

# Comparing Different Unit Test Generation Methods Through Different Mutation Operators

## Installing and loading packages

```
install.packages("ggpubr")
```

```
## Installing package into '/home/beatriz/R/x86_64-pc-linux-gnu-library/3.4'  
## (as 'lib' is unspecified)
```

```
install.packages("ggplot2")
```

```
## Installing package into '/home/beatriz/R/x86_64-pc-linux-gnu-library/3.4'  
## (as 'lib' is unspecified)
```

```
install.packages("reshape")
```

```
## Installing package into '/home/beatriz/R/x86_64-pc-linux-gnu-library/3.4'  
## (as 'lib' is unspecified)
```

```
install.packages("reshape2")
```

```
## Installing package into '/home/beatriz/R/x86_64-pc-linux-gnu-library/3.4'  
## (as 'lib' is unspecified)
```

```
install.packages("dplyr")
```

```
## Installing package into '/home/beatriz/R/x86_64-pc-linux-gnu-library/3.4'  
## (as 'lib' is unspecified)
```

```
library("ggplot2")  
library("reshape")  
library("reshape2")
```

```
##  
## Attaching package: 'reshape2'
```

```
## The following objects are masked from 'package:reshape':  
##  
## colsplit, melt, recast
```

```
library("ggpubr")
```

```
## Loading required package: magrittr
```

```
library("dplyr")
```

```
##  
## Attaching package: 'dplyr'
```

```
## The following object is masked from 'package:reshape':  
##  
##      rename
```

```
## The following objects are masked from 'package:stats':  
##  
##      filter, lag
```

```
## The following objects are masked from 'package:base':  
##  
##      intersect, setdiff, setequal, union
```

## Reading data

```
evoResults1 <- read.csv('/home/beatriz/Desktop/FinaldataSet/evoResultsEvoAvg.csv', sep = ",", header = TRUE)  
evoResults2 <- read.csv('/home/beatriz/Desktop/FinaldataSet/evoResults2EvoAvg.csv', sep = ",", header = TRUE)  
evoResults3 <- read.csv('/home/beatriz/Desktop/FinaldataSet/evoResults3EvoAvg.csv', sep = ",", header = TRUE)  
  
randoopResults1 <- read.csv('/home/beatriz/Desktop/FinaldataSet/randoopResultsRandoopAvg.csv', sep = ",", header = TRUE)  
randoopResults2 <- read.csv('/home/beatriz/Desktop/FinaldataSet/randoopResults2RandoopAvg.csv', sep = ",", header = TRUE)  
randoopResults3 <- read.csv('/home/beatriz/Desktop/FinaldataSet/randoopResults3RandoopAvg.csv', sep = ",", header = TRUE)  
  
manualResults1 <- read.csv('/home/beatriz/Desktop/FinaldataSet/manualSuitesResultsPITAvg.csv', sep = ",", header = TRUE)  
manualResults2 <- read.csv('/home/beatriz/Desktop/FinaldataSet/manualSuitesResults2PITAvg.csv', sep = ",", header = TRUE)  
manualResults3 <- read.csv('/home/beatriz/Desktop/FinaldataSet/manualSuitesResults3PITAvg.csv', sep = ",", header = TRUE)
```

## Merging data

```

evoResults12 <- merge(evoResults1, evoResults2, all = TRUE)
#evo12
evoResults123 <- merge(evoResults12, evoResults3, all = TRUE)
evoResults123

```

```

##      MutationCoverage ReturnValsMutator ConditionalsBoundaryMutator
## 1      0.2342513      0.3130890      0.1190476
## 2      0.2386252      0.3582857      0.1266667
## 3      0.3170927      0.4830937      0.3771800
## 4      0.3175885      0.3815475      0.1591111
## 5      0.4180821      0.5737705      0.3207164
## 6      0.4906863      0.5329545      0.3189189
## 7      0.5098667      0.6414414      0.3600000
## 8      0.5388335      0.6093407      0.4948718
## 9      0.6936607      0.6723262      0.6792952
## 10     0.7235751      0.8843750      0.4694581
##      IncrementsMutator NegateConditionalsMutator MathMutator LineCoverage
## 1      0.3000000      0.2519313 0.07142857 0.3844465
## 2      0.1000000      0.2315603 0.16400000 0.3330896
## 3      0.2903226      0.2956539 0.29569378 0.5184696
## 4      0.2000000      0.3388959 0.25837500 0.4458619
## 5      0.3668317      0.4334659 0.41811633 0.6117504
## 6      0.6909091      0.5434978 0.49718310 0.6814879
## 7      0.9000000      0.5099707 0.32592593 0.7715347
## 8      0.3411765      0.5413497 0.57931034 0.6424541
## 9      0.7691824      0.8096410 0.65455508 0.8792263
## 10     0.5200000      0.7894340 0.68588957 0.9378749
##      Project Tool InvertNegsMutator VoidMethodCallMutator
## 1      e-mail  evo      NA      0.10769231
## 2      csv    evo      NA      0.07419355
## 3      bcel   evo      NA      0.22429358
## 4      lang   evo      0.6500000 0.31827309
## 5      imaging evo      0.5833333 0.25390625
## 6      fileupload evo      NA      0.27710843
## 7      cli     evo      NA      0.32051282
## 8      validator evo      NA      0.16944444
## 9      codec   evo      NA      0.46905537
## 10     statistics evo      0.6657143      NA

```

```

randoopResults12 <- merge(randoopResults1, randoopResults2, all = TRUE)
#randoopResults12
randoopResults123 <- merge(randoopResults12, randoopResults3, all = TRUE)
#randoopResults123

manualResults12 <- merge(manualResults1, manualResults2, all = TRUE)
#manualResults12
manualResults123 <- merge(manualResults12, manualResults3, all = TRUE)
manualResults123

```

```
##      MutationCoverage ReturnValsMutator ConditionalsBoundaryMutator
## 1      0.3048661      0.3513514      0.3432282
## 2      0.4836810      0.4124688      0.4151261
## 3      0.6161790      0.5863874      0.2857143
## 4      0.6388889      0.5852273      0.4324324
## 5      0.7572283      0.7596154      0.6153846
## 6      0.8267091      0.8974359      0.5333333
## 7      0.8656351      0.8918300      0.5274074
## 8      0.8672619      0.8302139      0.6563877
## 9      0.8797280      0.8216146      0.7187192
## 10     0.8986667      0.8468468      0.6571429
##      IncrementsMutator NegateConditionalsMutator MathMutator LineCoverage
## 1      0.4946237      0.3615172      0.3803828      0.4786764
## 2      0.7116705      0.5370138      0.5001359      0.6528049
## 3      1.0000000      0.7467811      0.7142857      0.8151876
## 4      0.6363636      0.7982063      0.7464789      0.7638408
## 5      0.8529412      0.8269939      0.8505747      0.7782152
## 6      0.7272727      0.8936170      0.3673469      0.9180000
## 7      0.9535963      0.9640440      0.8100000      0.9531415
## 8      0.9056604      0.9353846      0.8781780      0.9208742
## 9      1.0000000      0.9671698      0.9259202      0.9228792
## 10     1.0000000      0.9618768      0.9259259      0.9669967
##      Project   Tool InvertNegsMutator VoidMethodCallMutator
## 1      bcel manual      NA      0.1904587
## 2      imaging manual    0.1666667      0.4241540
## 3      e-mail manual      NA      0.4358974
## 4      fileupload manual    NA      0.3253012
## 5      validator manual    NA      0.4166667
## 6      csv manual      NA      0.8225806
## 7      lang manual      0.9500000      0.8032129
## 8      codec manual      NA      0.8436482
## 9      statistics manual    0.9114286      NA
## 10     cli manual      NA      0.8717949
```

```
evoAndRandoopDataSet <- merge(evoResults123, randoopResults123, all = TRUE)
completeDataSet <- merge(evoAndRandoopDataSet, manualResults123, all = TRUE)
#completeDataSet
```

## Declaring common values

```
metrics1 = c("LineCoverage","MutationCoverage")
metrics2 = c("ReturnValsMutator","ConditionalsBoundaryMutator",
             "VoidMethodCallMutator","IncrementsMutator",
             "NegateConditionalsMutator","MathMutator", "InvertNegsMutator")

projects=c("BCEL","CLI","Codec", "CSV", "Email", "FileUpload", "Imaging", "Lang", "Statistics", "Validator")
```

## Melting data

```
completeDataSet.m1 <- melt(completeDataSet, id.var = "Tool", measure.vars = metrics1)
#completeDataSet.m1

completeDataSet.m2 <- melt(completeDataSet, id.var = "Tool", measure.vars = metrics2)
#completeDataSet.m2

completeDataSet.m3 <- melt(completeDataSet, id.vars = c("Tool", "Project"))
#completeDataSet.m3
```

## Casting data

```
completeDataSet.c1 <- cast(completeDataSet.m3, Tool~variable+Project)
completeDataSet.c1
```

```

##      Tool MutationCoverage_bcel MutationCoverage_cli
## 1      evo          0.3170927          0.5098667
## 2 randoop          0.1407471          0.3025333
## 3 manual           0.3048661          0.8986667
##      MutationCoverage_codec MutationCoverage_csv MutationCoverage_e-mail
## 1          0.6936607          0.2386252          0.2342513
## 2          0.6430357          0.4646481          0.4535284
## 3          0.8672619          0.8267091          0.6161790
##      MutationCoverage_fileupload MutationCoverage_validator
## 1          0.4906863          0.5388335
## 2          0.2506536          0.4467597
## 3          0.6388889          0.7572283
##      MutationCoverage_imaging MutationCoverage_lang
## 1          0.4180821          0.3175885
## 2          0.1402246          0.3148940
## 3          0.4836810          0.8656351
##      MutationCoverage_statistics ReturnValsMutator_bcel ReturnValsMutator_cli
## 1          0.7235751          0.4830937          0.6414414
## 2          0.7407383          0.2441633          0.5387387
## 3          0.8797280          0.3513514          0.8468468
##      ReturnValsMutator_codec ReturnValsMutator_csv ReturnValsMutator_e-mail
## 1          0.6723262          0.3582857          0.3130890
## 2          0.7030749          0.5828571          0.5732984
## 3          0.8302139          0.8974359          0.5863874
##      ReturnValsMutator_fileupload ReturnValsMutator_validator
## 1          0.5329545          0.6093407
## 2          0.3659091          0.5890110
## 3          0.5852273          0.7596154
##      ReturnValsMutator_imaging ReturnValsMutator_lang
## 1          0.5737705          0.3815475
## 2          0.2627561          0.3860130
## 3          0.4124688          0.8918300
##      ReturnValsMutator_statistics ConditionalsBoundaryMutator_bcel
## 1          0.8843750          0.3771800
## 2          0.8575521          0.1150278
## 3          0.8216146          0.3432282
##      ConditionalsBoundaryMutator_cli ConditionalsBoundaryMutator_codec
## 1          0.3600000          0.6792952
## 2          0.1114286          0.4964758
## 3          0.6571429          0.6563877
##      ConditionalsBoundaryMutator_csv ConditionalsBoundaryMutator_e-mail
## 1          0.1266667          0.1190476
## 2          0.2333333          0.2333333
## 3          0.5333333          0.2857143
##      ConditionalsBoundaryMutator_fileupload
## 1          0.3189189
## 2          0.1432432
## 3          0.4324324
##      ConditionalsBoundaryMutator_validator
## 1          0.4948718
## 2          0.3380342
## 3          0.6153846
##      ConditionalsBoundaryMutator_imaging ConditionalsBoundaryMutator_lang
## 1          0.32071636          0.1591111
## 2          0.06873185          0.1879867
## 3          0.41512605          0.5274074
##      ConditionalsBoundaryMutator_statistics IncrementsMutator_bcel

```

## 1	0.4694581	0.2903226
## 2	0.3197044	0.1591398
## 3	0.7187192	0.4946237
## IncrementsMutator_cli IncrementsMutator_codec IncrementsMutator_csv		
## 1	0.9000	0.7691824
## 2	0.4125	0.5981132
## 3	1.0000	0.9056604
## IncrementsMutator_e-mail IncrementsMutator_fileupload		
## 1	0.30	0.6909091
## 2	0.72	0.1590909
## 3	1.00	0.6363636
## IncrementsMutator_validator IncrementsMutator_imaging		
## 1	0.3411765	0.3668317
## 2	0.2911765	0.1247525
## 3	0.8529412	0.7116705
## IncrementsMutator_lang IncrementsMutator_statistics		
## 1	0.2000000	0.52
## 2	0.3703989	0.70
## 3	0.9535963	1.00
## NegateConditionalsMutator_bcel NegateConditionalsMutator_cli		
## 1	0.2956539	0.5099707
## 2	0.1658633	0.2504399
## 3	0.3615172	0.9618768
## NegateConditionalsMutator_codec NegateConditionalsMutator_csv		
## 1	0.8096410	0.2315603
## 2	0.7218462	0.5131206
## 3	0.9353846	0.8936170
## NegateConditionalsMutator_e-mail NegateConditionalsMutator_fileupload		
## 1	0.2519313	0.5434978
## 2	0.5206009	0.2677130
## 3	0.7467811	0.7982063
## NegateConditionalsMutator_validator NegateConditionalsMutator_imaging		
## 1	0.5413497	0.4334659
## 2	0.4101840	0.1556991
## 3	0.8269939	0.5370138
## NegateConditionalsMutator_lang NegateConditionalsMutator_statistics		
## 1	0.3388959	0.7894340
## 2	0.3648787	0.8641509
## 3	0.9640440	0.9671698
## MathMutator_bcel MathMutator_cli MathMutator_codec MathMutator_csv		
## 1	0.2956938	0.3259259
## 2	0.1167464	0.1259259
## 3	0.3803828	0.9259259
## MathMutator_e-mail MathMutator_fileupload MathMutator_validator		
## 1	0.07142857	0.4971831
## 2	0.32857143	0.2267606
## 3	0.71428571	0.7464789
## MathMutator_imaging MathMutator_lang MathMutator_statistics		
## 1	0.4181163	0.2583750
## 2	0.1175827	0.2149812
## 3	0.5001359	0.8100000
## LineCoverage_bcel LineCoverage_cli LineCoverage_codec LineCoverage_csv		
## 1	0.5184696	0.7715347
## 2	0.3889611	0.5235149
## 3	0.4786764	0.9669967
## LineCoverage_e-mail LineCoverage_fileupload LineCoverage_validator		
## 1	0.3844465	0.6814879
## 2	0.7168344	0.4693772

```

## 3      0.8151876      0.7638408      0.7782152
## LineCoverage_imaging LineCoverage_lang LineCoverage_statistics
## 1      0.6117504      0.4458619      0.9378749
## 2      0.3336928      0.5535309      0.9050557
## 3      0.6528049      0.9531415      0.9228792
## InvertNegsMutator_bcel InvertNegsMutator_cli InvertNegsMutator_codec
## 1      NA      NA      NA
## 2      NA      NA      NA
## 3      NA      NA      NA
## InvertNegsMutator_csv InvertNegsMutator_e-mail
## 1      NA      NA
## 2      NA      NA
## 3      NA      NA
## InvertNegsMutator_fileupload InvertNegsMutator_validator
## 1      NA      NA
## 2      NA      NA
## 3      NA      NA
## InvertNegsMutator_imaging InvertNegsMutator_lang
## 1      0.5833333      0.65
## 2      0.1500000      0.39
## 3      0.1666667      0.95
## InvertNegsMutator_statistics VoidMethodCallMutator_bcel
## 1      0.6657143      0.22429358
## 2      0.8142857      0.05893578
## 3      0.9114286      0.19045872
## VoidMethodCallMutator_cli VoidMethodCallMutator_codec
## 1      0.3205128      0.4690554
## 2      0.0965812      0.5638436
## 3      0.8717949      0.8436482
## VoidMethodCallMutator_csv VoidMethodCallMutator_e-mail
## 1      0.07419355      0.1076923
## 2      0.33548387      0.1675214
## 3      0.82258065      0.4358974
## VoidMethodCallMutator_fileupload VoidMethodCallMutator_validator
## 1      0.27710843      0.1694444
## 2      0.05301205      0.1333333
## 3      0.32530120      0.4166667
## VoidMethodCallMutator_imaging VoidMethodCallMutator_lang
## 1      0.25390625      0.3182731
## 2      0.05426136      0.1523728
## 3      0.42415403      0.8032129
## VoidMethodCallMutator_statistics
## 1      NA
## 2      NA
## 3      NA

```

## Plotting Box-plots

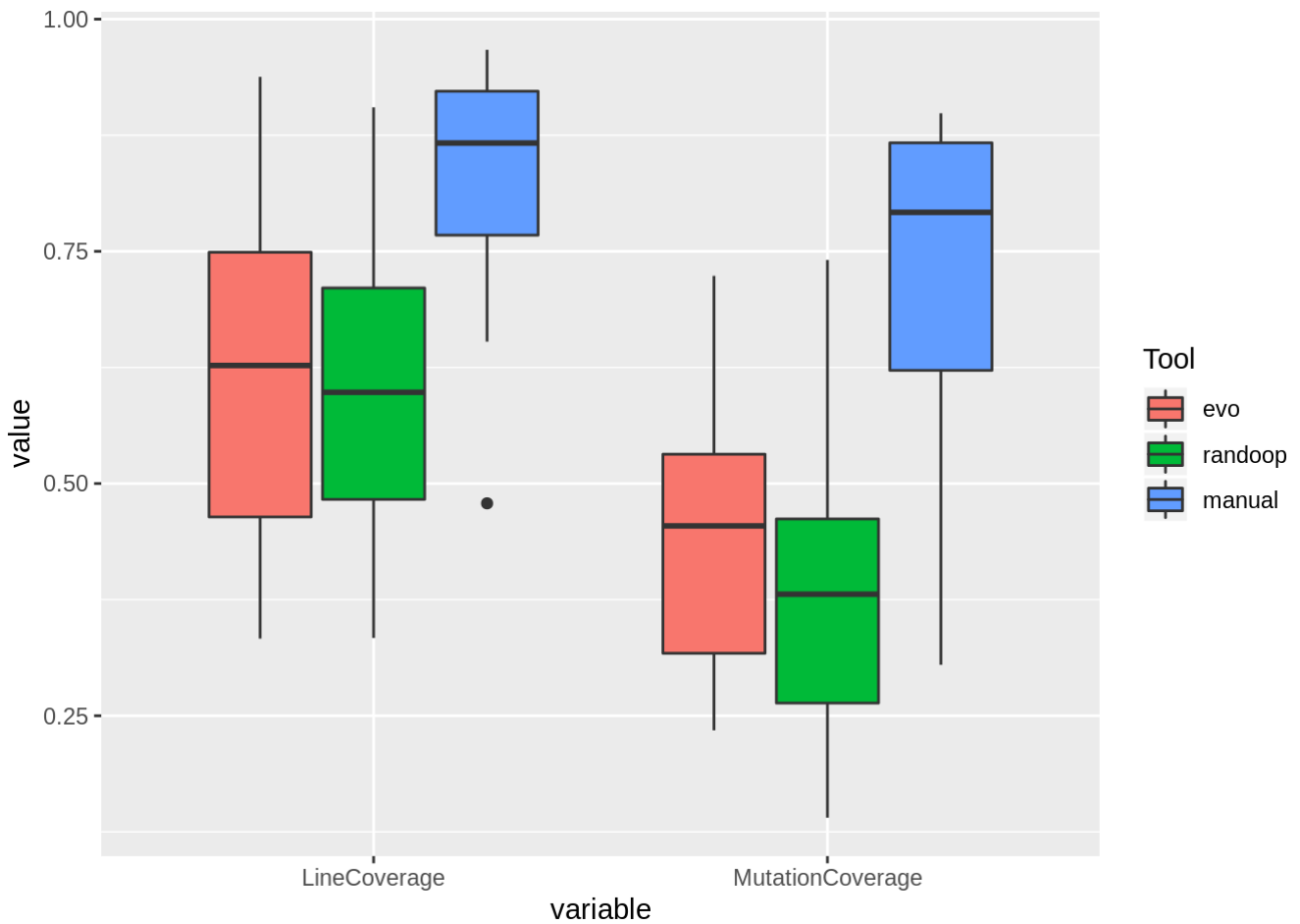
## Line Coverage and Mutation Coverage

```

ggplot(data = completeDataSet.m1, aes(x=variable, y=value)) +
  geom_boxplot(aes(fill=Tool)) +
  scale_x_discrete(labels= c("LineCoverage", "MutationCoverage"))

```

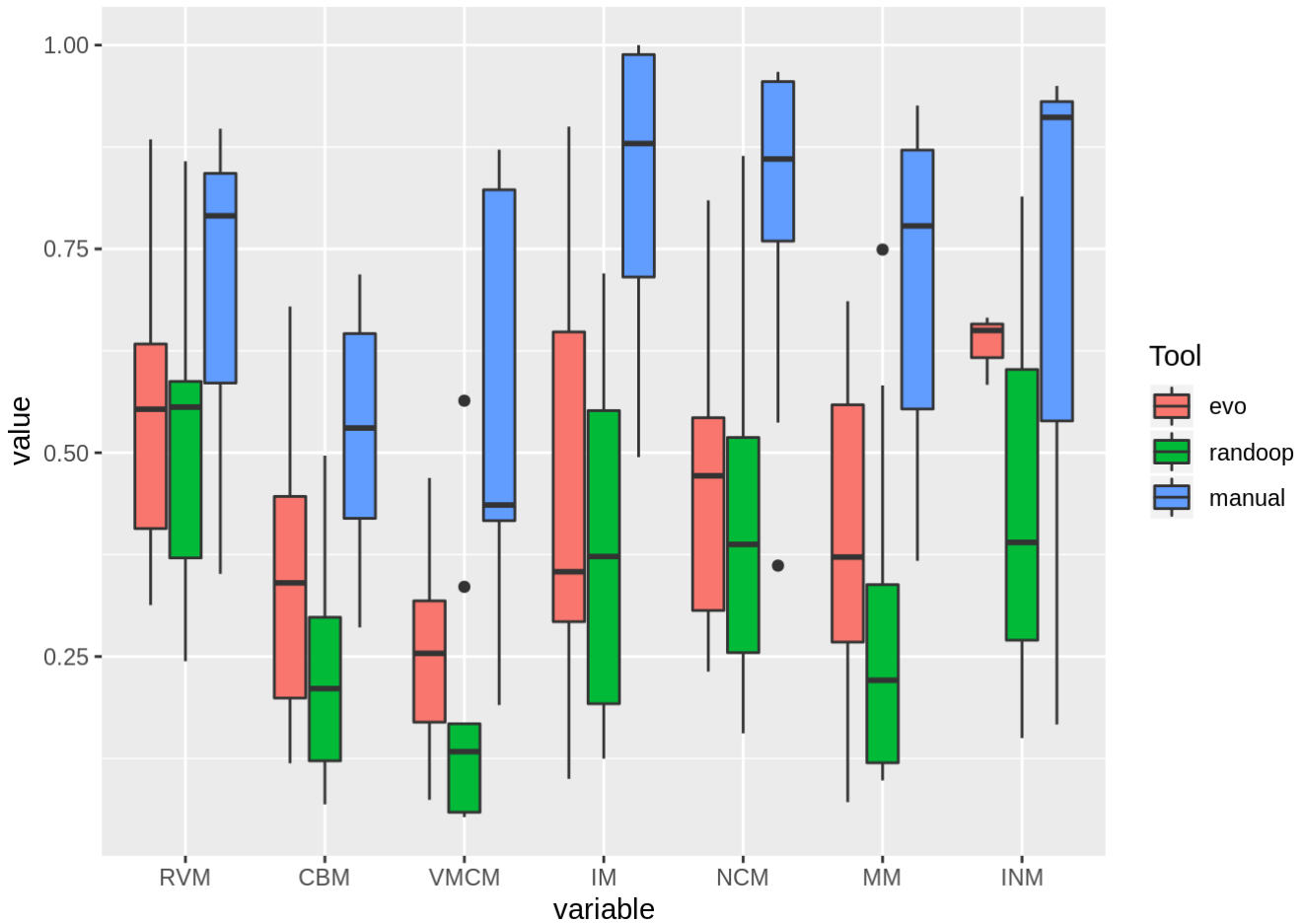




## PIT's Default Mutation Operators

```
ggplot(data = completeDataSet.m2, aes(x=variable, y=value)) +  
  geom_boxplot(aes(fill=Tool)) +  
  scale_x_discrete(labels= c(  
    "RVM", "CBM",  
    "VMCM", "IM",  
    "NCM", "MM", "INM"))
```

```
## Warning: Removed 24 rows containing non-finite values (stat_boxplot).
```



## Comparing Line Coverage per Tool

```
wilcox.test(evoResults123$LineCoverage, randoopResults123$LineCoverage, paired = TRUE
)
```

```
##
## Wilcoxon signed rank test
##
## data:  evoResults123$LineCoverage and randoopResults123$LineCoverage
## V = 32, p-value = 0.6953
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(evoResults123$LineCoverage, manualResults123$LineCoverage, paired = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  evoResults123$LineCoverage and manualResults123$LineCoverage
## V = 0, p-value = 0.001953
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(randoopResults123$LineCoverage, manualResults123$LineCoverage, paired = T
RUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  randoopResults123$LineCoverage and manualResults123$LineCoverage  
## V = 0, p-value = 0.001953  
## alternative hypothesis: true location shift is not equal to 0
```

## Comparing Mutation Coverage per Tool

```
wilcox.test(evoResults123$MutationCoverage, randoopResults123$MutationCoverage, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  evoResults123$MutationCoverage and randoopResults123$MutationCoverage  
## V = 53, p-value = 0.005859  
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(evoResults123$MutationCoverage, manualResults123$MutationCoverage, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  evoResults123$MutationCoverage and manualResults123$MutationCoverage  
## V = 0, p-value = 0.001953  
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(randoopResults123$MutationCoverage, manualResults123$MutationCoverage, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  randoopResults123$MutationCoverage and manualResults123$MutationCoverage  
## V = 0, p-value = 0.001953  
## alternative hypothesis: true location shift is not equal to 0
```

## Comparing ReturnValsMutator Coverage per Tool

```
wilcox.test(evoResults123$ReturnValsMutator, randoopResults123$ReturnValsMutator, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  evoResults123$ReturnValsMutator and randoopResults123$ReturnValsMutator  
## V = 38, p-value = 0.3223  
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(evoResults123$ReturnValsMutator, manualResults123$ReturnValsMutator, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  evoResults123$ReturnValsMutator and manualResults123$ReturnValsMutator  
## V = 1, p-value = 0.003906  
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(randoopResults123$ReturnValsMutator, manualResults123$ReturnValsMutator, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  randoopResults123$ReturnValsMutator and manualResults123$ReturnValsMutator  
## V = 1, p-value = 0.003906  
## alternative hypothesis: true location shift is not equal to 0
```

## Comparing ConditionalsBoundaryMutator Coverage per Tool

```
wilcox.test(evoResults123$ConditionalsBoundaryMutator, randoopResults123$ConditionalsBoundaryMutator, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  evoResults123$ConditionalsBoundaryMutator and randoopResults123$ConditionalsBoundaryMutator  
## V = 53, p-value = 0.005859  
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(evoResults123$ConditionalsBoundaryMutator, manualResults123$ConditionalsBoundaryMutator, paired = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  evoResults123$ConditionalsBoundaryMutator and manualResults123$Conditionals
BoundaryMutator
## V = 2, p-value = 0.005859
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(randoopResults123$ConditionalsBoundaryMutator, manualResults123$Condition
alsBoundaryMutator, paired = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  randoopResults123$ConditionalsBoundaryMutator and manualResults123$Conditio
nalsBoundaryMutator
## V = 0, p-value = 0.001953
## alternative hypothesis: true location shift is not equal to 0
```

## Comparing VoidMethodCallMutator Coverage per Tool

```
wilcox.test(evoResults123$VoidMethodCallMutator, randoopResults123$VoidMethodCallMuta
tor, paired = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  evoResults123$VoidMethodCallMutator and randoopResults123$VoidMethodCallMut
ator
## V = 35, p-value = 0.1641
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(evoResults123$VoidMethodCallMutator, manualResults123$VoidMethodCallMutat
or, paired = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  evoResults123$VoidMethodCallMutator and manualResults123$VoidMethodCallMuta
tor
## V = 0, p-value = 0.007813
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(randoopResults123$VoidMethodCallMutator, manualResults123$VoidMethodCallM
utator, paired = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  randoopResults123$VoidMethodCallMutator and manualResults123$VoidMethodCall
Mutator
## V = 0, p-value = 0.007813
## alternative hypothesis: true location shift is not equal to 0
```

## Comparing IncrementsMutator Coverage per Tool

```
wilcox.test(evoResults123$IncrementsMutator, randoopResults123$IncrementsMutator, pai
red = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  evoResults123$IncrementsMutator and randoopResults123$IncrementsMutator
## V = 33, p-value = 0.625
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(evoResults123$IncrementsMutator, manualResults123$IncrementsMutator, pair
ed = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  evoResults123$IncrementsMutator and manualResults123$IncrementsMutator
## V = 0, p-value = 0.001953
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(randoopResults123$IncrementsMutator, manualResults123$IncrementsMutator,
paired = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  randoopResults123$IncrementsMutator and manualResults123$IncrementsMutator
## V = 0, p-value = 0.001953
## alternative hypothesis: true location shift is not equal to 0
```

## Comparing NegateConditionalsMutator Coverage per Tool

```
wilcox.test(evoResults123$NegateConditionalsMutator, randoopResults123$NegateConditio
nalsMutator, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  evoResults123$NegateConditionalsMutator and randoopResults123$NegateConditionalsMutator  
## V = 48, p-value = 0.03711  
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(evoResults123$NegateConditionalsMutator, manualResults123$NegateConditionalsMutator, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  evoResults123$NegateConditionalsMutator and manualResults123$NegateConditionalsMutator  
## V = 0, p-value = 0.001953  
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(randoopResults123$NegateConditionalsMutator, manualResults123$NegateConditionalsMutator, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  randoopResults123$NegateConditionalsMutator and manualResults123$NegateConditionalsMutator  
## V = 0, p-value = 0.001953  
## alternative hypothesis: true location shift is not equal to 0
```

## Comparing MathMutator Coverage per Tool

```
wilcox.test(evoResults123$MathMutator, randoopResults123$MathMutator, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  evoResults123$MathMutator and randoopResults123$MathMutator  
## V = 48, p-value = 0.03711  
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(evoResults123$MathMutator, manualResults123$MathMutator, paired = TRUE)
```

```
##  
## Wilcoxon signed rank test  
##  
## data:  evoResults123$MathMutator and manualResults123$MathMutator  
## V = 1, p-value = 0.003906  
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(randoopResults123$MathMutator, manualResults123$MathMutator, paired = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  randoopResults123$MathMutator and manualResults123$MathMutator
## V = 0, p-value = 0.001953
## alternative hypothesis: true location shift is not equal to 0
```

## Comparing InvertNegsMutator Coverage per Tool

```
evoResults123WithoutNA <- evoResults123 %>% select(InvertNegsMutator) %>% filter(!is.na(InvertNegsMutator))
#evoResults123WithoutNA
randoopResults123WithoutNA <- randoopResults123 %>% select(InvertNegsMutator) %>% filter(!is.na(InvertNegsMutator))
#randoopResults123WithoutNA
manualResults123WithoutNA <- manualResults123 %>% select(InvertNegsMutator) %>% filter(!is.na(InvertNegsMutator))
#manualResults123WithoutNA

wilcox.test(evoResults123WithoutNA$InvertNegsMutator, randoopResults123WithoutNA$InvertNegsMutator, paired = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  evoResults123WithoutNA$InvertNegsMutator and randoopResults123WithoutNA$InvertNegsMutator
## V = 5, p-value = 0.5
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(evoResults123WithoutNA$InvertNegsMutator, manualResults123WithoutNA$InvertNegsMutator, paired = TRUE)
```

```
##
## Wilcoxon signed rank test
##
## data:  evoResults123WithoutNA$InvertNegsMutator and manualResults123WithoutNA$InvertNegsMutator
## V = 3, p-value = 1
## alternative hypothesis: true location shift is not equal to 0
```

```
wilcox.test(randoopResults123WithoutNA$InvertNegsMutator, manualResults123WithoutNA$InvertNegsMutator, paired = TRUE)
```



```
##  
## Wilcoxon signed rank test  
##  
## data:  randoopResults123WithoutNA$InvertNegsMutator and manualResults123WithoutNA  
$InvertNegsMutator  
## V = 0, p-value = 0.25  
## alternative hypothesis: true location shift is not equal to 0
```