College Data

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Packages:

```
# install the development version of vivid:
#devtools::install_github("AlanInglis/vivid")

# Load relevant packages:
library("vivid") # for visualisations
library("ISLR") # for data
library("mlr3") # to create model
library("mlr3learners") # to create model
library("randomForest") # to create model
library("condvis2") # for predict function
library("kknn") # to get knn model
library("dplyr")
```

Read in and setup data:

```
# Load data:
collegeData <- as.data.frame(College)

# Taking log values of skewed data:
collegeData <- collegeData %>%
   mutate(log(collegeData[,c(2:4,7:12)]))

# Split data into train and test
set.seed(101)
train <- sample(nrow(collegeData), round(.7*nrow(collegeData))) # split 70-30
collegeTrain <- collegeData[train, ]
collegeTest <- collegeData[-train, ]
xTest <- collegeTest$Enroll</pre>
```

Model fitting:

Fit a random forest an k-nearest neighbor models

```
# Fit a random forest model
# Used throughout Section 2:
set.seed(101)
```

```
rf <- randomForest(Enroll ~ ., data = collegeTrain)</pre>
# Check mse for rf model:
predRf <- predict(rf, newdata = collegeTest)</pre>
mspeRf <- mean((yTest - predRf)^2)</pre>
mspeRf
## [1] 0.03424027
# Fit an mlr3 knn model
# Used in Section 2.3:
knnT <- TaskRegr$new(id = "knn", backend = collegeTrain, target = "Enroll")</pre>
knnL <- lrn("regr.kknn")</pre>
knnMod <- knnL$train(knnT)</pre>
# Check mse for knn model:
pred <- predict(knnMod, newdata = collegeTest)</pre>
mspe <- mean((yTest - pred)^2)</pre>
mspe
## [1] 0.1031804
```

Create vivid matrix

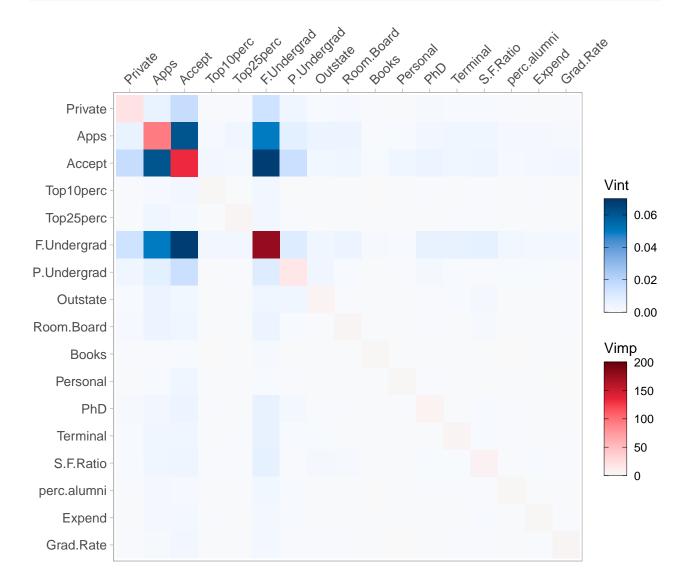
```
# Create unsorted vivid matrix for random forest fit:
# Used for Figure 1(a):
set.seed(101)
vividMatrixRF <- vivi(collegeTrain,</pre>
                       rf, "Enroll",
                       gridSize = 20,
                       reorder = FALSE)
# Sort matrix:
# Used for Figure 1(b):
vividMatrixRFSorted <- vividReorder(vividMatrixRF)</pre>
# Get agnostic VImp values instead of using random forests embedded VImps
# Used for Figure 2(b):
collegeVImps <- vivid:::vividImportance.default(rf,</pre>
  collegeTrain,
 "Enroll",
 importanceType = "agnostic",
 predictFun = CVpredict
# Update the matrix with the new VImp values and sort:
vividMatrixRFSorted_1 <- viviUpdate(vividMatrixRFSorted, collegeVImps)</pre>
vividMatrixRFSorted_1 <- vividReorder(vividMatrixRFSorted_1)</pre>
# Create vivid matrix for mlr3 knn fit using agnostic VImp
# Used for Figure 2(a):
set.seed(101)
```

```
knnMat <- vivi(
   fit = knnMod,
   data = collegeTrain,
   response = "Enroll",
   gridSize = 20,
   importanceType = "agnostic"
)</pre>
```

Visualisations for Section 2 -

Figure 1(a) & 1(b):

```
# Figure 1(a):
viviHeatmap(vividMatrixRF, angle = 45) # unsorted heatmap
# Figure 1(b):
viviHeatmap(vividMatrixRFSorted, angle = 45) # sorted heatmap
```



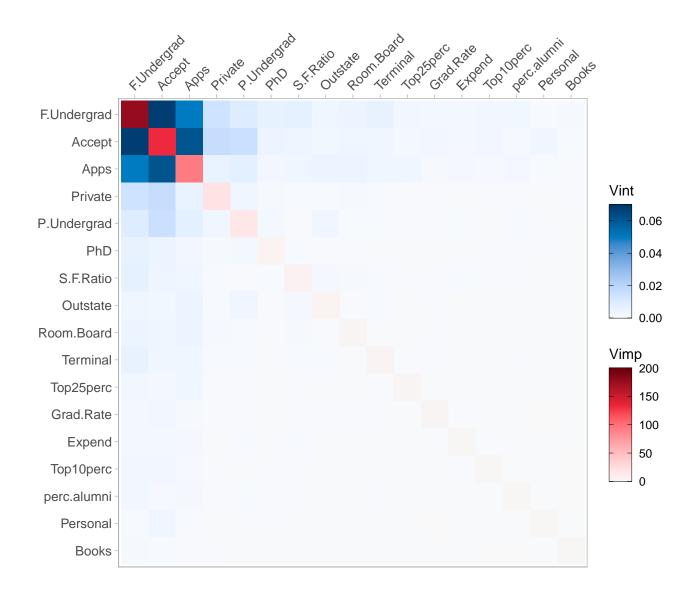
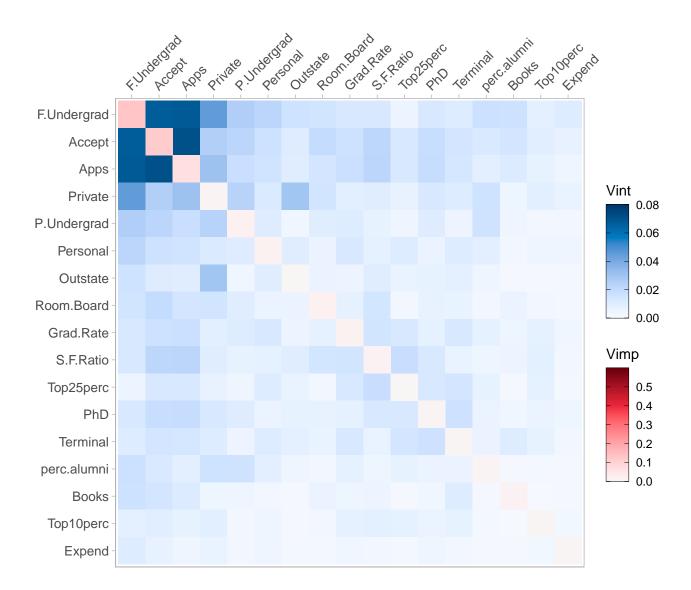


Figure 2(a) & 2(b):

```
# Figure 2(a):
viviHeatmap(knnMat, angle = 45, impLims = c(0, 0.6), intLims = c(0, 0.08)) # setting same VImp limits a
# Figure 2(b)
viviHeatmap(vividMatrixRFSorted_1, angle = 45, impLims = c(0, 0.6), intLims = c(0, 0.08)) # agnostic VI
```



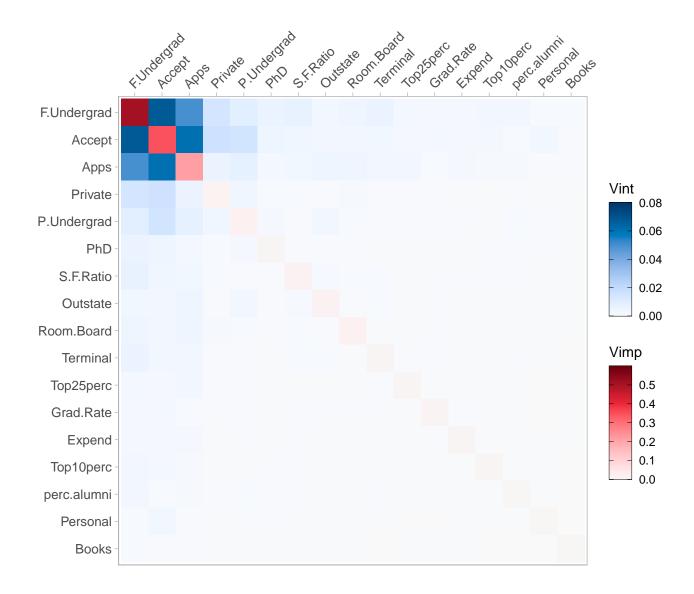
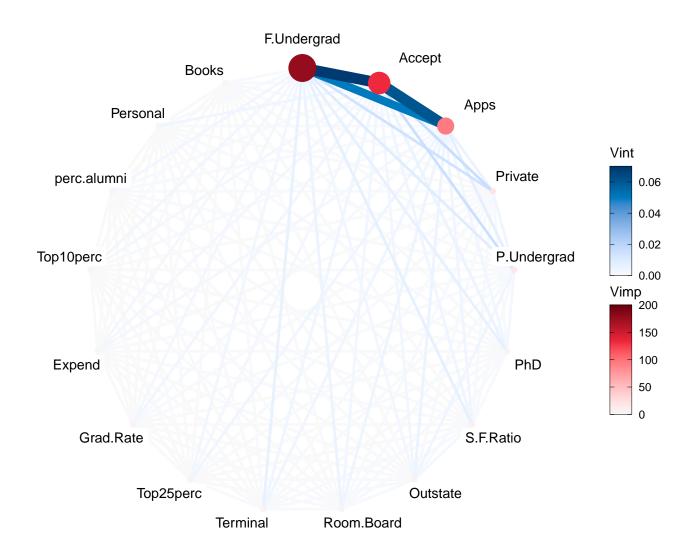
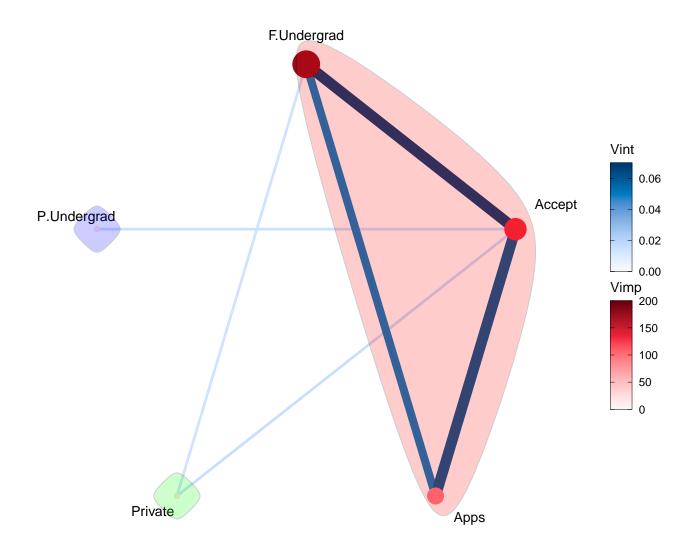


Figure 3(a) & 3(b):



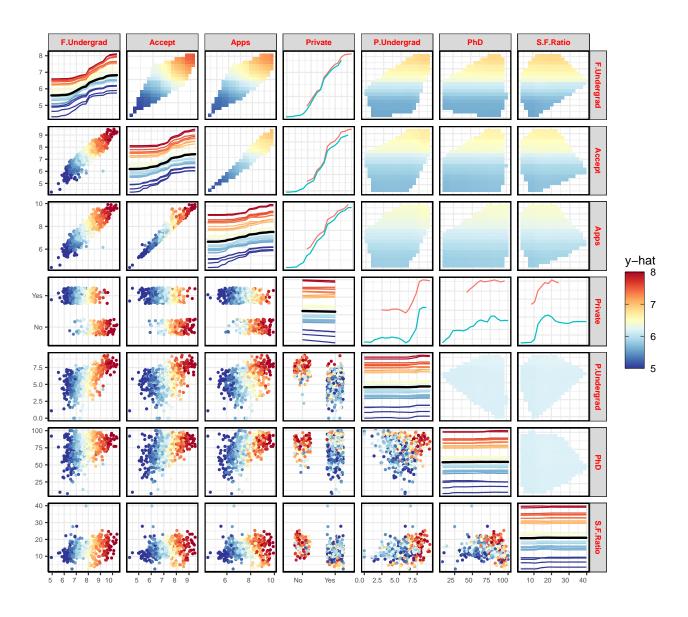


Visualisation for Section 3.2

Figure 4:

```
# Filter matrix:
nam <- colnames(vividMatrixRFSorted) # get names
nam <- nam[1:7] # filter names

# Create GPDP for Figure 4:
set.seed(101)
pdpPairs(collegeTrain,
    rf, "Enroll",
    gridSize = 20,
    vars = nam,
    convexHull = TRUE
)</pre>
```



Visualisation for Section 3.3 -

```
# Calculate the zpath using same threshold as Figure 3(b):
zpath <- zPath(vividMatrixRFSorted, 0.01)

# Create ZPDP using zpath for Figure 5:
set.seed(101)
pdpZen(collegeTrain,
    rf,
    "Enroll",
    gridSize = 20,
    zpath = zpath,
    convexHull = T
)</pre>
```

