

Software Testing Lab

Assignment 6

Submission Deadline: April 1st, 12:00

MUTATION TESTING

Mutation testing is a technique to measure the quality of a test suite by assessing its fault detection capabilities. Mutation testing starts with a *green* test suite, i.e. a test suite in which all the tests pass. First, a faulty version of the software is created by introducing faults into the system (*Mutation*). This is done by applying a known transformation (*Mutation Operator*) on a certain part of the code. After generating the faulty version of the software (*Mutant*), it is passed onto the test suite. If there is an error or failure during the execution of the test suite, the mutant is marked as killed (*Killed Mutant*). If all tests pass, it means that the test suite could not catch the fault, and the mutant has survived (*Survived Mutant*).

Mutation testing allows software engineers to monitor the fault detection capability of a test suite by means of mutation coverage (see Equation 1). If the output of a mutant for all possible inputs is the same as the original program, it is called an *equivalent mutant*. It is not possible to create a test case that passes for the original program and fails for an equivalent mutant, because the equivalent mutant has the same semantics as the original program. A test suite is said to achieve *full mutation test adequacy* whenever it can kill all the non-equivalent mutants, thus reaching a mutation coverage of 100%. Such test suite is called a *mutation-adequate test suite*.

$$\text{Mutation Coverage} = \frac{\text{Number of killed mutants}}{\text{Number of all non-equivalent mutants}} \quad (1)$$

In this assignment we will use LittleDarwin. LittleDarwin is a mutation testing tool to provide mutation testing within a continuous integration environment. It is designed to have a loose coupling with the test infrastructure, instead relying on the build system to run the test suite.

- **Exercise 1.** Use LittleDarwin¹ to analyze the original version of JPacMan. Explain the results. Now, use LittleDarwin to analyze *your* version of JPacMan. What differences can you see? **(Required, 10 points)**
- **Exercise 2.** Repeat the previous exercise, but this time use branch coverage (with JaCoCo) instead of mutation coverage. Look for a class with 100% branch coverage and less than 100% mutation coverage. Why did this happen? Take similar examples and explain the difference between two results. **(Required, 20 points)**
- **Exercise 3.** Find a killed mutant. Why was it killed? **(Required, 5 points)**
- **Exercise 4.** Take a survived mutant for class Engine, and write a test that kills it. Repeat the process until all survived mutants are covered. Move the previous results out of the way and rerun LittleDarwin to confirm. **(Required, 20 points)**
- **Exercise 5.** Can you find an example of an equivalent mutant? How do you know it is equivalent? **(Required, 5 points)**
- **Exercise 6.** Make a mutation-adequate test suite for JPacMan. How many tests did you have to write? How many equivalent mutants did you find? **(Required, 30 points)**
- **Exercise 7.** Repeat the exercises using PITest and Javalanche. Explain the difference in the results. **(Optional)**
- **Exercise 8.** What are the upsides and downsides of mutation testing? Explain your argument. **(Required, 10 points)**

¹LittleDarwin's man page: <https://littledarwin.parsai.net/release/littledarwin-manpage.pdf>