by including division by source lines of code, therefore this section also includes a continuation on correlation analysis. Before giving the anecdotal evidence for these results, there is an analysis to determine the influence of using a dataset which is either comprised of a single project or a set of projects. Since thresholds are outside of this scope, being able to use only a single project would still provide a more applicable method. Now it is more appropriate to provide initial anecdotal evidence that also

shows a few shortcomings which are addressed in the section about outliers for a scaled version of the previous metrics. Finding complementary aspects is the most important part of RQ2., the previous steps all provide information which support the final complementary analysis. This is done through the usage of set theory and a new metric attempting to provide an alternative approach.

### 5.2.1 visualising contextual information to better grasp where outliers might be found

Visualising contextual information is done through plotting the distribution of differences between two metrics for the complete Java dataset. This gives additional information about the relationship two specific metrics hold to each other without directly looking at the correlation. An example of this can be found at Figure 12. Note that this is purely a distribution of the cyclomatic complexity subtracted from the source lines of code for each single method. This also implies that there is no division by the source lines of code for this analysis. The original expectation, based on previous information surrounding the correlation and the functionality of the metrics, is that cyclomatic complexity will always be higher than source lines of code. This is visible in Figure 12 where the distribution is heavily shifted towards the right (more source lines of code than cyclomatic complexity). This implies that refactoring opportunities between these two metrics could occur on for two different reasons. Either the cyclomatic complexity is higher than the source lines of code, or the cyclomatic complexity is proportionally much larger than the source lines of code.

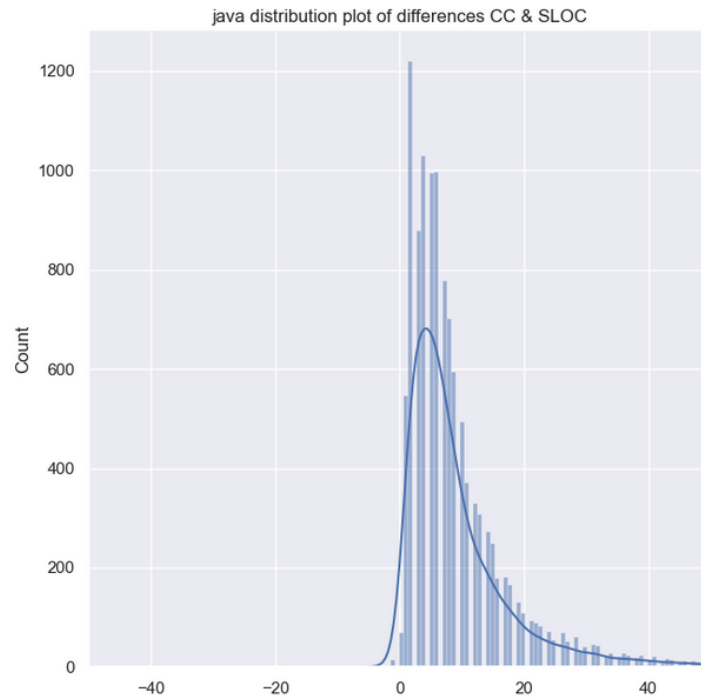


Figure 12: Distribution plot of cyclomatic complexity subtracted from source lines of code where the x-axis is the difference and the y-axis the count

If the same graph is used for mutant density subtracted from source lines of code, this results in Figure 13. From an intuitive perspective, in contrast to cyclomatic complexity, having more mutations than source lines of code is a possibility. Since a single line of code can have multiple different mutations depending on the chosen operators, these cases are assumed to occur more than a higher cyclomatic complexity compared to the lines of code.

Looking at the graph however, this is still a very infrequent presence. It occurs, but not as frequently as it might be possible. This is not that much of an issue since it might indicate the use good practices throughout the projects. The infrequent presence of these cases is not unexpected after the visualisations of the correlation. In the Java dataset, the mutant density and cyclomatic complexity were closely aligned. This means that despite it being possible of having methods with a much higher mutant density than source lines of code, the selected projects for this dataset do not contain it in large quantities. This would indicate that refactoring opportunities for mutant density in relation to source lines of code follow the same structure as with cyclomatic complexity.

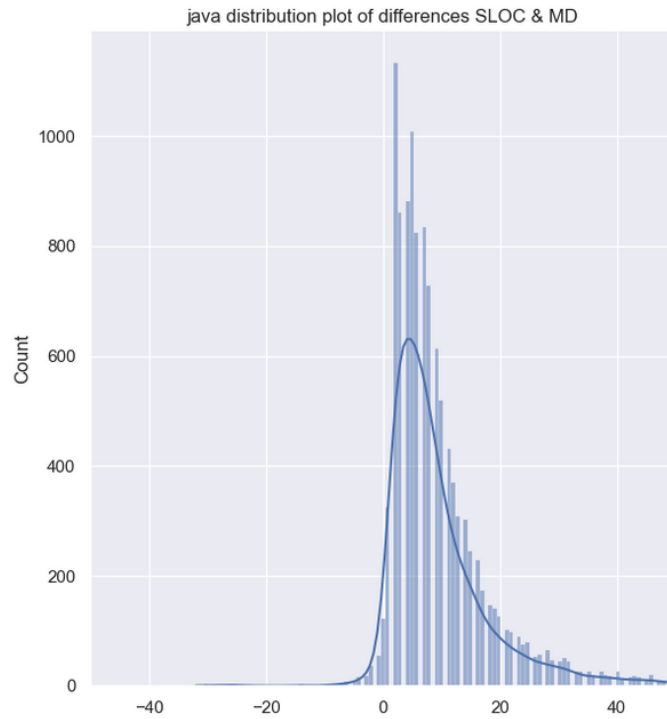


Figure 13: Distribution plot of mutant density subtracted from source lines of code where the x-axis is the difference and the y-axis the count

Finally, it is also possible to visualize the difference between mutant density and cyclomatic complexity. This can be seen in Figure 14. The graphs are relatively balanced with most of its methods having no, or a very small difference between cyclomatic complexity and mutant density. Possible refactoring opportunities in these cases would be that either of the metrics is proportionally small or large compared to the other. The same results over the datasets of different languages provide some differences yet again. The differences between cyclomatic complexity and source lines of code does not change over the different languages but any comparison with mutant density does. The change in results however falls in line with previously observed patterns.

The same differences between mutant density and source lines of codes for the C++ dataset have an almost even balance between the differences. This could be caused by the three causes that were discussed earlier, the differences between the tooling, general differences between implementations for languages and variation within the selection of the projects. It is hard to differentiate between those issues and pinpoint the cause within the scope of this work.

This large variation in mutant density for C++ is also present for the difference between the cyclomatic complexity and mutant density. It causes an extreme shift where most cases have a higher mutant density when compared to its corresponding cyclomatic complexity. This also changes the perspective on what an outlier could possibly be. Having different distributions over the languages is not optimal. It makes it very hard to generalize results and be certain about what the exact influences are on these metrics. Although it makes generalization very hard, it does not make the metrics and upcoming outlier detection bad. In the examples shown and the data resulting these, it all functions well within expectations. The difference in languages and the behaviour simply prompts careful consideration when looking at the possible generalization of singular results.

### 5.2.2 outliers for average cyclomatic complexity and mutant density

Using the simple difference between the two metrics is good for baseline indications but not for general usage. Therefore, the next step is looking at the average mutant density and average cyclomatic complexity. The results

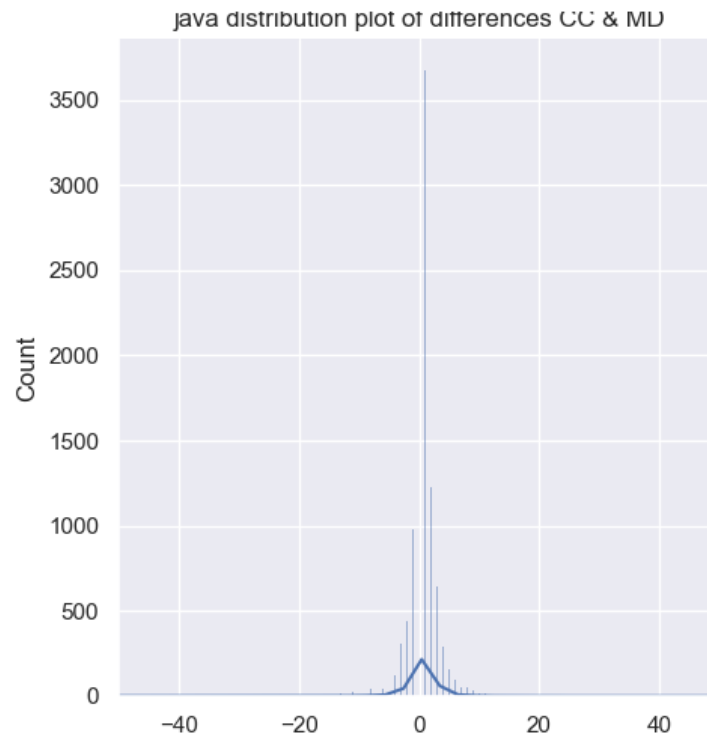


Figure 14: Distribution plot of mutant density subtracted from cyclomatic complexity where the x-axis is the difference and the y-axis the count

on more advanced metrics on the relation between cyclomatic complexity and mutant density will be discussed a bit later.

The average mutant density and cyclomatic complexity are averaged through division by the source lines of code. This prompts a further inspection in the correlation analysis by now including source lines of code. If there is too much correlation between those metrics, the redundancy would be visible in a different way. This would result in a division by almost the exact same metric, removing the effect that source lines of code should have.

The results for correlation between cyclomatic complexity and source lines of code will be taken from [18]. They provide extensive discussion on this subject with similar methods used. A short summary of the results shows a high positive correlation which is deemed not strong enough to cause redundancy between the methods. This implies that for average cyclomatic complexity the division does not influence the functionality of the metric.

Visualizing the relation between mutant density and source lines of code with a boxplot and scatterplot can be seen in Figure 15 and Figure 16. These graphs show a high correlation for lower source lines of code and more variation towards the tail of both the boxplot and scatterplot. At its core, this follows the principle where larger functions have more possible variation. This is amplified due to the nature of mutant density. The amount of cyclomatic complexity you can cause in a single line is not completely limited. It is always possible to increase it by adding more conditional operators. In reality, there is more of a limitation on how much cyclomatic complexity you can increase for every line of code. For example, there are also the lines of code within the bodies of for loops and if statements. Unless they contain nested statements, it does not contain statements increasing cyclomatic complexity any further. Most statements that increase cyclomatic complexity, also increase mutant density. This means that with a lower source lines of codes, a higher mutant density is more probable result.

The same results also vary with C++ like previous sections which can be seen in Figure 17. This shows a clear increase in variation and also a very different rate in which both proportionally increase. Just like previous examples however, this increase in variation does not take away from the positive correlation which is present. The overall results are taken back from Table 4 where every correlation remains positive despite potential changes over languages and between metrics pairs. Therefore it is concluded that the effect of these differences is also not enough to

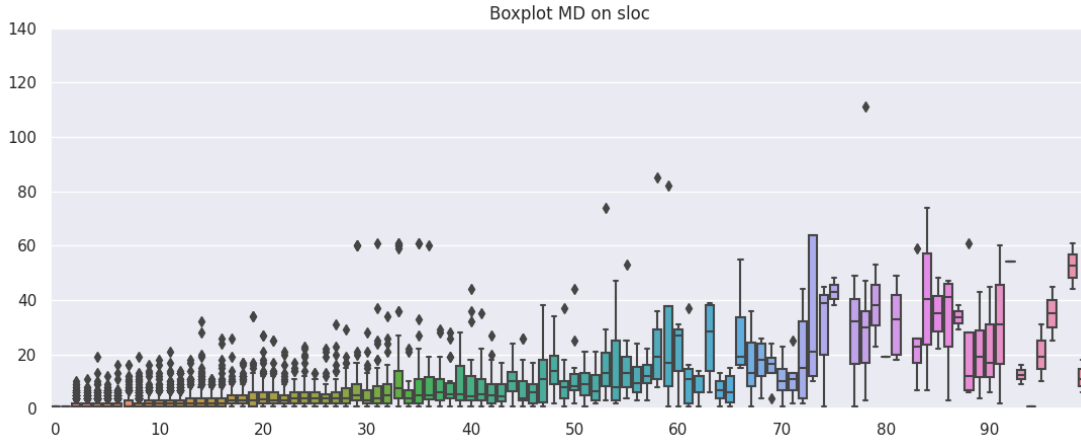


Figure 15: Boxplot on the correlation between mutant density (y-axis) and source lines of code (x-axis).

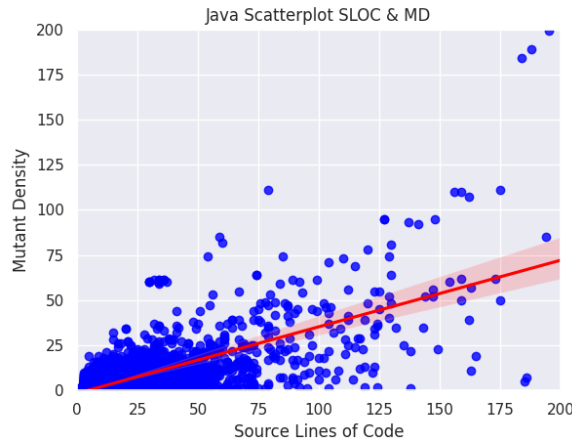


Figure 16: Scatterplot with a regression line on the correlation between mutant complexity (y-axis) and source lines of code (x-axis)

change the functionality of the average metrics.

Without any redundancy present, it is now possible to visualize outliers for the metrics. Using the results of average mutant density and average cyclomatic complexity in a distribution which shows the distances to mean, can be seen in Figure 18 and Figure 19. This provides initial insight in what potential cutoff values could be for refactoring targets. Based on the current data, there are slight differences between the two metrics but a current cutoff value will be placed at 4. All of this data is still generated over datasets comprised of multiple different projects even though it was mentioned in the methods that there might be certain variation caused by this. Up until now, this is not something blocking since the previous information functioned as a foundation. Still, before diving into further metrics and code example, there needs to be some verification surrounding this.

### 5.2.3 analysing the influence of using a dataset comprised of a single project or multiple projects

The largest potential risk surrounding the usage of a single project or a set of projects is the potential generalisation that occurs when making a larger dataset. The most simplistic way of verifying if generalisation takes place over a complete dataset is using basic outlier detection and verifying the differences between the two datasets.

These examples are based on the Java dataset, with Spring as the project that will be singled out. Generating outliers on average mutant density for the complete dataset gives 45 outliers that belong to the Spring project. Everything that is four standard deviations from the mean are considered outliers in this case. There are more outliers



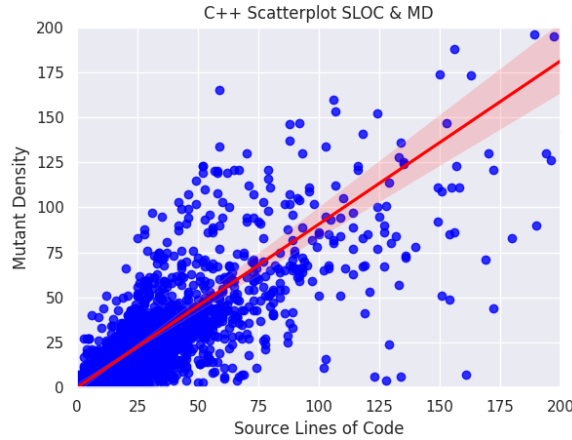


Figure 17: Scatterplot with a regression line on the correlation between mutant density (y-axis) and source lines of code (x-axis)

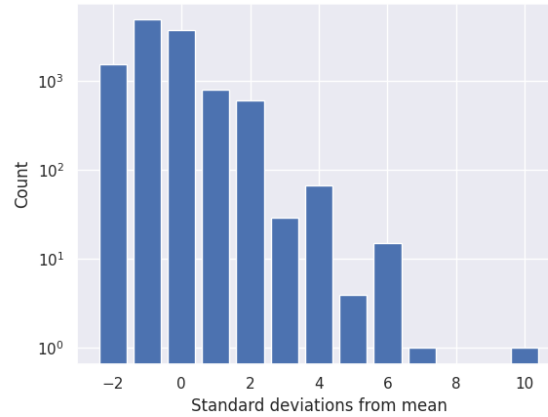


Figure 18: Distribution of outliers based on the average cyclomatic complexity

when doing this on the complete dataset but only 45 that belong to Spring. If the outliers are calculated based on Spring alone, this number increased to 76.

An important detail is that the increase of outliers does not cause a decrease of information. Every single one of those 45 outliers also occur in the 76 of only using the Spring project. This pattern is reoccurring, implying that using only a single project has no downsides. Staying in line with the original assumptions on this analysis, it is even safer to say that using more projects generalises and causes potential loss of information.

Looking more in depth on the cases that were added by only looking at Spring gives more insight on the situation. All the outliers that are added are relatively close to the edge of what was considered an outlier for this analysis. Every outlier that is close to five or even six times the standard deviation is caught in both scenarios. This further strengthens the idea that using more projects to use the mean and standard deviation from generalises. The cases visualized like the previous outliers can be seen in Figure 20. The orange cases are all the newly added outliers. As mentioned earlier, these are all very close to the edge of what can be considered an outliers so the usefulness is still a bit up to debate.

Since thresholding remains a difficult issue to address, it is also unsure whether these additional outliers contain the most useful information. Despite this, due to there not being any decrease in information, and using only a singular project reducing computational time, this is the approach from here on out.

One downside to this approach is that outliers are also more easily created due to the lack of more general

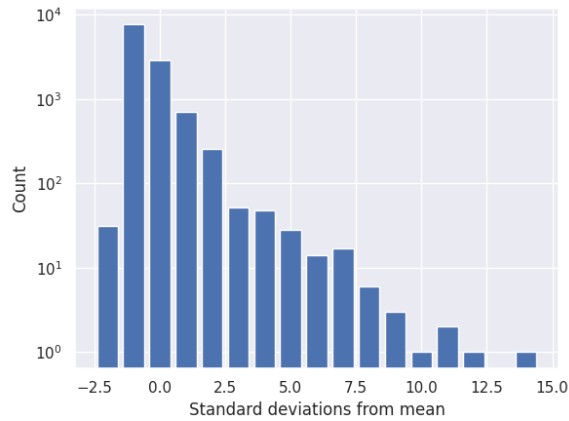


Figure 19: Distribution of outliers based on the average mutant density

data. If a project only has a few functions with a source lines of code above 50, these are much more prone to be outliers. Using a division by source lines of code and keeping the sizing of functions in mind reduces the impact of this. Still, even by using methods to reduce the influence of the sizing of methods, functions with higher source lines of code are more prone to be outliers. This is not a problematic occurrence. Functions that are too large in source lines of code compared to other functions within the same project might not be outliers in terms of cyclomatic complexity or mutant density, but the source lines of code still remains a complexity metric. Therefore, outliers based on source lines of code are more easily caught without specifically using this metric.

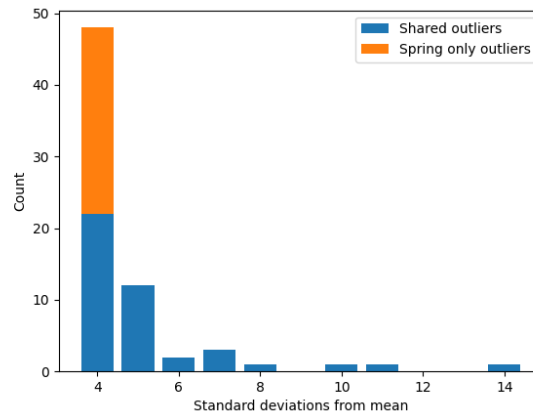


Figure 20: Distribution of outliers based on the difference between mutant density which shows the unique cases over a using a single project or a dataset of projects

#### 5.2.4 anecdotal evidence for outliers of average cyclomatic complexity and mutant density

With the previous analysis finished, it is now more appropriate to start diving into code examples on the previously discussed average mutant density and average cyclomatic complexity. The following results are all based on the Spring project, the reasoning behind this choice is that it's a large project with a large amount of variation in the domains it covers. The first example is the function used in the background information. This *isSockJsSpecialCharFunction* 4 on p. 35 is a severe outlier for both average mutant density and average cyclomatic complexity. The z-score for both of these metrics is above 10. This outliers stands out the most, especially compared to other outliers for average cyclomatic complexity. The second highest outlier for this metric has a z-score of 7.2, this is quite a large gap with the z-score of 10 but with the examples this gap is quite obvious.

The issue with the *isSockJsSpecialChar* is immediately visible. There are too many possible operations, too many conditional operators causing a high complexity on multiple aspects. In earlier sections it was mentioned that small functions like these, are not necessarily a threat to maintainability. It is written too complex and could certainly be refactored and simplified, but it still remains a small function separated from other functionality. This encapsulation of problematic functionality might simply be caused by the circumstances of usage, for example this calculation could be called at multiple locations. It could also be a conscious decision to create this encapsulation but despite the cause, it remains a reasonable refactoring opportunity.

---

```
private boolean isSockJsSpecialChar(char ch) {
    return (ch <= '\u001F') ||
        (ch >= '\u200C' && ch <= '\u200F') ||
        (ch >= '\u2028' && ch <= '\u202F') ||
        (ch >= '\u2060' && ch <= '\u206F') ||
        (ch >= '\uFFFO') || (ch >= '\uD800' && ch <= '\uDFFF');
}
```

---

Listing 4: Highest outlier for average mutant density

The *matches* Function 5 on p. 35 is the second highest outlier for average mutant density. This is for very similar reasons as the previous function, a long statement on a single lined function. There are less operators increasing cyclomatic complexity so it is not an outlier for that metric. Again, this is a very small function and most other cases with a high z-score based on average mutant density and cyclomatic complexity are less than 15 lines of code.

---

```
public boolean matches(Method m, @Nullable Class<?> targetClass) {
    return m.getName().equals("overload") &&
        m.getParameterCount() == 1 &&
        m.getParameterTypes()[0].equals(int.class);
}
```

---

Listing 5: Second highest outlier for average mutant density

Another result that is also visible through these metrics is that there is no filtering for test functions. This results in a few test functions appearing as outliers. Test functions are also slightly different in structure compared to normal methods, they often do not follow the same clean code aspects. This also means that the test functions are used for the calculation of the mean and the standard deviation. This is a relatively small issue but a detail to be kept in mind for further improvements and fine-tuning. These functions might influence the mean and standard deviation calculation but in a very limited sense. Especially since testing functions are in most projects much less present compared to normal functions.

## 5.2.5 outliers for scaled average cyclomatic complexity and mutant density

With the scaled metric, it is expected that functions which contain problematic patterns combined with larger source lines of codes are shown. This means that the worst rated methods might not be seen as the worst code, but they contain patterns that combined with size cause a problematic result. This does however reduce the amount of outliers that are found with a z-score higher than 4. For the scaled average cyclomatic complexity this gives only 2 outliers and on scaled average mutant density this decrease is less drastic. This change in distribution is visible in Figure 21 where the distribution is much more limited than the previous Figure 18. A first example of how this influences the obtained functions can be partially seen in Function 6 on p. 36 and fully in Function 16 on p. 51. For the Spring project, this is the worst rated function after using a scaled version of the average mutant density. At first sight it does not look like bad method, especially compared to the previous shorter examples. However, with more detailed inspection, it is visible that it contains more potential mutations. In terms of cyclomatic complexity, it is an outlier if the cutoff would be lowered to 2. This still means that the mutant density is much more present than the cyclomatic complexity.

The partial versions already shows this through a presence of multiple nested statements with even more potential mutations, the full function only expands on this. What mostly influences the scoring right now is the additional scaling that has been done. Due to the size of the function, and the higher than average cyclomatic complexity and mutant density, it becomes a prime target for refactoring. The smaller functions still have a relatively large z-score since they were such large outliers to begin with but this remains acceptable. The small functions are not expected to be completely dropped through the addition of scaling. A better balance in the given outliers is the main goal which is demonstrated by the obtained outliers.

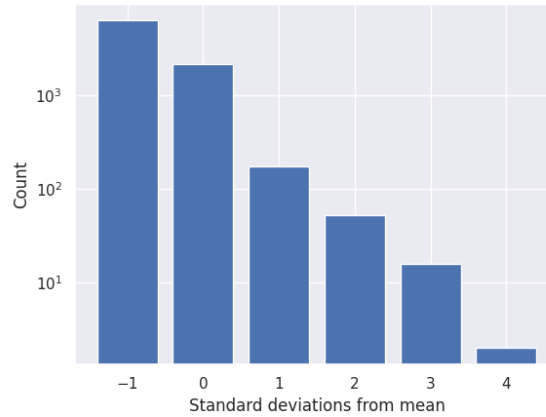


Figure 21: Distribution of outliers based on the scaled average cyclomatic complexity

---

```

final boolean merge( final SymbolTable symbolTable, final Frame dstFrame, final int
    catchTypeIndex) {
    ...
    for (int i = 0; i < numLocal; ++i) {
        int concreteOutputType;
        if (outputLocals != null && i < outputLocals.length) {
            int abstractOutputType = outputLocals[i];
            if (abstractOutputType == 0) {
                concreteOutputType = inputLocals[i];
            } else {
                concreteOutputType = getConcreteOutputType(abstractOutputType, numStack);
            }
        } else {
            concreteOutputType = inputLocals[i];
        }
        if (initializations != null) {
            concreteOutputType = getInitializedType(symbolTable, concreteOutputType);
        }
        frameChanged |= merge(symbolTable, concreteOutputType, dstFrame.inputLocals, i);
    }
    ...

```

---

Listing 6: Partial code example of worst function using scaled average mutant density

Since there are only two outliers for cyclomatic complexity after the scaling with a score higher than 4, choosing one is not hard. The first function is the *getOpcode* function which is partially shown at Function 7 on p. 36 and fully at Function 17 on p. 52. It is a large switch case statement that could use some simplification. Large switch case statements like these often exist for some sort of parser. This also means that it is not always problematic or avoidable. These types of outliers are similar to the very small functions in terms of how they should be interpreted. Despite being outliers, not every outlier is directly a bad function due to the functionality it implements. Enforcing refactoring on these might lead to solutions that could negatively influence maintainability and readability. Examples like this make it very clear that despite the usage of these metrics, it is always important to keep the context of implementation in mind. Certain types of complexity are simply unavoidable or have bigger upsides than downsides.

---

```

public int getOpcode(final int opcode) {
    if (opcode == Opcodes.IALOAD || opcode == Opcodes.IASTORE) {
        switch (sort) {
            case BOOLEAN:
            case BYTE:
                return opcode + (Opcodes.BALOAD - Opcodes.IALOAD);
            case CHAR:

```

---

```

        return opcode + (Opcodes.CALOAD - Opcodes.IALOAD);
    case SHORT:
        return opcode + (Opcodes.SALOAD - Opcodes.IALOAD);
    case INT:
        return opcode;
    case FLOAT:
        return opcode + (Opcodes.FALOAD - Opcodes.IALOAD);
    case LONG:
        return opcode + (Opcodes.LALOAD - Opcodes.IALOAD);
    case DOUBLE:
        return opcode + (Opcodes.DALOAD - Opcodes.IALOAD);
    case ARRAY:
    case OBJECT:
    case INTERNAL:
        return opcode + (Opcodes.AALOAD - Opcodes.IALOAD);
    ...
}

```

---

Listing 7: Partial code example for biggest outlier based on scaled average cyclomatic complexity

If the style of the outliers are closer to the example of mutant density, looking more like a standard implementation but higher complexity due to size inclusion, it would be more interesting. Not every project will necessarily have functions like these. Still, this does not mean this is a bad example, the function still contains an if else statement which could be separated in two different functions. Even though this would most likely create two outliers instead of one, the deviation from the mean would not be as large and it might increase readability.

The second outlier is closer to the mutant density example, this function is given in the appendix Function 18 on p. 53. It is an very large function that does not contain switch statements, the complexity purely comes from the size combined with a high amount of if else statements. This creates another very good example about the importance of the addition of the scaling.

### 5.2.6 investigating the complementary claim through set theory and altered metrics

The first steps of looking at the relation between outliers for mutant density and cyclomatic complexity is using set theory as explained earlier. As for the metrics, the previous scaled metrics are going to be used. One small variation is that instead of only taking from four standard deviations as outliers, this will be decreased to two. Four leaves too little outliers to apply set theory and after applying the scaling, two is still a reasonable cutoff. Let's go back to the visualization of how the outliers are distributed after applying the scaling. In Figure 21 it shows that two does not have too much cases but also not too little. Compared to the same graphs when there is no scaling applied, a cutoff of two would be too low and four is a better option. Due to the shifting of the general distribution, two is a reasonable option now.

Using this cutoff value provides 168 outliers based on mutant density and 70 based on cyclomatic complexity. This difference in amount of outliers is not new information. The larger possible variation on mutant density with higher source lines of code is the main cause for this. These two sets have an intersection of 46 cases, this is a large part of the cyclomatic complexity set. This means that a good portion is covered through mutant density but that the unique information still remains. Looking at the unique cases based on mutant density there are two general types of functions that are obtained. The first type is shown partially by Function 8 on p. 37 and fully by Function 19 on p. 54, these functions are relatively average looking and still contain cyclomatic complexity. The cyclomatic complexity is however not enough to raise it as an outlier for that metric. The second type of functions are similar to Function 9 on p. 38. These are short functions that often include a very small cyclomatic complexity or completely none. The same idea about short functions as mentioned earlier applies.

---

```

public Resource findLocalizedResource(String name, String extension, @Nullable Locale locale) {
    ...
    if (locale != null) {
        String lang = locale.getLanguage();
        String country = locale.getCountry();
        String variant = locale.getVariant();

        if (variant.length() > 0) {
            String location =

```

```

        name + this.separator + lang + this.separator + country + this.separator +
            variant + extension;
        resource = this.resourceLoader.getResource(location);
    }
    if ((resource == null || !resource.exists()) && country.length() > 0) {
        String location = name + this.separator + lang + this.separator + country + extension;
        resource = this.resourceLoader.getResource(location);
    }
    ...
}
...
}

```

---

Listing 8: First partial code example of unique information from average mutant density

---

```

public boolean isOverridable() {
    return !isStatic() && !isFinal() && !isPrivate();
}

```

---

Listing 9: Second code example of unique information from average mutant density

---

Then there are the type of functions that are only outliers for cyclomatic complexity. The following function might seem very similar to the previous large switch case statement but there is a fundamental difference. This *invoke* Function 10 on p. 38 (complete Function 20 on p. 55) has a lot less potential mutation operators which causes it to only be an outlier for cyclomatic complexity.

```

public Object invoke(Object proxy, Method method, Object[] args) throws Throwable {
    switch (method.getName()) {
        case "equals":
            return (proxy == args[0]);
        case "hashCode":
            return System.identityHashCode(proxy);
        case "getTargetConnection":
            return getTargetConnection(method);
        case "unwrap":
            if (((Class<?>) args[0]).isInstance(proxy)) {
                return proxy;
            }
            break;
        case "isWrapperFor":
            if (((Class<?>) args[0]).isInstance(proxy)) {
                return true;
            }
            break;
    }
    ...
}

```

---

Listing 10: Partial code example of outlier unique for average cyclomatic complexity

---

Now the second method to find the relation between these metrics is by using the scaled average difference between cyclomatic complexity and mutant density. The distribution of outliers can be visualized as Figure 22. With the cutoff still being at -2 or 2, this provides 156 outliers based on this metric. The important question for the results of this metric is how many outliers are found that are not found by the scaled average mutant density and cyclomatic complexity. There are 97 of those 156 outliers that are also found by these two other metrics. If we remove all similarities between the three metrics, only 59 completely new outliers are found. All of these outliers have a relatively low z-score, it still falls outside the cutoff value but they mostly fall between 2 and 4. This means that more extreme cases are already caught by the usage of only a single metric. Does this mean the metric lacks in performance? Not really, it still gives a more precise reasoning on why something is an outlier. The previous metrics had no notion of the direct relation between cyclomatic complexity or mutant density, now it can be shown whether an outlier is caused by a proportionally higher mutant density or cyclomatic complexity.

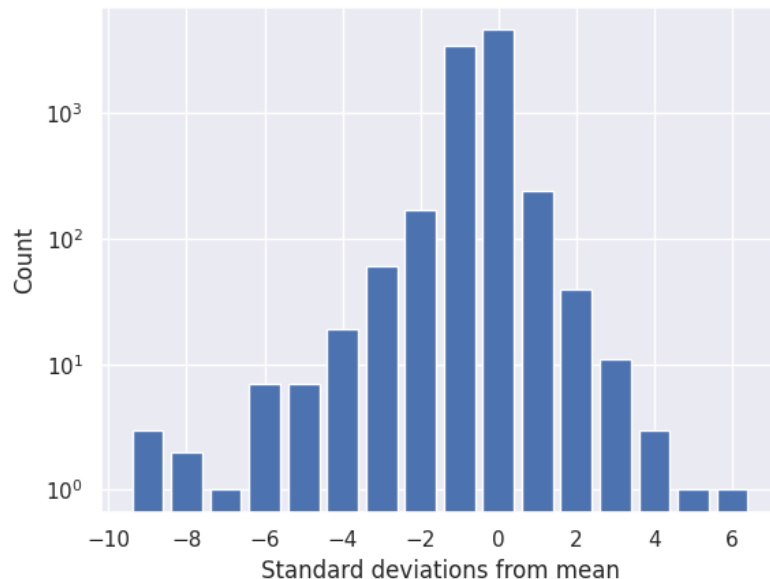


Figure 22: Distribution of outliers based on the scaled average CC & MD

The functions that are unique with a slightly lower z-score are still interesting. A first example is this *areBoxingCompatible* Function 11 on p. 39 (complete Function 21 on p. 56) which is an outlier due to a much higher cyclomatic complexity than mutant density. This is mainly due to a single reason. Often the conditional part of an if statement contains the possibility for quite some mutations. Most conditional statements of this function consist of a single true or false statement. On top of that, the bodies of those if else statements contain very few lines of code which also reduce the possible mutations. Due to these reasons, this is an outlier and also an interesting function for this case. This style of functions is very close to previous examples containing large switch case statements, therefore the same argumentation remains that not all of these cases will correspond to refactoring opportunities. This means that a proportionally higher cyclomatic complexity is not always a direct indication of the need for refactoring.

---

```

public static boolean areBoxingCompatible(String desc1, String desc2) {
    ...
    if (desc1.length() == 1) {
        if (desc1.equals("Z")) {
            return desc2.equals("Ljava/lang/Boolean");
        }
        else if (desc1.equals("D")) {
            return desc2.equals("Ljava/lang/Double");
        }
        else if (desc1.equals("F")) {
            return desc2.equals("Ljava/lang/Float");
        }
        else if (desc1.equals("I")) {
            return desc2.equals("Ljava/lang/Integer");
        }
        else if (desc1.equals("J")) {
            return desc2.equals("Ljava/lang/Long");
        }
    }
    ...
}

```

---

Listing 11: Partial code example of outlier based on a higher than average cyclomatic complexity compared to mutant density

---

```
private static int nextPowerOf2(int val) {
    val--;
    val = (val >> 1) | val;
    val = (val >> 2) | val;
    val = (val >> 4) | val;
    val = (val >> 8) | val;
    val = (val >> 16) | val;
    val++;
    return val;
}
```

---

Listing 12: Outlier based on a higher than average mutant density compared to cyclomatic complexity

A second example is this *nextPowerOf2* Function 12 on p. 40 function which was an outlier due to the higher mutant density compared to cyclomatic complexity. This function is quite a bit shorter but it is still clear why this is an outlier. There is no code that increases the cyclomatic complexity at all. Lacking cyclomatic complexity is not a bad indication, some functions simply need to execute a certain amount of operations. Unlike the previous function, it is easier to argue that a higher mutant density compared to cyclomatic complexity remains a good refactoring target. The example function has clear duplication which could be simplified. If functions like this occur with large source lines of code, it would be even better to provide these as refactoring targets. This metric does have a downside causing it to not be a better version of using set theory and the separate metrics. Since it relies on the difference between cyclomatic complexity and mutant density, if this distance is not present, there won't be an outlier. Even if both of these metrics are high, as long as the difference is smaller than a method will not be detected. Therefore the previous metrics retain their value as a metric since this metric emphasises other aspects. Another flaw was shown by the *areBoxingCompatible* Function 21 on p. 56. It is an outlier and for a clear reason but this does not make it a good target. Filtering these cases is also not that easy, at a certain point, a higher cyclomatic complexity might mean a good refactoring target even with a low mutant density. Deciding when these could be seen as refactoring opportunities is a non trivial issue.

How do these results answer **RQ2. Complementary - how can we utilize mutant density to provide additional insight into refactoring opportunities? Is there anecdotal evidence supporting this?**. The previous results have shown different aspects of the usage of mutant density and how it could possibly be applied. Although combining mutant density and cyclomatic complexity into a singular metric provides new interesting information, it leaves out other aspects and is hard to filter. This does not mean that it is impossible to do, just within the scope of this work this is not further expanded upon. The most interesting results are obtained by utilizing the scaled average mutant density and cyclomatic complexity. They provide a good insight into refactoring opportunities taking factors such as function size into account. The set theory analysis also proved that outside of the correlation between the metrics, there is also a solid similarity between the two. This strengthens the idea of using mutant density in a complementary manner. The anecdotal evidence is shown by the code examples, these examples also provide the same insight when revisiting them from the perspective of the DevOps tooling. As an example the previous *nextPowerOf2* Function 12 on p. 40 will be used, it has an original mutant density of 12 and a cyclomatic complexity of 1. CodeScene does not recommend any changes to this function based on its metrics or on based on cyclomatic complexity. Refactoring this as Function 13 on p. 41 heavily reduces the potential mutations by moving the reoccurring operation to a separate function. This changes nothing in terms of cyclomatic complexity but it does improve other aspects which are not covered.

This example is very small in size, it might not even be worth looking at functions of this size but the principles remain the same. Even if it would be contained within a larger function, functionality such as *nextPowerOf2* Function 12 on p. 40 is not viewed by cyclomatic complexity. CodeScene also frequently rely on other factors such as large conditionals and nested blocks to determine refactoring opportunities. Both of these rely on the presence of statements that also increase the cyclomatic complexity. This implies that also viewing complexity through the present operator is a viable way to analyze refactoring opportunities.



---

```

private static int nextPowerOf2_refactored(int val) {
    val--;
    val = val_operation(val, 1);
    val = val_operation(val, 2);
    val = val_operation(val, 4);
    val = val_operation(val, 8);
    val = val_operation(val, 16);
    val++;
    return val;
}

private static int val_operation(int val, int shift) {
    return (val >> shift) | val;
}

```

---

Listing 13: Refactoring of nextPowerOf2

### 5.3 RQ3. Fine Tuning

#### RQ3. Fine Tuning - what is the impact of fine-tuning mutant density on correlation and complementary?

The last results that need to be covered are those for operator specific operators which correspond to RQ3.. Due to the easier tooling in C and C++ using dextool, there will be no coverage across languages for this section. This might cause variation but the general takeaway from the results will remain the same. More specifically, OpenCV was chosen since it is the largest dataset amongst those chosen for C and C++. It's functionality is also focused around real time computer vision, therefore containing mathematical operations providing enough different operators.

#### 5.3.1 impact of mutation operators

Looking at the amount of outliers for each single mutation operator in Table 5, there are a few clear differences. Certain specific mutation operators have a much larger presence than others, this can be seen in the amount of total mutations and in the amount of methods that are influenced. Out of this data UOI (Unary operator insertion) has the least presence while DCR (Decision/Condition Requirement) and SDL (Statement Deletion) have the same presence. Despite this difference in presence of mutation operators, Dextool notes that a combination of ABS, UOI, LCR, AOR and ROR reaches a coverage of 99,5%. This means that despite the low presence of UOI, it still covers possible mutations that other metrics don't. Before looking at unique outliers, there are 24 overarching outliers. The following *ReadTagFallback* Function 14 on p. 41 is an example of those cases.

---

```

uint32_t CodedInputStream::ReadTagFallback(uint32_t first_byte_or_zero) {
    const int buf_size = BufferSize();
    if (buf_size >= kMaxVarintBytes ||
        (buf_size > 0 && !(buffer_end_[-1] & 0x80))) {
        GOOGLE_DCHECK_EQ(first_byte_or_zero, buffer_[0]);
        if (first_byte_or_zero == 0) {
            ++buffer_;
            return 0;
        }
        uint32_t tag;
        ::std::pair<bool, const uint8_t*> p =
            ReadVarint32FromArray(first_byte_or_zero, buffer_, &tag);
        if (!p.first) {
            return 0;
        }
        buffer_ = p.second;
        return tag;
    } else {
        if ((buf_size == 0) &&
            ((buffer_size_after_limit_ > 0) ||
             (total_bytes_read_ == current_limit_)) &&
            total_bytes_read_ - buffer_size_after_limit_ < total_bytes_limit_) {

```

---

```

        legitimate_message_end_ = true;
        return 0;
    }
    return ReadTagSlow();
}
}
}

```

Listing 14: Outlier over every single mutation operator

It is immediately clear that this function contains a high amount of possible mutations due to the presence of many different operators. There is also a good amount of variation visible in the types of operators, thus it being an outlier over every mutation operator. On top of that, these outliers are based on average mutant density, there is no additional scaling for the sizing of the functions. Despite there not being any scaling for this, all of the functions in this category have a size large enough to consider them good refactoring targets. This is caused by the structure of this analysis. Being an outlier over a multitude of different mutation operators requires a fair amount of operators in the code that conform to each of the types. This is less feasible with smaller sized functions, directly causing the functions the increase in size without specifically including this in the metric.

Looking at the outliers that are unique for a single mutation operators is a very different way of working then the previous analysis. In Table 6 it is shown how many unique outliers there are for each mutation operator. Most functions have at least some form of overlap in terms of mutation operators. Going from multiple thousand mutations, only a very few remain when looking at pure unique cases. Most operators do not even have unique cases remaining, this might not be optimal since some operators also have overlap but it is how it will be interpreted for now. What is interesting is how most of the unique cases are for operators that do not have a very large coverage. The mutation operators with less coverage are more likely to be more different to other thus causing more potential unique cases.

Table 5: Mutation Statistics

Mutations	Methods	Total Mutants
rorp	958	5920
aor	628	7906
dcr	2946	20776
cr	1449	5877
lcr	692	6004
aors	641	2051
sdl	2946	20776
ror	958	5920
lcrb	441	4198
uoi	422	829

There are not that many code example too choose from, the examples can also be generally divided in two groups. One of the examples is the following *opj\_jp2\_start\_copmress* Function 15 on p. 42, obtained as an outliers of UOI.

```

OPJ_B00L opj_jp2_start_compress(opj_jp2_t *jp2,
                                opj_stream_private_t *stream,
                                opj_image_t * p_image,
                                opj_event_mgr_t * p_manager
                                )
{
    assert(jp2 != 00);
    assert(stream != 00);
    assert(p_manager != 00);
    if (! opj_jp2_setup_encoding_validation(jp2, p_manager)) {

```

```

    return OPJ_FALSE;
}
if (! opj_jp2_exec(jp2, jp2->m_validation_list, stream, p_manager)) {
    return OPJ_FALSE;
}
if (! opj_jp2_setup_header_writing(jp2, p_manager)) {
    return OPJ_FALSE;
}
if (! opj_jp2_exec(jp2, jp2->m_procedure_list, stream, p_manager)) {
    return OPJ_FALSE;
}
return opj_j2k_start_compress(jp2->j2k, stream, p_image, p_manager);
}

```

---

Listing 15: Outlier for only a single mutation operator (Uoi)

This function is not as small as previous examples and contains a clear limit on the types of operators it contains. Therefore, functions such as these could be placed in the first category, medium sized functions with a simple limit on the operators. Note that there is only a small amount of operators in this function. UOI has a much smaller reach and potential targets in comparison to other operators as shown earlier. Since the threshold for calculating the outliers are now based on singular metrics, this means that the mean and standard deviation vary a lot between the different operators. This information is given in Table 7, where the difference is clearly visible. A mutation operator with more reach does not directly correlate to higher means or standard deviations. DCR has the highest amount of mutations but a relatively low mean and standard deviation. In the case of UOI the mean is very low, outliers are thus much more easily created.

Table 6: Single Mutation Outliers

Mutations	Outliers
rorp	0
aor	0
dcr	0
cr	30
lcr	8
aors	3
sdl	0
ror	0
lcrb	0
uoi	24

The second type of functions obtained through this analysis are small functions with a few operators of the corresponding type. When looking at methods that are outliers for each method, this mostly gave larger functions as explained earlier. It is not unexpected that this analysis provides opposite results due to its opposite approach.

Mutations	Mean	Standard Deviation	Total Mutants
rorp	0.433	0.492	5920
aor	0.803	0.833	7906
dcr	0.544	0.410	20776
cr	0.329	1.461	5877
lcr	0.432	0.392	6004
aors	0.204	0.211	2051
sdl	0.544	0.410	20776
ror	0.433	0.491	5920
lcrb	0.380	0.291	4198
uoi	0.087	0.179	829

Table 7: Table of mutations with mean and standard deviation

How does this answer **RQ3. Fine Tuning - what is the impact of fine-tuning mutant density on correlation and complementary?** The impact of fine-tuning within this work is limited due to the taken approach. Instead of analyzing specific combinations of mutation operators, this work focused on two more defined scenarios. The most interesting part of these analyses is using outliers that occur for each individual mutation operator. It does not directly benefit from the modularity of the metric in a way that was originally expected but it does provide some direction for future attempts at this. Instead of calculating outliers that occur over each mutation operator, it might be possible to create a metric that is used the amount of mutation operators contain an outlier to give a more defined way of finding refactoring opportunities.

## 6 Threats to Validity

### 6.1 Construct validity

The construct validity discusses whether the operational measures reflect the goal and intentions of this work. In this work, there are a few different aspects which deviated from the original expectations and intentions. Most of these aspects occurred in what was expected to be the preparational phase. More specifically, this concerns the creation of the dataset and getting functional tooling to obtain all the metrics needed for the analyses. Issues surrounding the tooling also directly influenced the creation of the dataset. Mutant density is not a widely adapted metric, on top of that, the calculation of this metric is not nearly as simple as source lines of code and cyclomatic complexity. As mentioned in Section 4.1, the only real tool available for the direct calculation of mutant density is Chaosmeter. There are no other tools that provide this functionality for other languages, causing the need for new or altered tools.

New tools falls completely out of scope, although improved tooling would be at the core of potential future work, this is not the intention of this work. Altered tooling was thus the only other option, this is however not without any challenges. Tools for mutation testing exist for different languages since this is a slightly more common procedure. However, not every tool is open source, decently functioning or written in a way that it can be altered for different purposes. Trying different tools that were able to run on the machines available took a lot more time than originally anticipated. The tools described which were used for all the analyses function mostly at this point but there are still remaining flaws that were not covered due to time constraints. This does create an initial risk of inconsistencies within the data created by the tooling.

Creating datasets was also very time consuming due to difficulties with the tooling. Mutpy, after the alterations, still had some functionality in place which caused the need for all the requirements for the project to be in place. This functionality is needed when running the mutation tests but not for the general calculation of the amount of mutations. This is an issue that was not solved since it took more time than intended which needed to be spent on generating and analyzing the results. A similar problem is present for Dextool, the projects that need to be analyzed for this tool need to be able to be build on your machine. In the case of python, is this more easy to work around but for C and C++ this is not always an easy task. Looking through github for opensource C and C++ projects that are sufficient in size and easily compiled on the machine used was a time consuming task. Larger C and C++ projects also take quite some time to build, this implies that difficulties with a chosen project might only arise later in the building process after already spending quite some time on this. At a certain point the choice was made to continue with the data that was available at the time instead of continuing to fine-tune everything surrounding the tooling. This provides a second risk in the available data, the datasets might not always represent similar structures amongst the different programming languages giving a potential reason for the discrepancy between languages present in the results.

Within the selection and creation of metrics that are used for outliers there are also some potential risks present. The first potential risk lies within Section 5.2.1 providing contextual information for outliers. It relies on the differences between metrics and its corresponding visualisation to provide initial context on how and what outliers could potentially be. Taking the difference between two metrics is combining two metrics in a way that they should not necessarily be combined. This could result in potentially wrong results but the effect is limited due to these differences only being used for some initial context and guidelines.

The scaling factor used to force more influence on larger methods combined with average mutant density and average cyclomatic complexity could also be seen as a small threat. This scaling factor is not directly based on existing literature. Although the anecdotal evidence obtained through the usage of this scaling factor were within expectations, this method does not have empirical evidence backing it.

### 6.2 Internal validity

The internal validity discusses if there are any factors which may affect the obtained results. This mainly concerns uncertainty about the exact calculation used by each tool for the different metrics such as source lines of code, mutant density and cyclomatic complexity. Pinpointing the exact influence of this was not within scope but this also means the variation could fall outside of expectations.

### 6.3 External validity

The external validity discusses the extent to which the results of this work can be generalized. For the research that is done in this work, this applies mostly to generalizing across different languages and different types of projects. The

aspect of applying the results over different languages and projects has been covered from the beginning of the project. Especially at the start, there is coverage over Java, C, C++ and Python. Within those languages there is also some variation within the selected projects. Although throughout the work more emphasis is placed on Java, results from other languages that differ from this are discussed.

This work tries to provide a foundation for working with mutant density alongside cyclomatic complexity. Based on the results, there is additional information that can be obtained through the approaches that are mentioned. There are however multiple aspects that hinder or influence the possible generalization of this work.

Choosing the dataset within a language is the first aspect which influences this. In earlier sections, there has been analysis on how working with a single project or a set of projects influences the results. If working with a set of projects were the best option, this would prompt additional research in the selection of those projects. It was shown however that for the data provided, there was no loss of information when only working with singular projects. This approach was therefore used in further analyses. Using only a singular project removes the influence from selecting the projects partially. Complete removal of this influence is however not possible by only using a singular project. Applying metrics such as cyclomatic complexity and mutant density which are directly based on specific parts of the code is always effected heavily by the project in question. Projects containing a more mathematical solution to a problem have higher potential mutant density compared to those that provide a completely different type of solution. This is however, not necessarily something that hinders generalization. Since the outlier detection is based on the same project, more mathematical functions in a project will also influence the mean and standard deviation. Therefore, the influence is limited on this aspect.

What could be more influential is the composition of the projects themselves. What is the balance of mathematical methods compared to those that have only function calls for example? An imbalanced distribution between types of methods might hide a specific outliers. If there is an abnormally large presence of mathematical methods, the mean would accommodate this. The presence of functions with a very low amount of mathematical operations are thus much more likely to be considered outliers based on the current approach. Although the distribution of methods in projects are not likely to behave this way, it remains an influential aspect. Again, this does not completely hinder the generalization, it is more of a reminder of the potential variation that might happen. This specific issue could be addressed in two general ways. The first one would be to investigate some form of thresholding such as those applied for cyclomatic complexity. The second method is more in depth analysis of internal project structures and how it influences the results of applying specific metrics.

The largest potential threat against generalization is choosing the programming language on which the results are applied. In earlier sections, it was already mentioned that there is a difference in the distribution of mutant density over different languages. The same goes for other metrics, especially something like source lines of code. Average mutant density uses both of these metrics and the variation of languages could have a potentially large influence. Over all the languages that are covered in this work the difference is very noticeable. Languages that are not covered in this work might have even more variation. Older languages such as Fortran could give completely different results. The following statements are purely assumptions since the data generated from this work is not sufficient and broad enough to be completely certain.

It is expected that this issue is partially countered by the nature of mutant density itself. The increase in mutant density variation when moving from Java to C or C++, was argued to be caused by the complexity of the languages themselves. Languages such as Java have more built in operators that might not always exist in older or lower level languages. The need for more operations to reach the same functionality is however the same over the complete project. Since the outlier detection works based on the mean and standard deviation, changes throughout the whole project are taken into account this way. Working with average mutant density could also slightly alleviate this issue. This is based on the same reasoning, more operations needed could also correspond to more source lines of code implementing this. Giving a potentially better balance between these two metrics.

These statements do need proper verification but investigating these aspects is beyond the scope of this project itself. This would cause the need for more in depth analysis of internal project structure and of the differences between languages in terms of specific code patterns. Researching proper thresholding for mutant density or any of the other metrics utilizing this concept could also provide a solution. This is a very complex issue which would be sufficient for a work on its own. Having this can circumvent the most prominent problem for generalization at this time. Currently problematic methods or refactoring targets are defined based on outlier detection within a singular project. When starting to work on a new project, there are simply not enough methods to generate outliers upon. This means this method of detecting refactoring targets is currently only suited for projects that already have a significant code base.

Growing a project from the start while using this metric is thus not a realistic approach. Mutant density could still be used to outline the amount of mutations, this would allow developers to use this information on their own accord. A threshold would allow this to be used for the start while giving some clear indications. Due to code variation and general complexity, using a single threshold to define when a function might be problematic, is difficult.

## **6.4 Reliability**

The reliability discusses to degree to which the results depends on the tooling used. As discussed frequently in previous sections every single piece of data is obtained through different tooling. There is no certainty about exact commonalities between the tooling over different languages and the rules they apply internally. This resulted in different tooling that was either supporting the intentions of this work or needed to be customized to adapt this. Within mutpy there are some remaining bugs surrounding requirements needing to be present to generate the mutations which was not removed. This was mostly due to time limitations but this only effects the amount of data returned, not the specifics of the data.

The largest threat is how different tools implement different metrics, there might be small variations possible within the calculation for lines of code. Another potential variation is in the different types of mutation operators that are supported. Dextool provides an easy selection for using different mutation operators which is not present in other tooling. This is not directly an issue within the current context but for expansion on the modularity this does need addressing.

## 7 Conclusion

Cyclomatic complexity has a long standing reputation, although this depends on who you ask, this is not always a positive reputation. Despite this, it has been used for a long time in multiple different industrial tools. As not to pile onto the critique this metric has already received throughout the years, the emphasis was on acknowledgement and supplementing the gaps in this metric. The chosen approach was based on mutant density, a metric that takes principles from mutation testing to determine the code complexity. Calculating the amount of possible mutations gives information on the difficulty of testing and based on how complexity is often described, the code. The goal was to show how well mutant density can be used as the complementary metric to provide additional insight into refactoring opportunities. Not only would this allow more detailed reasoning behind complex functions, it would also allow to show that complex functions can occur in different ways and that mutant density is able to cover others that cyclomatic complexity does not.

This lead to an approach based on the following research questions.

**RQ1.:** *Correlation - what is the correlation between cyclomatic complexity and mutant density?*

- The results provided by the correlation analysis fall in line with the expectations, there is a positive correlation between mutant density and the other metrics but not enough to argue for the redundancy of either of the methods. The correlation is still positive enough to acknowledge a strong sense of similarity between the different metrics. This is reflected in the ability to more directly compare cyclomatic complexity and mutant density. A higher correlation between these two allow the argument of a strong similar basis while providing individual unique results to function within the context of this work.

**RQ2.:** *Complementary - how can we utilize mutant density to provide additional insight into refactoring opportunities? Is there anecdotal evidence supporting this?*

- Using the fundamental ideas provided by the correlation analysis, outliers are calculated by using the z-score on a variety of different metrics. Out of these, the most important individual metrics were the scaled average mutant density and cyclomatic complexity. They provided an initial overview of the individual performance of these metrics in detecting outliers. Combined with the set theory approach, these results also directly show that mutant density views a type of complexity that cyclomatic complexity does not. This is also supported by functions found through this approach.

Applying a more direct metric to measure the relation between cyclomatic complexity and mutant density also gave some interesting perspectives. It purely emphasises the direct relation between those two metrics, it does however skip outlier based on a single metric. The metric does cover a similar basis while also adding a few unique scenarios. It gives functions that do not occur for a singular metric but do occur when they are put in relation to each other. These metrics have been summarized in Table 1 but due to multiple potential threats, this answer cannot be given with full confidence. In specific contexts the metrics do show usable results but due to the lack of generalization it remains hard to pinpoint the exact performance.

**RQ3.:** *Fine Tuning - what is the impact of fine-tuning mutant density on correlation and complementary?*

- Modularity was also mentioned as an important aspect of mutant density. Fine-tuning the possible mutation operators would allow more precise complexity definitions. This fine-tuning allows for vast variety in combinations but most emphasis was placed on more simple applications of this modularity. Running the same outlier detection for each possible mutation operator separately is a much more time consuming method which should be kept in mind. Looking at outliers that only occurred for a single mutation operator did not provide interesting results. It mostly showed the difference in presence of certain operators within a projects. The most interesting results were obtained by looking at methods that were outliers for each single mutation operator. These methods were easily seen as complex due to the presence of too many of each single operator types. Not only that, due to the inclusion of operators of different types, it also affected the size of the functions found without directly incorporating a scaling factor. This opens up a lot more potential surrounding this approach. Although it would make for a computationally heavy metric, creating a metric based on the amount of operators a method is an outlier for, has potential.

In the end, these results provide the foundation for mutant density as a provider for additional information about refactoring opportunities alongside cyclomatic complexity. It has been shown that due to a different approach to complexity, mutant density allows the finding of new unique information without being redundant due too high similarity. Different base metrics have been given that use a combination of cyclomatic complexity, mutant density



and source lines of code. These combinations have their own different advantages and disadvantages that have been provided in earlier sections. Despite the disadvantages, the advantages show to possibilities surrounding mutant density. This results in varied metrics that can find refactoring opportunities which also relate to testing effort.

Although, the possibilities and an original foundation has been shown to exist, there are still a lot of possible improvements. Some improvements are even necessities before larger scale applications of these metrics can exist. These issues have been partially discussed throughout the whole work but with all of the results finished, it is easier to obtain a complete picture on the state of this topic.

The different directions for the future work can be divide among three different target audiences. Tool builders are required for the most prominent part of future work which is the tooling that is available. There is almost no tooling available specifically for calculating mutant density, even popular tools like SonarQube and CodeScene have no support. Meaning most of the tools that are applied, are derived from tooling for mutation testing. This is for C, C++ and partially for python, not done in a static manner. In the context of mutation testing this is appropriate but for mutant density it is not the most efficient solution. A large first step for this topic would be the creation of a general tooling solution that applies static mutant density calculation to different languages. This would also allow for the streamlining of, source lines of code calculation and the variation in mutation operators.

The second type of future work would fall under researches and this is further investigation on the influence of these results under a more precisely defined dataset. Right now the dataset is not large enough and has too many uncertainties to correctly evaluate the generalization of the results. Creating a larger dataset with either a more defined construction, or a size large enough for general results, is quite a difficult task. It is however necessary for more appropriate evaluation of the results presented in this work. Easier and more efficient static tooling would also simplify this issue and would allow for the usage of repository mining and the automation of this process.

Then, there is also the issue of scaling these metrics onto a threshold to allow the usage on smaller sized projects. The usage of outlier detection to determine refactoring targets only works on larger projects as described in this work. Cyclomatic complexity has threshold for determining which functions should be seen as complex. This has however, very little analytical reasoning. It is thus very hard to apply the same or similar thresholding without proper investigation and analysis in this issue. This is also a difficult issue due to all the differences between different projects and programming languages as shown in the results. A singular threshold for across all possible applications is therefor not a really feasible solution.

Another possible approach for utilizing these metrics on smaller scale projects is moving back to calculating the mean and standard deviations across a large set of projects. This would allow the usage of a very general mean that does not need a large project to function upon since it can be used directly. Using a complete dataset would however reintroduce the issue of generalization. It also falls back to the need for a much larger and thought out dataset creation.

If the previous improvements are in place, it becomes easier to further investigate the exact performance of the metrics provided. It also might show certain aspects that could still use slight tweaking. Further analysis of the modular aspect also becomes more feasible after the improvement of the tooling. This metric is very dependant on the implementation and efficiency of the tool so this would be a fundamental step before continuing in this direction.

Lastly, there is future work that influences practitioners that rely on DevOps tooling and certain metrics to apply refactorings. The anecdotal evidence shown throughout this work verifies the aspects of complexity that the current tooling might leave out. Utilizing metrics in a complementary manners shows that different metrics can support each other while also providing new information. On top of that, refactoring opportunities found by multiple metrics at the same time provide more context as to why a refactoring should take place. Refactoring in this expanded manner therefore shows potential for managing maintainability and testability in a new way.

## 8 Relation to previous works

This section will shortly cover my previous research and the relationship with this work if present.

### 8.1 Semantic annotation and navigation in office environments

**Supervisor:** Steven Latré

**Assistant:** Kevin Mets

**Content:** This research looked training networks for navigation in office environments based on semantic annotation. This work has no relation to the current research since it lies in a completely different field of research.

### 8.2 Attention, only intuitive or also an explanation?

**Supervisor:** Toon Calders

**Assistant:** Ewoenam Kwaku Tokpo

**Content:** This research looked into the possibility of using attention in neural networks to provide explanation on decisions in the NLP context. This work has no relation to the current research since it lies in a completely different field of research.

### 8.3 Automatic recognition of molecular graphical sketches

**Supervisor:** José Antonio Oramas Mogrovejo

**Content:** This thesis tried to use convolutional neural networks and other machine learning techniques to attempt to transform drawings of molecular structure to a corresponding formula. This work has no relation to the current research since it lies in a completely different field of research.

## Appendix

---

```
final boolean merge( final SymbolTable symbolTable, final Frame dstFrame, final int
    catchTypeIndex) {
    boolean frameChanged = false;
    int numLocal = inputLocals.length;
    int numStack = inputStack.length;
    if (dstFrame.inputLocals == null) {
        dstFrame.inputLocals = new int[numLocal];
        frameChanged = true;
    }
    for (int i = 0; i < numLocal; ++i) {
        int concreteOutputType;
        if (outputLocals != null && i < outputLocals.length) {
            int abstractOutputType = outputLocals[i];
            if (abstractOutputType == 0) {
                concreteOutputType = inputLocals[i];
            } else {
                concreteOutputType = getConcreteOutputType(abstractOutputType, numStack);
            }
        } else {
            concreteOutputType = inputLocals[i];
        }
        if (initializations != null) {
            concreteOutputType = getInitializedType(symbolTable, concreteOutputType);
        }
        frameChanged |= merge(symbolTable, concreteOutputType, dstFrame.inputLocals, i);
    }
    if (catchTypeIndex > 0) {
        for (int i = 0; i < numLocal; ++i) {
            frameChanged |= merge(symbolTable, inputLocals[i], dstFrame.inputLocals, i);
        }
        if (dstFrame.inputStack == null) {
            dstFrame.inputStack = new int[1];
            frameChanged = true;
        }
        frameChanged |= merge(symbolTable, catchTypeIndex, dstFrame.inputStack, 0);
        return frameChanged;
    }
    int numInputStack = inputStack.length + outputStackStart;
    if (dstFrame.inputStack == null) {
        dstFrame.inputStack = new int[numInputStack + outputStackTop];
        frameChanged = true;
    }
    for (int i = 0; i < numInputStack; ++i) {
        int concreteOutputType = inputStack[i];
        if (initializations != null) {
            concreteOutputType = getInitializedType(symbolTable, concreteOutputType);
        }
        frameChanged |= merge(symbolTable, concreteOutputType, dstFrame.inputStack, i);
    }
    for (int i = 0; i < outputStackTop; ++i) {
        int abstractOutputType = outputStack[i];
        int concreteOutputType = getConcreteOutputType(abstractOutputType, numStack);
        if (initializations != null) {
            concreteOutputType = getInitializedType(symbolTable, concreteOutputType);
        }
        frameChanged |= merge(symbolTable, concreteOutputType, dstFrame.inputStack, numInputStack+i);
    }
    return frameChanged;
}
```

---

Listing 16: Full code example of worst function using scaled average mutant density

---

```

public int getOpcode(final int opcode) {
    if (opcode == Opcodes.IALOAD || opcode == Opcodes.IASTORE) {
        switch (sort) {
            case BOOLEAN:
            case BYTE:
                return opcode + (Opcodes.BALOAD - Opcodes.IALOAD);
            case CHAR:
                return opcode + (Opcodes.CALOAD - Opcodes.IALOAD);
            case SHORT:
                return opcode + (Opcodes.SALOAD - Opcodes.IALOAD);
            case INT:
                return opcode;
            case FLOAT:
                return opcode + (Opcodes.FALOAD - Opcodes.IALOAD);
            case LONG:
                return opcode + (Opcodes.LALOAD - Opcodes.IALOAD);
            case DOUBLE:
                return opcode + (Opcodes.DALOAD - Opcodes.IALOAD);
            case ARRAY:
            case OBJECT:
            case INTERNAL:
                return opcode + (Opcodes.AALOAD - Opcodes.IALOAD);
            case METHOD:
            case VOID:
                throw new UnsupportedOperationException();
            default:
                throw new AssertionError();
        }
    } else {
        switch (sort) {
            case VOID:
                if (opcode != Opcodes.IRETURN) {
                    throw new UnsupportedOperationException();
                }
                return Opcodes.RETURN;
            case BOOLEAN:
            case BYTE:
            case CHAR:
            case SHORT:
            case INT:
                return opcode;
            case FLOAT:
                return opcode + (Opcodes.FRETURN - Opcodes.IRETURN);
            case LONG:
                return opcode + (Opcodes.LRETURN - Opcodes.IRETURN);
            case DOUBLE:
                return opcode + (Opcodes.DRETURN - Opcodes.IRETURN);
            case ARRAY:
            case OBJECT:
            case INTERNAL:
                if (opcode != Opcodes.ILOAD && opcode != Opcodes.ISTORE && opcode != Opcodes.IRETURN) {
                    throw new UnsupportedOperationException();
                }
                return opcode + (Opcodes.ARETURN - Opcodes.IRETURN);
            case METHOD:
                throw new UnsupportedOperationException();
            default:
                throw new AssertionError();
        }
    }
}
}

```

---

Listing 17: Full code example for biggest outlier based on scaled average cyclomatic complexity

---

```

protected Mono<Object> readBody(MethodParameter bodyParam, @Nullable MethodParameter actualParam,
    boolean isBodyRequired, BindingContext bindingContext, ServerWebExchange exchange) {

    ResolvableType bodyType = ResolvableType.forMethodParameter(bodyParam);
    ResolvableType actualType = (actualParam != null ?
        ResolvableType.forMethodParameter(actualParam) : bodyType);
    Class<?> resolvedType = bodyType.resolve();
    ReactiveAdapter adapter = (resolvedType != null ?
        getAdapterRegistry().getAdapter(resolvedType) : null);
    ResolvableType elementType = (adapter != null ? bodyType.getGeneric() : bodyType);
    isBodyRequired = isBodyRequired || (adapter != null && !adapter.supportsEmpty());

    ServerHttpRequest request = exchange.getRequest();
    ServerHttpResponse response = exchange.getResponse();

    MediaType contentType;
    HttpHeaders headers = request.getHeaders();
    try {
        contentType = headers.getContentType();
    }
    catch (InvalidMediaTypeException ex) {
        throw new UnsupportedMediaTypeStatusException(
            "Can't parse Content-Type [" + headers.getFirst("Content-Type") + "]: " +
            ex.getMessage(),
            getSupportedMediaTypes(elementType));
    }

    MediaType mediaType = (contentType != null ? contentType :
        MediaType.APPLICATION_OCTET_STREAM);
    Object[] hints = extractValidationHints(bodyParam);

    if (mediaType.isCompatibleWith(MediaType.APPLICATION_FORM_URLENCODED)) {
        if (logger.isDebugEnabled()) {
            logger.debug("Form data is accessed via ServerWebExchange.getFormData() in WebFlux.");
        }
        return Mono.error(new ResponseStatusException(HttpStatus.UNSUPPORTED_MEDIA_TYPE));
    }

    if (logger.isDebugEnabled()) {
        logger.debug(exchange.getLogPrefix() + (contentType != null ?
            "Content-Type:" + contentType :
            "No Content-Type, using " + MediaType.APPLICATION_OCTET_STREAM));
    }

    for (HttpMessageReader<?> reader : getMessageReaders()) {
        if (reader.canRead(elementType, mediaType)) {
            Map<String, Object> readHints = Hints.from(Hints.LOG_PREFIX_HINT,
                exchange.getLogPrefix());
            if (adapter != null && adapter.isMultiValue()) {
                if (logger.isDebugEnabled()) {
                    logger.debug(exchange.getLogPrefix() + "0..N [" + elementType + "]");
                }
                Flux<?> flux = reader.read(actualType, elementType, request, response, readHints);
                flux = flux.onErrorResume(ex -> Flux.error(handleReadError(bodyParam, ex)));
                if (isBodyRequired) {
                    flux = flux.switchIfEmpty(Flux.error(() -> handleMissingBody(bodyParam)));
                }
                if (hints != null) {
                    flux = flux.doOnNext(target ->
                        validate(target, hints, bodyParam, bindingContext, exchange));
                }
                return Mono.just(adapter.fromPublisher(flux));
            }
            else {

```

```

        // Single-value (with or without reactive type wrapper)
        if (logger.isDebugEnabled()) {
            logger.debug(exchange.getLogPrefix() + "0..1 [" + elementType + "]");
        }
        Mono<?> mono = reader.readMono(actualType, elementType, request, response,
            readHints);
        mono = mono.onErrorResume(ex -> Mono.error(handleReadError(bodyParam, ex)));
        if (isBodyRequired) {
            mono = mono.switchIfEmpty(Mono.error(() -> handleMissingBody(bodyParam)));
        }
        if (hints != null) {
            mono = mono.doOnNext(target ->
                validate(target, hints, bodyParam, bindingContext, exchange));
        }
        return (adapter != null ? Mono.just(adapter.fromPublisher(mono)) : Mono.from(mono));
    }
}

// No compatible reader but body may be empty.

HttpMethod method = request.getMethod();
if (contentType == null && SUPPORTED_METHODS.contains(method)) {
    Flux<DataBuffer> body = request.getBody().doOnNext(buffer -> {
        DataBufferUtils.release(buffer);
        // Body not empty, back to 415..
        throw new UnsupportedMediaTypeStatusException(
            mediaType, getSupportedMediaTypes(elementType), elementType);
    });
    if (isBodyRequired) {
        body = body.switchIfEmpty(Mono.error(() -> handleMissingBody(bodyParam)));
    }
    return (adapter != null ? Mono.just(adapter.fromPublisher(body)) : Mono.from(body));
}

return Mono.error(new UnsupportedMediaTypeStatusException(
    mediaType, getSupportedMediaTypes(elementType), elementType));
}

```

---

Listing 18: Full code example for second biggest outlier based on scaled average cyclomatic complexity

---

```

public Resource findLocalizedResource(String name, String extension, @Nullable Locale locale) {
    Assert.notNull(name, "Name must not be null");
    Assert.notNull(extension, "Extension must not be null");

    Resource resource = null;

    if (locale != null) {
        String lang = locale.getLanguage();
        String country = locale.getCountry();
        String variant = locale.getVariant();

        if (variant.length() > 0) {
            String location =
                name + this.separator + lang + this.separator + country + this.separator +
                variant + extension;
            resource = this.resourceLoader.getResource(location);
        }
        if ((resource == null || !resource.exists()) && country.length() > 0) {
            String location = name + this.separator + lang + this.separator + country + extension;
            resource = this.resourceLoader.getResource(location);
        }
        if ((resource == null || !resource.exists()) && lang.length() > 0) {
            String location = name + this.separator + lang + extension;

```

```

        resource = this.resourceLoader.getResource(location);
    }
}
if (resource == null || !resource.exists()) {
    String location = name + extension;
    resource = this.resourceLoader.getResource(location);
}
return resource;
}
}

```

---

Listing 19: First full code example of unique information from average mutant density

---

```

public Object invoke(Object proxy, Method method, Object[] args) throws Throwable {
    switch (method.getName()) {
        case "equals":
            return (proxy == args[0]);
        case "hashCode":
            return System.identityHashCode(proxy);
        case "getTargetConnection":
            return getTargetConnection(method);
        case "unwrap":
            if (((Class<?>) args[0]).isInstance(proxy)) {
                return proxy;
            }
            break;
        case "isWrapperFor":
            if (((Class<?>) args[0]).isInstance(proxy)) {
                return true;
            }
            break;
    }
    if (!hasTargetConnection()) {
        switch (method.getName()) {
            case "toString":
                return "Lazy Connection proxy for target DataSource [" + getTargetDataSource() + "]";
            case "getAutoCommit":
                if (this.autoCommit != null) {return this.autoCommit;}
                break;
            case "setAutoCommit":
                this.autoCommit = (Boolean) args[0];
                return null;
            case "getTransactionIsolation":
                if (this.transactionIsolation != null) {return this.transactionIsolation;}
                break;
            case "setTransactionIsolation":
                this.transactionIsolation = (Integer) args[0];
                return null;
            case "isReadOnly":
                return this.readOnly;
            case "setReadOnly":
                this.readOnly = (Boolean) args[0];
                return null;
            case "getHoldability":
                return this.holdability;
            case "setHoldability":
                this.holdability = (Integer) args[0];
                return null;
            case "commit":
            case "rollback":
                return null;
            case "getWarnings":
            case "clearWarnings":
                return null;
            case "close":

```

```

        this.closed = true;
        return null;
    case "isClosed":
        return this.closed;
    default:
        if (this.closed) {
            throw new SQLException("Illegal operation: connection is closed");
        }
    }
}
...
}

```

---

Listing 20: Full code example of outlier unique for average cyclomatic complexity

---

```

public static boolean areBoxingCompatible(String desc1, String desc2) {
    if (desc1.equals(desc2)) {
        return true;
    }
    if (desc1.length() == 1) {
        if (desc1.equals("Z")) {
            return desc2.equals("Ljava/lang/Boolean");
        }
        else if (desc1.equals("D")) {
            return desc2.equals("Ljava/lang/Double");
        }
        else if (desc1.equals("F")) {
            return desc2.equals("Ljava/lang/Float");
        }
        else if (desc1.equals("I")) {
            return desc2.equals("Ljava/lang/Integer");
        }
        else if (desc1.equals("J")) {
            return desc2.equals("Ljava/lang/Long");
        }
    }
    else if (desc2.length() == 1) {
        if (desc2.equals("Z")) {
            return desc1.equals("Ljava/lang/Boolean");
        }
        else if (desc2.equals("D")) {
            return desc1.equals("Ljava/lang/Double");
        }
        else if (desc2.equals("F")) {
            return desc1.equals("Ljava/lang/Float");
        }
        else if (desc2.equals("I")) {
            return desc1.equals("Ljava/lang/Integer");
        }
        else if (desc2.equals("J")) {
            return desc1.equals("Ljava/lang/Long");
        }
    }
    return false;
}

```

---

Listing 21: Full code example of outlier based on a higher than average cyclomatic complexity compared to mutant density



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