

# Shooting justification algorithm

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```
#importing dataset into R
setwd("D:/Google Drive/RESEARCH/Older projects/APLS - Short form/Study 2")
library(readxl)
S2 <- read_excel("S2.xlsx")
```

```
## New names:
## • `` -> `...22`
```

*#Filtering out participants who did not answer the justification question*

```
library(dplyr)
```

```
## Warning: package 'dplyr' was built under R version 4.2.3
```

```
##
## Attaching package: 'dplyr'
##
## The following objects are masked from 'package:stats':
## 
##     filter, lag
## 
## The following objects are masked from 'package:base':
## 
##     intersect, setdiff, setequal, union
```

```

S2_filtered <- S2[complete.cases(S2$justif), ]
S <- S2_filtered

#####
#recoding the variables
S <- S %>% mutate(edc=recode(edc, `1` = "Some High School", `2` = "GED",
                                '3' = "High School Diploma", '4'="Associates Degree",
                                '5' = "Some college", '6' = "Bachelor Degree",
                                '7' = "Masters Degree", '8' = "Doctoral Degree"))

S <- S %>% mutate(income=recode(income, `1` = "Less than $10,000", `2` = "$10,001 to $20,000",
                                '3' = "$20,001 to $30,000", '4'="$30,001 to $40,000",
                                '5' = "$40,001 to $50,000", '6' = "$50,001 to $60,000",
                                '7' = "$60,001 to $70,000", '8' = "$70,001 to $80,000",
                                '9' = "$80,001 to $90,000", '10' = "$90,001 to $100,000",
                                '11' = "More than $100,000"))

S <- S %>% mutate(race.recoded=recode(race.recoded, "0" = "White",
                                         "2" = "Asian",
                                         "3"="Black",
                                         "4" = "Hispanic",
                                         "5" = "Middle_Eastern",
                                         "7" = "Other",
                                         "8" = "Multi"))

S <- S %>% mutate(sex=recode(sex, `1` = "Male", `2` = "Female"))

S <- S %>% mutate(justif=recode(justif, `1` = "Justified", `0` = "Not_justified"))

S<-rename(S, APLS_SF.Rasch = APLS.R)

#factoring variables

S$sex <- factor(S$sex)

S$justif <- factor(S$justif)

S$justif<-relevel(S$justif, ref = "Not_justified")

S$race.recoded<-factor(S$race.recoded)

S$income <- factor(S$income, ordered = TRUE)

library(forcats)
S$income <- fct_relevel(S$income, "Less than $10,000", "$10,001 to $20,000", "$20,001 to $30,000",
                        "$30,001 to $40,000", "$40,001 to $50,000", "$50,001 to $60,000",
                        "$60,001 to $70,000", "$70,001 to $80,000", "$80,001 to $90,000",
                        "$90,001 to $100,000", "More than $100,000")

```

```

"$60,001 to $70,000", "$70,001 to $80,000", "$80,001 to $90,000",
"$90,001 to $100,000", "More than $100,000")

S$edc<-factor(S$edc, ordered = TRUE)

S$edc<-fct_relevel(S$edc, "Some High School", "GED", "High School Diploma", "Some college", "Associates Degree",
                     "Bachelor Degree", "Masters Degree", "Doctoral Degree")

#creating the training and test sets
set.seed(1234)

S <- S %>% select(age, edc, income, sex, race.recoded, SD0.avg, RWA.avg, APLS_SF.Rasch,
                  apls1, apls2, apls3, apls4, apls5, apls6, apls7, apls8, apls9,
                  apls10, apls11,
                  SD01, SD02, SD03, SD04, SD05, SD06, SD07, SD08, SD09, SD010,
                  SD011, SD012, SD013, SD014, SD015, SD016,
                  RWA1, RWA3, RWA4, RWA5, RWA6, RWA7, RWA8, RWA9,
                  RWA10, RWA11, RWA12, RWA13, RWA14, RWA15, justif)

sample_set <- sample(nrow(S), round(nrow(S)*.8), replace = FALSE)

S_train <- S[sample_set, ]

S_test <- S[-sample_set, ]

```

```

#Creating the decision tree model for the training set
library(caret)

```

```
## Warning: package 'caret' was built under R version 4.2.3
```

```
## Loading required package: ggplot2
```

```
## Loading required package: lattice
```

```

mod1<- train(justif ~ .,
              data=S_train,
              metric="Kappa",
              method= "rpart",
              trControl= trainControl(method="cv", number=3),
              tuneLength=20
)
mod1

```

```
## CART
##
## 255 samples
## 49 predictor
## 2 classes: 'Not_justified', 'Justified'
##
## No pre-processing
## Resampling: Cross-Validated (3 fold)
## Summary of sample sizes: 171, 170, 169
## Resampling results across tuning parameters:
##
##     cp          Accuracy   Kappa
## 0.00000000  0.7214058  0.3416466
## 0.01199201  0.7214058  0.3416466
## 0.02398401  0.7214058  0.3416466
## 0.03597602  0.7214058  0.3416466
## 0.04796802  0.7488568  0.3729921
## 0.05996003  0.7567933  0.4067084
## 0.07195203  0.7567933  0.4067084
## 0.08394404  0.7448885  0.3949031
## 0.09593604  0.7448885  0.3949031
## 0.10792805  0.7369998  0.3860990
## 0.11992005  0.7331238  0.4272221
## 0.13191206  0.7331238  0.4272221
## 0.14390406  0.7331238  0.4272221
## 0.15589607  0.7331238  0.4272221
## 0.16788807  0.7331238  0.4272221
## 0.17988008  0.7098680  0.2671760
## 0.19187209  0.7098680  0.2671760
## 0.20386409  0.7098680  0.2671760
## 0.21585610  0.7098680  0.2671760
## 0.22784810  0.7098680  0.2671760
##
## Kappa was used to select the optimal model using the largest value.
## The final value used for the model was cp = 0.1678881.
```

```
summary(mod1)
```

```

## Call:
## (function (formula, data, weights, subset, na.action = na.rpart,
##   method, model = FALSE, x = FALSE, y = TRUE, parms, control,
##   cost, ...)
## {
##   Call <- match.call()
##   if (is.data.frame(model)) {
##     m <- model
##     model <- FALSE
##   }
##   else {
##     indx <- match(c("formula", "data", "weights", "subset"),
##       names(Call), nomatch = 0)
##     if (indx[1] == 0)
##       stop("a 'formula' argument is required")
##     temp <- Call[c(1, indx)]
##     temp$na.action <- na.action
##     temp[[1]] <- quote(stats::model.frame)
##     m <- eval.parent(temp)
##   }
##   Terms <- attr(m, "terms")
##   if (any(attr(Terms, "order") > 1))
##     stop("Trees cannot handle interaction terms")
##   Y <- model.response(m)
##   wt <- model.weights(m)
##   if (any(wt < 0))
##     stop("negative weights not allowed")
##   if (!length(wt))
##     wt <- rep(1, nrow(m))
##   offset <- model.offset(m)
##   X <- rpart.matrix(m)
##   nobs <- nrow(X)
##   nvar <- ncol(X)
##   if (missing(method)) {
##     method <- if (is.factor(Y) || is.character(Y))
##       "class"
##     else if (inherits(Y, "Surv"))
##       "exp"
##     else if (is.matrix(Y))
##       "poisson"
##     else "anova"
##   }
##   if (is.list(method)) {
##     mlist <- method
##     method <- "user"
##     init <- if (missing(parms))
##       mlist$init(Y, offset, wt = wt)
##     else mlist$init(Y, offset, parms, wt)
##     keep <- rpartcallback(mlist, nobs, init)
##     method.int <- 4
##     parms <- init$parms
##   }

```

```

##      else {
##          method.int <- pmatch(method, c("anova", "poisson", "class",
##                                         "exp"))
##          if (is.na(method.int))
##              stop("Invalid method")
##          method <- c("anova", "poisson", "class", "exp")[method.int]
##          if (method.int == 4)
##              method.int <- 2
##          init <- if (missing(parms))
##                  get(paste("rpart", method, sep = "."), envir = environment())(Y,
##                                         offset, , wt)
##          else get(paste("rpart", method, sep = "."), envir = environment())(Y,
##                                         offset, parms, wt)
##          ns <- asNamespace("rpart")
##          if (!is.null(init$print))
##              environment(init$print) <- ns
##          if (!is.null(init$summary))
##              environment(init$summary) <- ns
##          if (!is.null(init$text))
##              environment(init$text) <- ns
##      }
##      Y <- init$y
##      xlevels <- .getXlevels(Terms, m)
##      cats <- rep(0, ncol(X))
##      if (!is.null(xlevels)) {
##          xlevels <- xlevels[names(xlevels) %in% colnames(X)]
##          cats[match(names(xlevels), colnames(X))] <- unlist(lapply(xlevels,
##                                         length))
##      }
##      extraArgs <- list(...)
##      if (length(extraArgs)) {
##          controlargs <- names(formals(rpart.control))
##          indx <- match(names(extraArgs), controlargs, nomatch = 0)
##          if (any(indx == 0))
##              stop(gettextf("Argument %s not matched", names(extraArgs)[indx ==
##                                         0]), domain = NA)
##      }
##      controls <- rpart.control(...)
##      if (!missing(control))
##          controls[names(control)] <- control
##      xval <- controls$xval
##      if (is.null(xval) || (length(xval) == 1 && xval == 0) ||
##          method == "user") {
##          xgroups <- 0
##          xval <- 0
##      }
##      else if (length(xval) == 1) {
##          xgroups <- sample(rep(1:xval, length.out = nobs), nobs,
##                            replace = FALSE)
##      }
##      else if (length(xval) == nobs) {
##          xgroups <- xval

```

```

##      xval <- length(unique(xgroups))
##    }
##  else {
##    if (!is.null(attr(m, "na.action"))) {
##      temp <- as.integer(attr(m, "na.action"))
##      xval <- xval[-temp]
##      if (length(xval) == nobs) {
##        xgroups <- xval
##        xval <- length(unique(xgroups))
##      }
##      else stop("Wrong length for 'xval'")
##    }
##    else stop("Wrong length for 'xval'")
##  }
##  if (missing(cost))
##    cost <- rep(1, nvar)
##  else {
##    if (length(cost) != nvar)
##      stop("Cost vector is the wrong length")
##    if (any(cost <= 0))
##      stop("Cost vector must be positive")
##  }
##  tfun <- function(x) if (is.matrix(x))
##    rep(is.ordered(x), ncol(x))
##  else is.ordered(x)
##  labs <- sub("^`(.*)`$", "\\\1", attr(Terms, "term.labels"))
##  isord <- unlist(lapply(m[labs], tfun))
##  storage.mode(X) <- "double"
##  storage.mode(wt) <- "double"
##  temp <- as.double(unlist(init$parms))
##  if (!length(temp))
##    temp <- 0
##  rpf <- .Call(C_rpart, ncat = as.integer(cats * !isord),
##    method = as.integer(method.int), as.double(unlist(controls)),
##    temp, as.integer(xval), as.integer(xgroups), as.double(t(init$y)),
##    X, wt, as.integer(init$numy), as.double(cost))
##  nsplit <- nrow(rpf$isplit)
##  ncat <- if (!is.null(rpf$csplit))
##    nrow(rpf$csplit)
##  else 0
##  if (nsplit == 0)
##    xval <- 0
##  numcp <- ncol(rpf$cptable)
##  temp <- if (nrow(rpf$cptable) == 3)
##    c("CP", "nsplit", "rel error")
##  else c("CP", "nsplit", "rel error", "xerror", "xstd")
##  dimnames(rpf$cptable) <- list(temp, 1:numcp)
##  tname <- c("<leaf>", colnames(X))
##  splits <- matrix(c(rpf$isplit[, 2:3], rpf$dsplit), ncol = 5,
##    dimnames = list(tname[rpf$isplit[, 1] + 1], c("count",
##      "ncat", "improve", "index", "adj")))
##  index <- rpf$inode[, 2]

```

```

##      nadd <- sum(isord[rpfit$isplit[, 1]])
##      if (nadd > 0) {
##          newc <- matrix(0, nadd, max(cats))
##          cvar <- rpfit$isplit[, 1]
##          indx <- isord[cvar]
##          cdir <- splits[indx, 2]
##          ccut <- floor(splits[indx, 4])
##          splits[indx, 2] <- cats[cvar[indx]]
##          splits[indx, 4] <- ncat + 1:nadd
##          for (i in 1:nadd) {
##              newc[i, 1:(cats[(cvar[indx])[i]])] <- -as.integer(cdir[i])
##              newc[i, 1:ccut[i]] <- as.integer(cdir[i])
##          }
##          catmat <- if (ncat == 0)
##              newc
##          else {
##              cs <- rpfit$csplit
##              ncs <- ncol(cs)
##              ncc <- ncol(newc)
##              if (ncs < ncc)
##                  cs <- cbind(cs, matrix(0, nrow(cs), ncc - ncs))
##              rbind(cs, newc)
##          }
##          ncat <- ncat + nadd
##      }
##      else catmat <- rpfit$csplit
##      if (nsplit == 0) {
##          frame <- data.frame(row.names = 1, var = "<leaf>", n = rpfit$inode[, 5],
##          wt = rpfit$dnode[, 3], dev = rpfit$dnode[, 1],
##          yval = rpfit$dnode[, 4], complexity = rpfit$dnode[, 2],
##          ncompete = 0, nsurrogate = 0)
##      }
##      else {
##          temp <- ifelse(index == 0, 1, index)
##          svar <- ifelse(index == 0, 0, rpfit$isplit[temp, 1])
##          frame <- data.frame(row.names = rpfit$inode[, 1], var = tname[svar + 1],
##          n = rpfit$inode[, 5], wt = rpfit$dnode[, 3],
##          dev = rpfit$dnode[, 1], yval = rpfit$dnode[, 4],
##          complexity = rpfit$dnode[, 2], ncompete = pmax(0,
##          rpfit$inode[, 3] - 1), nsurrogate = rpfit$inode[, 4])
##      }
##      if (method.int == 3) {
##          numclass <- init$numresp - 2
##          nodeprob <- rpfit$dnode[, numclass + 5]/sum(wt)
##          temp <- pmax(1, init$counts)
##          temp <- rpfit$dnode[, 4 + (1:numclass)] %*% diag(init$parms$prior/temp)
##          yprob <- temp/rowSums(temp)
##          yval2 <- matrix(rpfit$dnode[, 4 + (0:numclass)], ncol = numclass + 1)
##          frame$yval2 <- cbind(yval2, yprob, nodeprob)
##      }

```

```

##      else if (init$numresp > 1)
##          frame$yval2 <- rpfit$dnode[, -(1:3), drop = FALSE]
##      if (is.null(init$summary))
##          stop("Initialization routine is missing the 'summary' function")
##      functions <- if (is.null(init$print))
##          list(summary = init$summary)
##      else list(summary = init$summary, print = init$print)
##      if (!is.null(init$text))
##          functions <- c(functions, list(text = init$text))
##      if (method == "user")
##          functions <- c(functions, mlist)
##      where <- rpfit$which
##      names(where) <- row.names(m)
##      ans <- list(frame = frame, where = where, call = Call, terms = Terms,
##                  cptable = t(rpfit$cptable), method = method, parms = init$parms,
##                  control = controls, functions = functions, numresp = init$numresp)
##      if (nsplit)
##          ans$splits = splits
##      if (ncat > 0)
##          ans$csplit <- catmat + 2
##      if (nsplit)
##          ans$variable.importance <- importance(ans)
##      if (model) {
##          ans$model <- m
##          if (missing(y))
##              y <- FALSE
##      }
##      if (y)
##          ans$y <- Y
##      if (x) {
##          ans$x <- X
##          ans$wt <- wt
##      }
##      ans$ordered <- isord
##      if (!is.null(attr(m, "na.action")))
##          ans$na.action <- attr(m, "na.action")
##      if (!is.null(xlevels))
##          attr(ans, "xlevels") <- xlevels
##      if (method == "class")
##          attr(ans, "ylevels") <- init$ylevels
##      class(ans) <- "rpart"
##      ans
## })(formula = .outcome ~ ., data = list(c(44, 52, 26, 27, 36,
## ## 33, 37, 36, 59, 26, 36, 38, 33, 62, 34, 26, 57, 28, 53, 40, 23,
## ## 26, 19, 59, 50, 21, 22, 39, 20, 23, 52, 20, 44, 68, 26, 31, 26,
## ## 46, 36, 35, 36, 52, 43, 43, 40, 63, 35, 19, 20, 31, 27, 30, 44,
## ## 22, 51, 26, 39, 30, 34, 41, 36, 40, 48, 44, 57, 30, 54, 58, 20,
## ## 35, 54, 46, 41, 25, 48, 35, 23, 27, 36, 28, 33, 26, 36, 32, 30,
## ## 58, 25, 23, 41, 31, 21, 37, 25, 25, 21, 24, 49, 39, 25, 23, 73,
## ## 26, 39, 19, 44, 30, 33, 57, 38, 53, 27, 31, 34, 32, 40, 49, 49,
## ## 27, 26, 40, 53, 29, 71, 66, 64, 35, 26, 73, 55, 39, 48, 20, 36,
## ## 19, 27, 34, 58, 50, 36, 24, 56, 29, 28, 41, 52, 19, 26, 55, 30,

```

```

## 40, 53, 26, 44, 54, 39, 70, 44, 38, 40, 22, 50, 36, 27, 50, 32,
## 40, 42, 25, 28, 29, 26, 24, 48, 43, 32, 21, 31, 44, 27, 61, 20,
## 30, 30, 23, 57, 35, 42, 27, 64, 46, 42, 39, 24, 61, 32, 33, 39,
## 31, 36, 47, 29, 26, 21, 25, 29, 32, 30, 41, 24, 41, 31, 59, 22,
## 41, 49, 33, 34, 28, 59, 34, 29, 55, 66, 34, 33, 20, 27, 62, 40,
## 22, 40, 54, 47, 20, 39, 60, 25, 45, 22, 19, 25, 42, 32, 74, 23,
## 20, 37, 36, 41, 51, 39, 30, 36, 55, 35), c(0.231455024943138,
## -0.231455024943138, 0.231455024943138, -0.231455024943138, 0.077151674981046,
## -0.231455024943138, -0.38575837490523, -0.0771516749810459, 0.077151674981046,
## -0.540061724867322, 0.540061724867322, -0.0771516749810459, 0.077151674981046,
## 0.231455024943138, -0.0771516749810459, 0.231455024943138, 0.540061724867322,
## -0.0771516749810459, -0.231455024943138, 0.231455024943138, 0.077151674981046,
## -0.540061724867322, -0.231455024943138, 0.231455024943138, -0.0771516749810459,
## 0.231455024943138, 0.231455024943138, -0.231455024943138, -0.0771516749810459,
## 0.077151674981046, 0.231455024943138, -0.0771516749810459, 0.38575837490523,
## 0.077151674981046, 0.231455024943138, 0.231455024943138, 0.231455024943138,
## 0.231455024943138, 0.38575837490523, -0.0771516749810459, 0.077151674981046,
## -0.231455024943138, -0.0771516749810459, -0.231455024943138,
## -0.0771516749810459, 0.231455024943138, -0.0771516749810459,
## -0.231455024943138, -0.0771516749810459, 0.38575837490523, 0.38575837490523,
## -0.0771516749810459, -0.0771516749810459, 0.231455024943138,
## -0.0771516749810459, 0.231455024943138, 0.231455024943138, 0.231455024943138,
## -0.38575837490523, 0.38575837490523, 0.077151674981046, -0.0771516749810459,
## -0.231455024943138, -0.0771516749810459, -0.231455024943138,
## 0.077151674981046, 0.231455024943138, 0.38575837490523, 0.231455024943138,
## 0.38575837490523, -0.231455024943138, -0.231455024943138, -0.0771516749810459,
## 0.231455024943138, 0.231455024943138, 0.077151674981046, 0.38575837490523,
## 0.231455024943138, 0.38575837490523, 0.231455024943138, 0.231455024943138,
## 0.231455024943138, -0.0771516749810459, 0.231455024943138, 0.231455024943138,
## 0.231455024943138, -0.0771516749810459, -0.0771516749810459,
## 0.38575837490523, 0.231455024943138, -0.0771516749810459, 0.231455024943138,
## -0.231455024943138, 0.077151674981046, 0.077151674981046, -0.540061724867322,
## 0.077151674981046, 0.231455024943138, -0.0771516749810459, 0.077151674981046,
## -0.231455024943138, 0.231455024943138, 0.231455024943138, 0.38575837490523,
## 0.38575837490523, 0.231455024943138, 0.231455024943138, 0.540061724867322,
## -0.231455024943138, -0.231455024943138, -0.0771516749810459,
## 0.38575837490523, 0.231455024943138, -0.231455024943138, -0.0771516749810459,
## 0.231455024943138, -0.0771516749810459, 0.077151674981046, -0.231455024943138,
## -0.0771516749810459, -0.231455024943138, 0.231455024943138, -0.0771516749810459,
## 0.231455024943138, 0.231455024943138, -0.0771516749810459, 0.077151674981046,
## 0.231455024943138, 0.38575837490523, 0.231455024943138, -0.0771516749810459,
## 0.231455024943138, 0.38575837490523, 0.231455024943138, 0.077151674981046,
## 0.077151674981046, 0.231455024943138, -0.0771516749810459, 0.231455024943138,
## -0.231455024943138, 0.231455024943138, -0.231455024943138, 0.38575837490523,
## 0.38575837490523, 0.231455024943138, 0.231455024943138, 0.077151674981046,
## -0.231455024943138, 0.231455024943138, -0.38575837490523, 0.38575837490523,
## 0.38575837490523, -0.231455024943138, 0.231455024943138, -0.0771516749810459,
## 0.231455024943138, 0.38575837490523, 0.231455024943138, 0.077151674981046,
## 0.077151674981046, 0.231455024943138, -0.0771516749810459, 0.231455024943138,
## -0.231455024943138, 0.231455024943138, -0.231455024943138, 0.38575837490523,
## 0.38575837490523, 0.231455024943138, 0.231455024943138, 0.231455024943138,
## -0.0771516749810459, -0.0771516749810459, 0.231455024943138,
```

```

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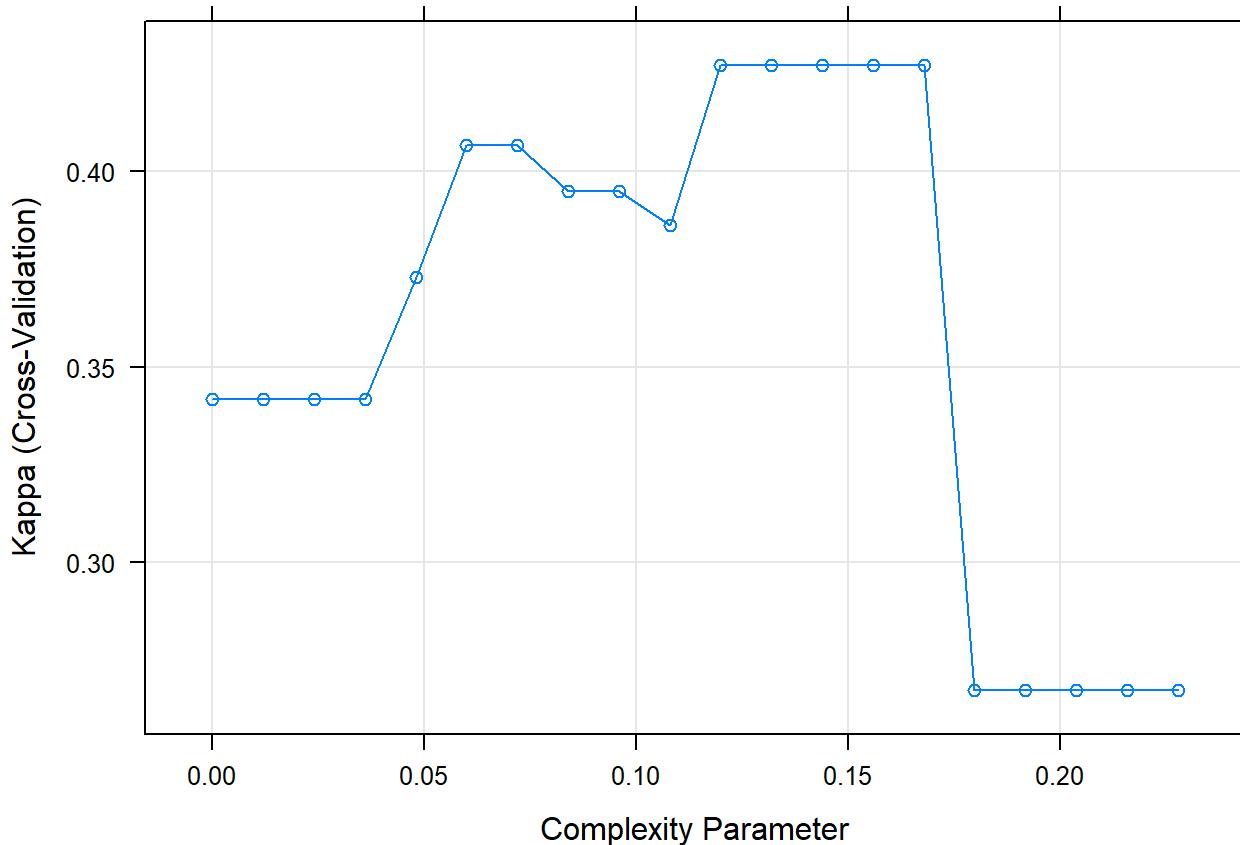
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## 1, 1, 2, 1, 1, 2, 1, 1, 1, 2, 1, 1, 1, 1, 1, 1, 1, 2, 1, 1, 2,
## 2, 1, 2, 2, 1, 1, 1, 1, 2, 2, 2, 1, 1, 1, 2, 2, 2, 1, 1, 1,
## 1, 1, 1, 1, 1, 1, 2, 1, 1, 1, 2, 2, 1, 1, 1, 2, 1, 1, 2, 2,
## 1, 2, 1, 1, 1, 2, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
## 1, 2, 2, 1, 2, 1, 1, 1, 1, 1, 1, 1, 1, 2, 1, 2, 1, 2, 2, 1, 1,
## 1, 1, 1, 2, 2, 1, 1, 2, 1, 1, 1, 2, 1, 1, 1, 2, 2, 1, 1, 1, 2,
## 1, 1, 1, 2, 1, 2, 1, 1, 1, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
## 1, 1, 2, 1, 2, 1, 1, 1, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
## 1, 1, 1, 2, 1, 2, 1, 1, 1, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
## 1, 1, 2, 1, 2, 1, 1, 1, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1),
## control = list(
##   20, 7, 0, 4, 5, 2, 0, 30, 0))
## n= 255
##
##          CP nsplit rel error
## 1 0.2278481      0 1.0000000
## 2 0.1678881      1 0.7721519
##
## Variable importance
##           apls2 APLS_SF.Rasch      apls5      apls4      apls3
##             24            19            17            14            14
##           apls1
##             13
##
## Node number 1: 255 observations,    complexity param=0.2278481
## predicted class=Not_justified  expected loss=0.3098039  P(node) =1
##   class counts:  176    79
##   probabilities: 0.690  0.310
## left son=2 (151 obs) right son=3 (104 obs)
## Primary splits:
##   apls2      < 3.5      to the left,  improve=26.90006, (0 missing)
##   APLS_SF.Rasch < 55.395 to the left,  improve=25.25502, (0 missing)
##   apls5      < 3.5      to the left,  improve=22.40698, (0 missing)
##   apls8      < 3.5      to the left,  improve=20.37007, (0 missing)
##   apls10     < 2.5      to the left,  improve=19.90602, (0 missing)
## Surrogate splits:
##   APLS_SF.Rasch < 55.395 to the left,  agree=0.914, adj=0.788, (0 split)
##   apls5      < 3.5      to the left,  agree=0.878, adj=0.702, (0 split)
##   apls4      < 3.5      to the left,  agree=0.843, adj=0.615, (0 split)
##   apls3      < 3.5      to the left,  agree=0.827, adj=0.577, (0 split)
##   apls1      < 3.5      to the left,  agree=0.824, adj=0.567, (0 split)
##
## Node number 2: 151 observations
## predicted class=Not_justified  expected loss=0.1192053  P(node) =0.5921569
##   class counts:  133    18

```

```
##   probabilities: 0.881 0.119
##
## Node number 3: 104 observations
##   predicted class=Justified      expected loss=0.4134615  P(node) =0.4078431
##   class counts:    43     61
##   probabilities: 0.413  0.587
```

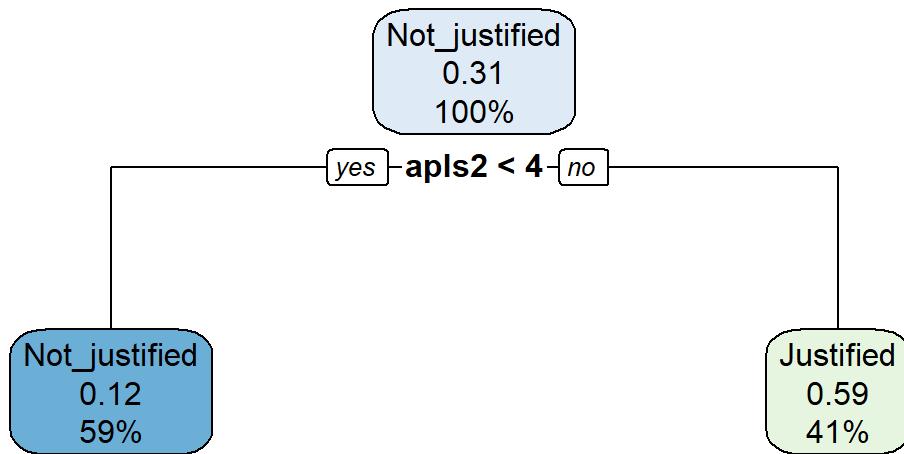
```
plot(mod1)
```



```
library(rpart.plot)
```

```
## Loading required package: rpart
```

```
rpart.plot(mod1$finalModel)
```



```
#building the final model in rpart based on the training results
library(rpart)
```

```
mod1.r <- rpart(justif ~ .,
                  data=S_train,
                  method = "class",
                  cp = .167,
                  maxdepth= 10,
                  minbucket = 10
      )
```

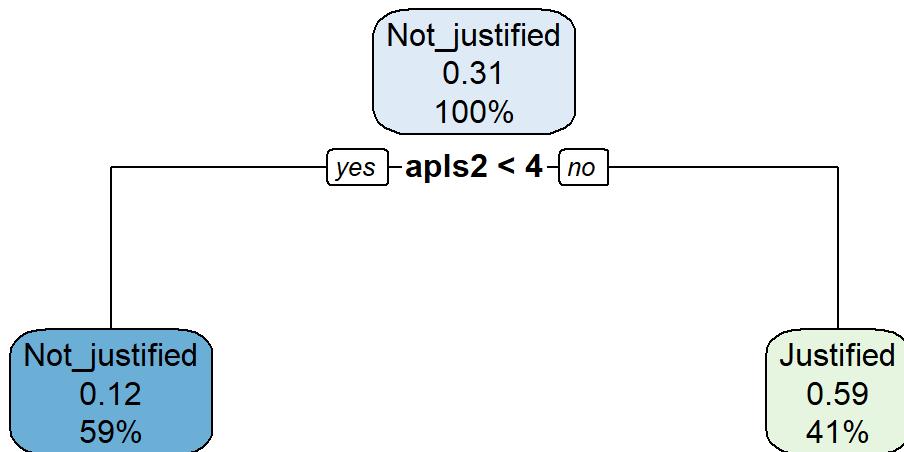
```
mod1.r
```

```
## n= 255
##
## node), split, n, loss, yval, (yprob)
##       * denotes terminal node
##
## 1) root 255 79 Not_justified (0.6901961 0.3098039)
##   2) apls2< 3.5 151 18 Not_justified (0.8807947 0.1192053) *
##   3) apls2>=3.5 104 43 Justified (0.4134615 0.5865385) *
```

```
summary(mod1.r)
```

```
## Call:
## rpart(formula = justif ~ ., data = S_train, method = "class",
##       cp = 0.167, maxdepth = 10, minbucket = 10)
## n= 255
##
##          CP nsplit rel error      xerror      xstd
## 1 0.2278481      0 1.0000000 1.0000000 0.09347010
## 2 0.1670000      1 0.7721519 0.8860759 0.09020647
##
## Variable importance
##          apls2 APLS_SF.Rasch      apls5      apls4      apls3
##            24           19           17           14           14
##          apls1
##            13
##
## Node number 1: 255 observations,    complexity param=0.2278481
## predicted class=Not_justified  expected loss=0.3098039  P(node) =1
##   class counts: 176 79
##   probabilities: 0.690 0.310
## left son=2 (151 obs) right son=3 (104 obs)
## Primary splits:
##   apls2 < 3.5 to the left,  improve=26.90006, (0 missing)
##   APLS_SF.Rasch < 55.395 to the left,  improve=25.25502, (0 missing)
##   apls5 < 3.5 to the left,  improve=22.40698, (0 missing)
##   apls8 < 3.5 to the left,  improve=20.37007, (0 missing)
##   apls10 < 2.5 to the left,  improve=19.90602, (0 missing)
## Surrogate splits:
##   APLS_SF.Rasch < 55.395 to the left,  agree=0.914, adj=0.788, (0 split)
##   apls5 < 3.5 to the left,  agree=0.878, adj=0.702, (0 split)
##   apls4 < 3.5 to the left,  agree=0.843, adj=0.615, (0 split)
##   apls3 < 3.5 to the left,  agree=0.827, adj=0.577, (0 split)
##   apls1 < 3.5 to the left,  agree=0.824, adj=0.567, (0 split)
##
## Node number 2: 151 observations
## predicted class=Not_justified  expected loss=0.1192053  P(node) =0.5921569
##   class counts: 133 18
##   probabilities: 0.881 0.119
##
## Node number 3: 104 observations
## predicted class=Justified    expected loss=0.4134615  P(node) =0.4078431
##   class counts: 43 61
##   probabilities: 0.413 0.587
```

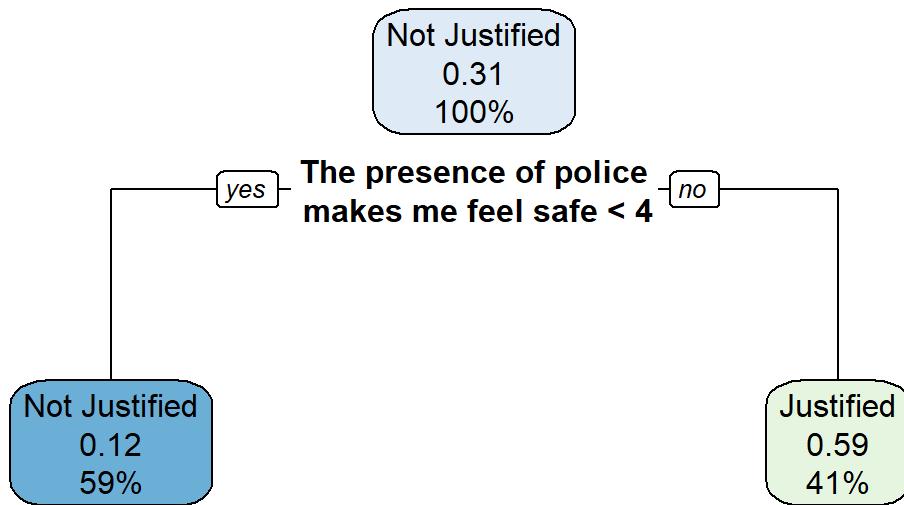
```
library(rpart)
library(rpart.plot)
rpart.plot(mod1.r)
```



```
split.fun <- function(x, labs, digits, varlen, faclen)
{
  labs <- gsub("apl$2",      "The presence of police\\n makes me feel safe", labs)
  labs
}

node.fun <- function(x, labs, digits, varlen)
{
  labs <- gsub("Not_justified", "Not Justified", labs)
  labs
}

rpart.plot(mod1.r, node.fun=node.fun, split.fun=split.fun, type = 2)
```



```

setwd("D:/Google Drive/RESEARCH/LAB STUFF/Officer involved shooting sudy") #setting working directory to folder in D drive
pdf(file="plot1.pdf")
rpart.plot(mod1.r, node.fun=node.fun, split.fun=split.fun, type = 2)
dev.off()
  
```

```

## png
## 2
  
```

```

setwd("D:/Google Drive/RESEARCH/LAB STUFF/Officer involved shooting sudy") #setting working directory to folder in D drive
png(file="plot1.png")
rpart.plot(mod1.r, node.fun=node.fun, split.fun=split.fun, type = 2)
dev.off()
  
```

```

## png
## 2
  
```

```

mod1.r.pred<-predict(mod1.r, newdata = S_test, type = "class")
confusionMatrix(data=mod1.r.pred, reference=S_test$justif, positive = "Justified", mode = "everything")
  
```

```

## Confusion Matrix and Statistics
##
##             Reference
## Prediction      Not_justified Justified
## Not_justified          36         7
## Justified              2        19
##
##                 Accuracy : 0.8594
##                 95% CI : (0.7498, 0.9336)
## No Information Rate : 0.5938
## P-Value [Acc > NIR] : 3.792e-06
##
##                 Kappa : 0.6994
##
## McNemar's Test P-Value : 0.1824
##
##                 Sensitivity : 0.7308
##                 Specificity : 0.9474
## Pos Pred Value : 0.9048
## Neg Pred Value : 0.8372
##                 Precision : 0.9048
##                 Recall : 0.7308
##                 F1 : 0.8085
##                 Prevalence : 0.4062
## Detection Rate : 0.2969
## Detection Prevalence : 0.3281
## Balanced Accuracy : 0.8391
##
## 'Positive' Class : Justified
##

```

```

# calculate area under curve
library(pROC)

```

```

## Type 'citation("pROC")' for a citation.

```

```

##
## Attaching package: 'pROC'

```

```

## The following objects are masked from 'package:stats':
##
##     cov, smooth, var

```

```

mod1.r.pred2<-predict(mod1.r, S_test, type = "prob")

mod1.r.rpart_ROC <- multiclass.roc(S_test$justif, mod1.r.pred2)

auc(mod1.r.rpart_ROC)

```

```
## Multi-class area under the curve: 0.8391
```

#####

#apls item 2 - The presence of police makes me feel safe.  
#the algorithm predicts that, on a scale from 1-5, those who answer the question, "The presence  
of police makes me feel safe" with 1-3 will perceive a shooting as Not justified.  
#for those who answer it 4 or 5, the algorithm predicts that they would view it as justified  
#Remarkably, using this single item, the algorithm achieves a balanced accuracy of .8391, an AUC  
of .8391, and a Kappa of .6994