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Code ▼
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Hide
library(caret)
 library(ranger)
 library(tidyverse)
 library(e1071)
library(readxl)
red <- read_excel("C:/Users/sarah/Desktop/info8000/winequalityred.xlsx",</pre>
                col_types = c("numeric", "numeric",
   "numeric", "numeric",
   "numeric", "numeric",
   "numeric", "numeric",
   "numeric", "numeric",
   "numeric", "numeric")
 library(caret)
 library(ranger)
 library(tidyverse)
library(e1071)
 set.seed(65468)
 inTraining<-createDataPartition(red$quality,p=.75,list = FALSE)</pre>
training<-red[inTraining,]
 testing<-red[-inTraining,]
 fitControl<-trainControl(## 10-fold CV
        method = "cv",
         number=10,
        ##repeated ten times
        repeats = 10)
  `repeats` has no meaning for this resampling method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hide
rfFit1<- train(quality ~.,data=training,
                                                                           trControl = fitControl)
rfFit1
Random Forest
1200 samples
        11 predictor
No pre-processing
Resampling: Cross-Validated (10 fold)
 Summary of sample sizes: 1080, 1079, 1080, 1081, 1082, 1080, \dots
Resampling results across tuning parameters:
        mtry RMSE
                                                                                      Rsquared MAE
                              0.5908197 0.4927346 0.4410006
                                   0.5928324 0.4795846 0.4372829
        11 0.5943479 0.4758715 0.4349772
RMSE was used to select the optimal % \left( 1\right) =\left( 1\right) \left( 
   model using the smallest value.
  The final value used for the model was
   mtry = 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hide
 save(file='rffit.model',rfFit1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hide
 rf2fit<- train(quality ~ alcohol + citricacid + volatileacidity + totalsulfurdioxide + sulphates + fixedacidity + density,
                                                                      data=training,
                                                                     method="rf".
                                                                     trControl=fitControl)
```

performs best when all variables are included! Still not great though...