# Survey Analysis

## (1) Load the datasets

```
ds <- read.csv("second-survey-js.csv", head=T, sep=",")</pre>
atom ds <- read.csv("s02-atoms.csv", head=T, sep=",")
nrow(ds)
## [1] 1839
colnames(ds)
## [1] "rid"
                      "sid"
                                    "qid"
                                                  "atom"
                                                               "time"
## [6] "correct"
                      "experience" "education"
                                                 "total"
                                                               "ref"
## [11] "empty"
                      "ssid"
nrow(atom_ds)
## [1] 48
colnames(atom_ds)
## [1] "atomId"
                          "atomDescription"
(2) Filter the datasets
# sqldf("select rid, count(*) from ds group by rid")
# sqldf("select ref, count(*) from ds group by ref")
# sqldf("select total, count(*) from ds group by total")
subSet <- ds[ds$ref=="reddit" & ds$total == "YES",]</pre>
summary(subSet$total)
##
      Length
                  Class
                             Mode
##
        1500 character character
squareLengths <- tapply(subSet$time,subSet$rid,length)</pre>
completeCases <- names(squareLengths)[squareLengths==24]</pre>
ds <- ds[is.element(el = ds$rid, set = completeCases),]</pre>
dim(ds)
## [1] 816 12
sqldf("select rid, count(*) from ds group by rid")
      rid count(*)
## 1
                24
       1
## 2
       13
## 3
                24
       16
## 4
       18
                 24
## 5
                 24
       19
```

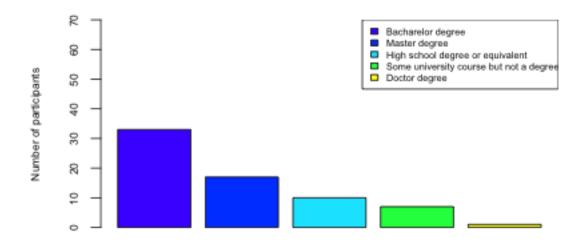
```
24
## 6
                24
## 7
       25
## 8
       28
                24
## 9
       35
                24
## 10 47
                24
## 11 48
                24
## 12 51
## 13 55
                24
## 14 57
                24
## 15 59
                24
## 16 60
                24
                24
## 17
      61
                24
## 18 63
## 19 65
                24
## 20 79
                24
## 21 81
                24
## 22 87
                24
## 23 89
                24
## 24 91
                24
                24
## 25 95
## 26 97
                24
## 27 98
                24
## 28 103
                24
## 29 104
                24
                24
## 30 105
## 31 112
                24
## 32 113
                24
## 33 115
                24
## 34 117
                24
```

## (3) Exploratory Data Analysis

#### Demographics

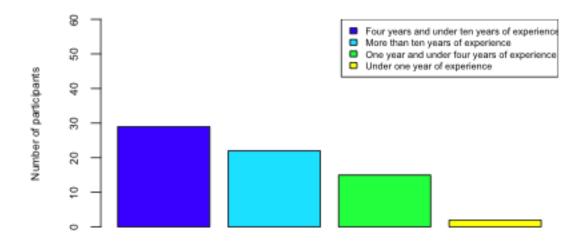
#### Education

```
##
                                 description total
## 1
                           Bacharelor degree
## 2
                               Master degree
                                                 17
## 3
            High school degree or equivalent
                                                 10
## 4 Some university course but not a degree
                                                 7
## 5
                               Doctor degree
                                                 1
barplot(demEducation$total, col=topo.colors(5),
        ylim = c(0, 70), cex=0.7, cex.lab = 0.7, cex.axis=0.7,
        ylab="Number of participants")
legend("topright", legend=demEducation$description, fill=topo.colors(5), cex=0.6)
```



```
from demExperience a, experience_ds b
where a.experience = b.id
order by 3 desc")
demExperience
```

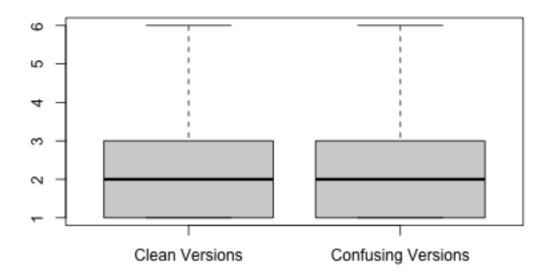
#### Experience



## Total number of correct answers (Table III)

```
codeWithAtoms["atomId"] = codeWithAtoms$qid %% 24
codeWithoutAtoms["atomId"] = codeWithoutAtoms$qid %% 24
merged <- sqldf("select c.atomDescription as Atom,
                       a.confuseCode as 'Confusing Versions',
                       b.cleanCode as 'Clean Versions',
                       (b.cleanCode * 100 / a.confuseCode) -100 as 'Delta (%)'
                from codeWithAtoms a, codeWithoutAtoms b, atom ds c
                where a.atomId = b.atomId and a.atomId = c.atomId
                order by 4 desc")
xtable(merged)
## % latex table generated in R 4.2.0 by xtable 1.8-4 package
## % Wed Oct 5 10:35:23 2022
## \begin{table}[ht]
## \centering
## \begin{tabular}{rlrrr}
##
    \hline
## & Atom & Confusing Versions & Clean Versions & Delta (\%) \
##
    \hline
## 1 & Type Conversion &
                          2 &
                                6 & 200 \\
##
    2 & Change Literal Encoding &
                                    3 &
                                        7 & 133 \\
##
    3 & Comma Operator &
                           7 & 12 & 71 \\
##
    4 & Arithmetic As Logic &
                                3 & 5 & 66 \\
##
    5 & Indentation No Braces & 4 & 6 & 50 \\
##
    6 & Assignment As Value &
                                5 & 7 & 40 \\
##
    7 & Repurposed Variables &
                               3 &
                                     4 & 33 \\
##
    8 & Post Increment & 13 & 15 & 15 \\
##
    9 & Arrow Function &
                          7 &
                                8 & 14 \\
##
    10 & Ommited Curly Braces & 18 & 20 & 11 \\
    11 & Array Destructuring & 18 & 20 & 11 \\
##
##
    12 & Logic As Control Flow & 21 & 23 & 9 \\
    13 & Indentation With Braces & 19 & 20 & 5 \
##
##
    14 & Infix Operator Precedence &
                                       6 % 6 % 0 \\
##
    15 & Ternary Operator & 23 & 23 &
                                          0 \\
    16 & Constant Variables & 22 & 21 & -5 \\
##
    17 & Dead Unreachable Repeated & 23 & 22 & -5 \\
##
    18 & Property Access & 22 & 21 & -5 \\
##
    19 & Array Spread & 25 & 21 & -16 \\
##
##
    20 & Implicit Predicate & 21 & 15 & -29 \\
    21 & Automatic Semicolon Insertion & 2 &
##
                                                1 & -50 \\
##
    22 & Object Spread &
                          5 &
                               2 & -60 \\
     \hline
##
## \end{tabular}
## \end{table}
wrongAnswers = ds[ds$correct == 'WRONG', ]
wrongAnswersByStudentTreatment <- aggregate(rid~sid+atom,</pre>
                                           data = wrongAnswers,
                                           FUN=length)
boxplot(wrongAnswersByStudentTreatment$rid ~ wrongAnswersByStudentTreatment$atom
```

```
, ylab = "", xlab = "", main = "",
names = c("Clean Versions", "Confusing Versions"))
```

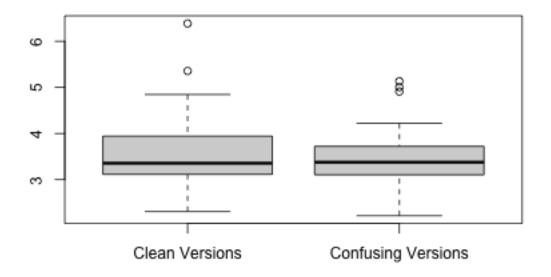


#### Average time for correct answers (Table IV)

```
merged <- sqldf("select c.atomDescription as Atom,</pre>
                        a.timeConfuseCode as 'Confuse Code',
                        b.timeCleanCode as 'Clean Code',
                        (b.timeCleanCode * 100 / a.timeConfuseCode) -100 as 'Delta (%)'
                 from codeWithAtoms a, codeWithoutAtoms b, atom_ds c
                 where a.atomId = b.atomId and a.atomId = c.atomId
                 order by 4")
xtable(merged)
## % latex table generated in R 4.2.0 by xtable 1.8-4 package
## % Wed Oct 5 10:35:24 2022
## \begin{table}[ht]
## \centering
## \begin{tabular}{rlrrr}
     \hline
##
  & Atom & Confuse Code & Clean Code & Delta (\%) \\
##
##
     \hline
## 1 & Change Literal Encoding & 82.30 & 13.36 & -83.77 \\
     2 & Infix Operator Precedence & 43.98 & 24.24 & -44.89 \\
##
##
     3 & Comma Operator & 64.03 & 35.84 & -44.03 \
##
     4 & Constant Variables & 24.59 & 13.98 & -43.14 \\
##
    5 & Type Conversion & 50.20 & 29.76 & -40.72 \
##
    6 & Object Spread & 66.40 & 43.85 & -33.96 \\
    7 & Indentation With Braces & 36.39 & 28.74 & -21.02 \\
```

```
8 & Automatic Semicolon Insertion & 63.78 & 55.77 & -12.57 \\
##
##
     9 & Dead Unreachable Repeated & 18.50 & 17.08 & -7.66 \\
     10 & Arrow Function & 43.55 & 40.44 & -7.15 \\
##
     11 & Repurposed Variables & 69.92 & 66.76 & -4.52 \
##
##
     12 & Ommited Curly Braces & 25.77 & 26.07 & 1.15 \\
##
     13 & Assignment As Value & 41.67 & 45.88 & 10.11 \\
##
     14 & Arithmetic As Logic & 28.57 & 33.36 & 16.76 \\
     15 & Array Destructuring & 22.48 & 28.68 & 27.59 \\
##
##
     16 & Post Increment & 40.90 & 54.84 & 34.09 \\
##
     17 & Array Spread & 39.02 & 52.47 & 34.49 \\
     18 & Logic As Control Flow & 37.26 & 50.84 & 36.43 \\
##
     19 & Indentation No Braces & 23.01 & 40.40 & 75.61 \\
     20 & Property Access & 34.64 & 66.65 & 92.40 \\
##
     21 & Ternary Operator & 27.39 & 87.85 & 220.69 \\
##
     22 & Implicit Predicate & 30.12 & 252.82 & 739.29 \\
##
##
      \hline
## \end{tabular}
## \end{table}
correctAnswers = ds[ds$correct == 'CORRECT', ]
correctAnswersByStudentTreatment <- aggregate(time~sid+atom,</pre>
                                            data = correctAnswers,
                                            FUN=mean)
boxplot(log(correctAnswersByStudentTreatment$time) ~ correctAnswersByStudentTreatment$atom
        , ylab = "", xlab = "", main = "Number of wrong answers of each participant",
        names = c("Clean Versions", "Confusing Versions"))
```

## Number of wrong answers of each participant



```
prop.table(table(ds$correct,ds$atom),margin = 2)
##
##
                    NO
                             YES
##
     CORRECT 0.7181373 0.6740196
##
     WRONG
           0.2818627 0.3259804
chisq.test(ds$correct,ds$atom)
##
  Pearson's Chi-squared test with Yates' continuity correction
##
## data: ds$correct and ds$atom
## X-squared = 1.6741, df = 1, p-value = 0.1957
(4) Regression Analysis (correctness-atom + experience + education)
experience <- as.factor(ds$experience)</pre>
education <- as.factor(ds$education)</pre>
contrr < -matrix(c(rep(1,4),c(1/2,1/2,-1/2,-1/2),c(1,-1,0,0),c(0,0,1,-1)), byrow=TRUE, nrow=4)
contrasts(experience)<-solve(contrr)[,2:4]</pre>
ds$atom <- as.factor(ds$atom)</pre>
ds$correct <- as.factor(ds$correct)</pre>
mod <- glm(ds$correct~ds$atom+experience+education,family = "binomial")</pre>
summary(mod)
##
## Call:
## glm(formula = ds$correct ~ ds$atom + experience + education,
##
       family = "binomial")
##
## Deviance Residuals:
                     Median
##
       Min
                 10
                                   30
                                           Max
## -1.3941 -0.8035 -0.7149 1.1472
                                        1.9437
##
## Coefficients:
##
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.01341   0.21274 -0.063   0.9497
              0.22303
## ds$atomYES
                           0.15768
                                     1.414
                                             0.1572
## experience1 0.04762
                           0.26637
                                     0.179
                                             0.8581
                           0.48657 -1.080
## experience2 -0.52572
                                             0.2799
## experience3 -0.22857
                           0.18971 - 1.205
                                             0.2283
                           0.30118 -1.481
## education2 -0.44600
                                             0.1386
                           0.22200 -5.578 2.44e-08 ***
## education3 -1.23823
## education4 -1.30554
                           0.25864 -5.048 4.47e-07 ***
## education5 -1.80216
                           0.80437 -2.240
                                            0.0251 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 1002.29 on 815 degrees of freedom
```

```
## Residual deviance: 950.84 on 807 degrees of freedom
## ATC: 968.84
##
## Number of Fisher Scoring iterations: 4
car::Anova(mod, type=3)
## Analysis of Deviance Table (Type III tests)
##
## Response: ds$correct
##
            LR Chisq Df Pr(>Chisq)
## ds$atom
              2.006 1
                            0.1567
## experience 4.657 3
                            0.1987
## education 42.510 4 1.308e-08 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

## Impact considering Developer Experience

```
for(i in 1:4){
   print(paste("Experience ", i ))
   print(prop.table(table(ds$correct[ds$experience==i],ds$atom[ds$experience==i]),margin=2))
   print(chisq.test(table(ds$correct[ds$experience==i],ds$atom[ds$experience==i])))
}
```

## Impact considering Developer Education

```
for(i in 1:5){
  print(paste("Education ", i ))
  print(prop.table(table(ds$correct[ds$education==i],ds$atom[ds$education==i]),margin=2))
  print(chisq.test(table(ds$correct[ds$education==i],ds$atom[ds$education==i])))
}
```

#### Impact of Individual Atoms and Hypotheses Testing

```
chiTest <- c()</pre>
oddsRatio <- c()
oddsRatioTest <- c()
ci25 <- c()
ci975 <- c()
mannWhitneyTest <- c()</pre>
cliffDelta <- c()</pre>
for(i in 1:10){
  subSet \leftarrow ds[is.element(el = ds$qid, set = c(i,i+10)),]
  print(atomDescription[i])
  tableCorrectness <- table(subSet$correct,subSet$atom)</pre>
  tableTime <- aggregate(ds$time, by=list(atom=ds$atom), FUN=sum)</pre>
  print(tableCorrectness)
  print(tableTime)
  test <- chisq.test(tableCorrectness)</pre>
  print(test)
```

```
chiTest[i] <- format.pval(test$p.value)</pre>
  oddsRatio[i] <- odds.ratio(tableCorrectness)$OR</pre>
  oddsRatioTest[i] <- format.pval(odds.ratio(tableCorrectness)$p)</pre>
  ci25[i] <- odds.ratio(tableCorrectness, level=0.95)$"2.5 %"
  ci975[i] <- odds.ratio(tableCorrectness, level=0.95)$"97.5 %"</pre>
  mannWhitneyTest[i] <- format.pval(wilcox.test(subSet$time~as.factor(subSet$atom))$p.value)</pre>
  cliffDelta[i] <- cliff.delta(subSet$time~as.factor(subSet$atom))$estimate</pre>
  experience <- as.factor(subSet$experience)</pre>
  contrr < -matrix(c(rep(1,4),c(1/2,1/2,-1/2,-1/2),c(1,-1,0,0),c(0,0,1,-1)), byrow=TRUE, nrow=4)
  contrasts(experience)<-solve(contrr)[,2:4]</pre>
  mod <- glm(subSet$correct ~ subSet$atom + experience,family="binomial")</pre>
  print(summary(mod))
  print(prop.table(table(subSet$correct,subSet$experience),margin = 2))
analysis_df <- data.frame(atomDescription, chiTest, oddsRatio,</pre>
                            oddsRatioTest, ci25, ci975, mannWhitneyTest, cliffDelta)
analysis_df <- analysis_df[order(-oddsRatio),]</pre>
colnames(analysis_df) <- c("Atom", "ChiTest",</pre>
                             "Odds Ratio Correctness",
                             "p-value", "CI 2.5%", "CI 97.5%",
                             "Wilcox Test (Time)",
                             "Cliff Delta")
xtable(analysis_df)
```