## Mechanistic interpretation of feature importance

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Feature importance analysis from Random forest using Gini index (i.e., variance of the responses).

Creating a R function to perform feature importance

```
randomForest_feature_importance <- function(x) {</pre>
  library(doSNOW)
  library(foreach)
  library(parallel)
  cl <- makeCluster(8)</pre>
  registerDoSNOW(cl)
  results <- list(100)
  results <- foreach (i = 1:100) %dopar% {
    x \leftarrow na.omit(x)
    para <- dplyr::sample_n(x, size = 2571, replace = TRUE)</pre>
    in_train_para <- sample(nrow(para),</pre>
                               size = as.integer(nrow(para) * 0.8),
                               replace = FALSE)
    Train <- para[in_train_para, ]</pre>
    Test <- para[-in_train_para, ]</pre>
    rm(in_train_para)
    rm(Test)
    model <- ranger::ranger(pIC50~., data = Train, importance = 'impurity',</pre>
                                     write.forest = TRUE, save.memory = TRUE)
    rm(Train)
    importance <- model$variable.importance</pre>
    results[[i]] <- importance</pre>
  return(results)
  stopCluster(cl)
}
```

Generating the plot of feature importance

```
input <- readRDS(file = "data.Rds")
SubStructure_fingerPrintCount <- input$Substructure_fingerPrintCount
set.seed(10)
results_feature_importance_RF <- randomForest_feature_importance(SubStructure_fingerPrintCount)
data1 <- data.frame(results_feature_importance_RF)
data1 <- cbind(features = rownames(data1), data1)
library(reshape2)
data_melt <- suppressWarnings(melt(data1, id.vars = "features"))
#data_melt <- melt(data1, id.vars = "features")
data_melt$features <- factor(data_melt$features)
library(ggplot2)
set.seed(1)</pre>
```

```
plot_feature <- ggplot(data_melt, aes(x = reorder(features, value, FUN = median), y = value)) +</pre>
 geom_boxplot(fill = "#F8766D", colour = "black", alpha = 0.5) +
 theme_bw() + xlab("") + ylab("Gini Index") + coord_flip() + theme(
   axis.text.y = element_text(size = 20, colour = "black"),
   axis.text.x = element_text(size = 20, colour = "black"),
   #axis.title.x = element_blank(),
   plot.margin = unit(c(1, 1, 1, 1), "cm"),
   panel.border = element_rect(linetype = "solid", colour = "black", fill = NA, size = 1),
   axis.title = element_text(size = 25, face = "bold", colour = "black")
 )
plot_feature
  SubFPC2
SubFPC302
SubFPC274
 SubFPC32
SubFPC295
SubFPC275
  SubFPC3
  SubFPC1
SubFPC181
 SubFPC26
SubFPC137
SubFPC153
 SubFPC88
 SubFPC18
SubFPC298-
  SubFPC4
SubFPC135
 SubFPC49
SubFPC100
SubFPC171
  SubFPC5
SubFPC303
SubFPC182
                  1
SubFPC180
 SubFPC28-
```

200

400

**Gini Index** 

600

800

SubFPC85-