# Y-permutation Plot

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June 10, 2016

#### Function For Creating the Scrambling Plot

```
scrambling_R2 <- function(x) {</pre>
    library(doSNOW)
    library(foreach)
    library(parallel)
    cl <- makeCluster(8)</pre>
    registerDoSNOW(cl)
    results <- list(50)
    results <- foreach(i = 1:50) %dopar% {
        x \leftarrow na.omit(x)
        para <- dplyr::sample_n(x, size = 2571, replace = TRUE)</pre>
        in_train_para <- sample(nrow(para), size = as.integer(nrow(para) * 0.8),</pre>
             replace = FALSE)
        Train <- para[in_train_para, ]</pre>
        Test <- para[-in_train_para, ]</pre>
        pIC50 <- gtools::permute(Train$pIC50)
        pIC50 <- data.frame(pIC50)
        fake_data <- cbind(pIC50, Train[, 2:ncol(Train)])</pre>
        # ctrl <- caret::trainControl(method = 'repeatedcv', number = 10, repeats =
         # 1) tune <- caret::train(pIC50~., data = fake_data, method = 'rf',
         # trControl = ctrl, tuneLength = 10)
        fit <- ranger::ranger(pIC50 ~ ., data = fake_data, write.forest = TRUE,</pre>
             save.memory = TRUE)
         # actual <- train$Activity
        prediction <- predict(fit, Train)</pre>
        prediction <- prediction$predictions</pre>
        value <- data.frame(obs = Train$pIC50, pred = prediction)</pre>
        rm(fit)
        rm(prediction)
        labeling <- c("obs", "pred")</pre>
        colnames(value) <- labeling</pre>
        results[[i]] <- caret::defaultSummary(value)</pre>
    }
    R2 <- data.frame(results)
    R2 \leftarrow t(R2)
    R2 <- as.numeric(R2[, 2])
    R2 \leftarrow round(R2, digits = 5)
    return(R2)
    stopCluster(cl)
real_R2 <- function(x) {</pre>
```

```
x <- na.omit(x)
    para <- dplyr::sample_n(x, size = 2571, replace = TRUE)</pre>
    in_train_para <- sample(nrow(para), size = as.integer(nrow(para) * 0.8),</pre>
        replace = FALSE)
    Train <- para[in_train_para, ]</pre>
    Test <- para[-in_train_para, ]</pre>
    # ctrl <- caret::trainControl(method = 'repeatedcv', number = 10, repeats =</pre>
    # 1) tune <- train(pIC50~., data = x, method = 'rf', trControl = ctrl,
    # tuneLength = 10)
    fit <- ranger::ranger(pIC50 ~ ., data = Train, write.forest = TRUE, save.memory = TRUE)
    # actual <- train$Activity</pre>
    prediction <- predict(fit, Train)</pre>
    prediction <- prediction$predictions</pre>
    value <- data.frame(obs = Train$pIC50, pred = prediction)</pre>
    labeling <- c("obs", "pred")</pre>
    colnames(value) <- labeling</pre>
    result <- caret::defaultSummary(value)
    R2 <- as.data.frame(result)
    R2 \leftarrow R2[2,]
    R2 <- round(R2, digits = 5)
    return(R2)
}
scrambling_Q2 <- function(x) {</pre>
    library(doSNOW)
    library(foreach)
    library(parallel)
    cl <- makeCluster(8)</pre>
    registerDoSNOW(cl)
    results <- list(50)
    results <- foreach(i = 1:50) %dopar% {
        para <- dplyr::sample_n(x, size = 2571, replace = TRUE)</pre>
        in_train_para <- sample(nrow(para), size = as.integer(nrow(para) * 0.8),
             replace = FALSE)
        Train <- para[in_train_para, ]</pre>
        Test <- para[-in_train_para, ]</pre>
        pIC50 <- gtools::permute(Train$pIC50)</pre>
        pIC50 <- data.frame(pIC50)
        myData <- cbind(pIC50, Train[, 2:ncol(x)])</pre>
        index <- sample(1:k, nrow(myData), replace = TRUE)</pre>
        folds <- 1:k
        myRes = data.frame()
        for (j in 1:k) {
             training <- subset(myData, index %in% folds[-j])</pre>
             testing <- subset(myData, index %in% c(j))</pre>
             \# pIC50 <- gtools::permute(training$pIC50) pIC50 <- data.frame(pIC50)
             # fake_data <- cbind(pIC50, training[2:ncol(training)]) ctrl <-</pre>
             # caret::trainControl(method = 'repeatedcv', number = 10, repeats = 1) tune
             # <- caret::train(pIC50~., data = training, method = 'rf', trControl = ctrl,
             # tuneLength = 10)
             fit <- ranger::ranger(pIC50 ~ ., data = training, write.forest = TRUE,
```

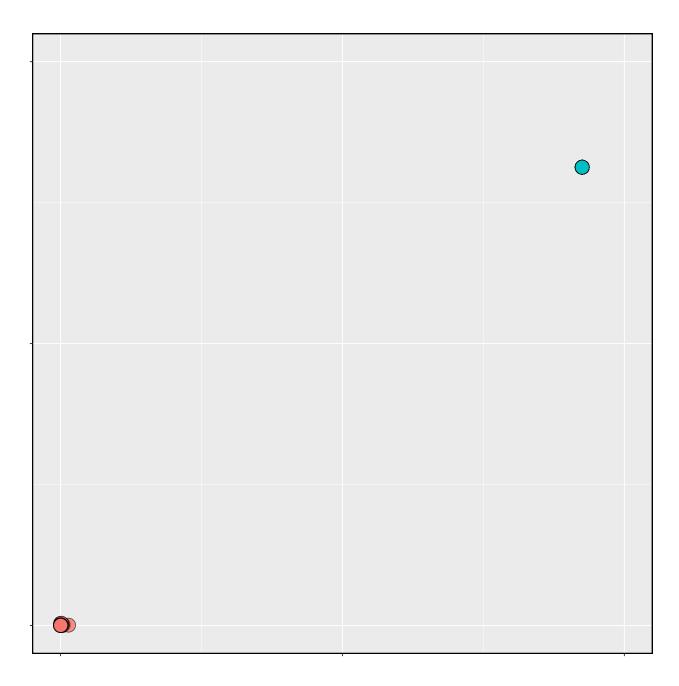
```
save.memory = TRUE)
             # actual <- train$Activity</pre>
             prediction <- predict(fit, testing)</pre>
             prediction <- prediction$predictions</pre>
             value <- data.frame(obs = testing$pIC50, pred = prediction)</pre>
             # fit <- randomForest::randomForest(pIC50~., data = training, mtry =
             # tune$bestTune[[1]]) prediction <- predict(fit, testing) value <-
             # data.frame( obs = testing$pIC50, pred = prediction)
             myRes <- rbind(myRes, value)</pre>
        value <- myRes
        labeling <- c("obs", "pred")</pre>
        rm(tune)
        rm(fit)
        rm(prediction)
        rm(ctrl)
        colnames(value) <- labeling</pre>
        results[[i]] <- caret::defaultSummary(value)</pre>
    Q2 <- data.frame(results)
    Q2 \leftarrow t(Q2)
    Q2 <- as.numeric(Q2[, 2])
    Q2 <- round(Q2, digits = 5)
    return(Q2)
    stopCluster(cl)
}
real_Q2 <- function(x) {</pre>
    para <- dplyr::sample_n(x, size = 2571, replace = TRUE)</pre>
    in_train_para <- sample(nrow(para), size = as.integer(nrow(para) * 0.8),</pre>
        replace = FALSE)
    Train <- para[in_train_para, ]</pre>
    Test <- para[-in_train_para, ]</pre>
    myData <- Train
    k = 10
    index <- sample(1:k, nrow(myData), replace = TRUE)</pre>
    folds <- 1:k
    myRes <- data.frame()</pre>
    for (j in 1:k) {
        training <- subset(myData, index %in% folds[-j])</pre>
        testing <- subset(myData, index %in% c(j))</pre>
        # ctrl <- caret::trainControl(method = 'repeatedcv', number = 10, repeats =
        # 1) tune <- train(pIC50~., data = training, method = 'rf', trControl =
        # ctrl, tuneLength = 10)
        fit <- ranger::ranger(pIC50 ~ ., data = training, write.forest = TRUE,</pre>
             save.memory = TRUE)
         # actual <- train$Activity
        prediction <- predict(fit, testing)</pre>
        prediction <- prediction$predictions</pre>
        value <- data.frame(obs = testing$pIC50, pred = prediction)</pre>
         # fit <- randomForest(pIC50~., data = training, mtry = tune$bestTune[[1]])
```

```
# prediction <- predict(fit, testing) value <- data.frame(obs =</pre>
                                              # testing$pIC50, pred = prediction)
                                             myRes <- rbind(myRes, value)</pre>
                      value <- myRes</pre>
                      labeling <- c("obs", "pred")</pre>
                       colnames(value) <- labeling</pre>
                      result <- caret::defaultSummary(value)</pre>
                      Q2 <- as.data.frame(result)
                      Q2 \leftarrow Q2[2,]
                      Q2 \leftarrow round(Q2, digits = 5)
                      return(Q2)
}
data_pre <- function(x) {</pre>
                      fake_R2 <- scrambling_R2(x)</pre>
                      real_R2 <- real_R2(x)</pre>
                      fake_Q2 <- scrambling_Q2(x)</pre>
                      real_Q2 <- real_Q2(x)</pre>
                      fake_R2 <- as.data.frame(fake_R2)</pre>
                      fake_R2$Label <- "Fake"</pre>
                      real_R2 <- as.data.frame(real_R2)</pre>
                      real_R2$Label <- "Real"</pre>
                      fake_Q2 <- as.data.frame(fake_Q2)</pre>
                      fake Q2$Label <- "Fake"</pre>
                      real Q2 <- as.data.frame(real Q2)</pre>
                      real Q2$Label <- "Real"</pre>
                      fake <- cbind(fake_R2, fake_Q2)</pre>
                      real <- cbind(real_R2, real_Q2)</pre>
                       combine <- data.frame(fake, real)</pre>
                      return(combine)
plot_scrambling <- function(x) {</pre>
                       ok <- data_pre(x)</pre>
                       colnames(ok) <- c("R2", "Label", "Q2", "Label", "R2", "Label", "Q2", "Label")
                      R2 \leftarrow ok[c(1, 5)]
                      R2 <- reshape2::melt(R2)
                      R2 <- R2$value
                      R2 <- data.frame(R2)
                      Q2 \leftarrow ok[c(3, 7)]
                      Q2 <- reshape2::melt(Q2)
                      Q2 <- Q2$value
                      Q2 <- data.frame(Q2)
                      Label <- c("Fake", "Fake", "Fa
                                              "Fake", "Fake", "Fake", "Fake", "Fake", "Fake", "Fake", "Fake",
                                              "Fake", "Fake", "Fake", "Fake", "Fake", "Real", "Real", "Real",
                                              "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real"
                                              "Real", "Real"
                                              "Real", "Real"
```

```
"Real", "Real"
              "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real", "Real",
              "Real", "Real")
data <- cbind(R2, Q2, Label)
library(ggplot2)
plot <- ggplot(data, aes(x = R2, y = Q2, colour = Label)) + geom_point(size = 7,</pre>
              colour = "black", aes(fill = factor(Label)), pch = 21, alpha = 0.8) +
              theme(legend.position = ("none"), axis.text = element_blank(), panel.border = element_rect(line
                           colour = "black", fill = NA, size = 1)) + xlab("") + ylab("") +
              # labs(y = expression(paste(italic(Q^2)))) + labs(x =
# expression(paste(italic(R^2)))) +
scale_x_continuous(limits = c(0, 1), breaks = seq(from = 0, to = 1, by = 0.5)) +
              scale_y_continuous(limits = c(0, 1), breaks = seq(from = 0, to = 1,
                           by = 0.5)
\# coord\_cartesian(ylim = c(0, 1), xlim = c(0, 1))
return(plot)
```

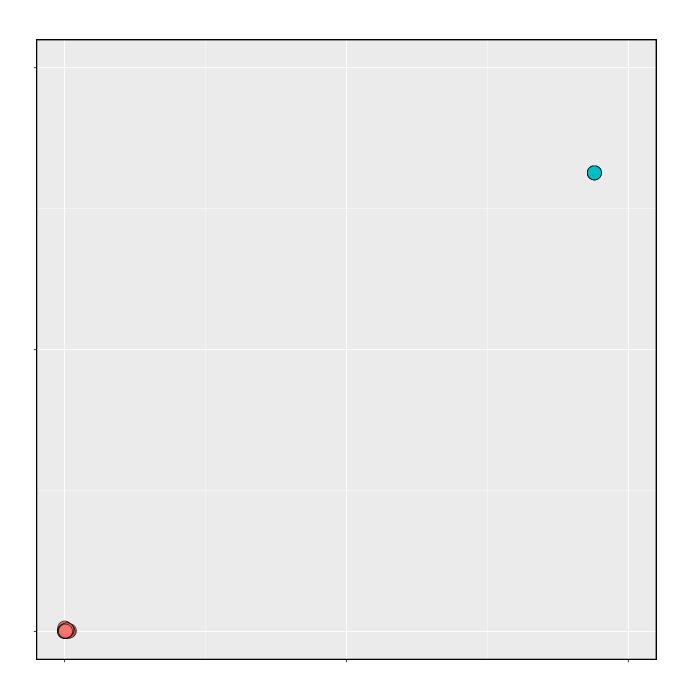
#### CDK

```
input <- readRDS("data.Rds")
CDK_Fingerprint <- input$FingerPrinter
plot_scrambling(CDK_Fingerprint)</pre>
```



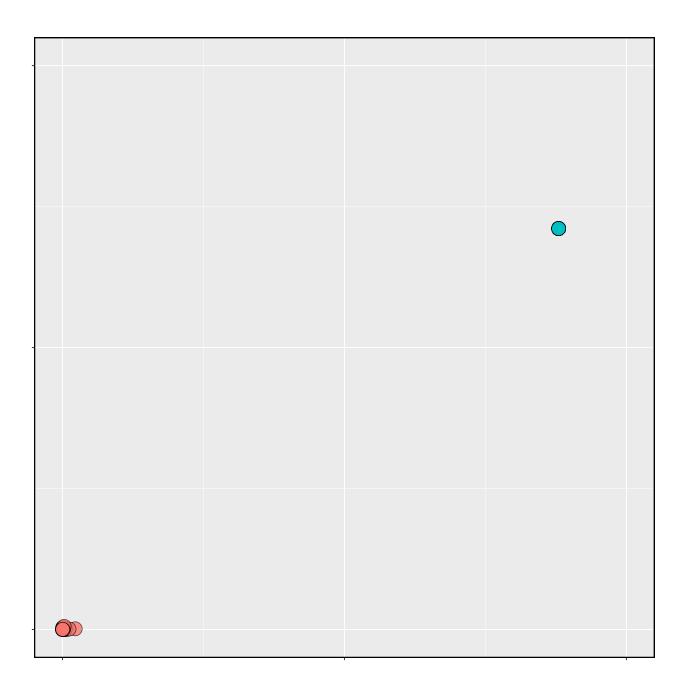
## CDK Extended

```
input <- readRDS("data.Rds")
Extended_CDK_Fingerprint <- input$Extended_finterPrinter
plot_scrambling(Extended_CDK_Fingerprint)</pre>
```



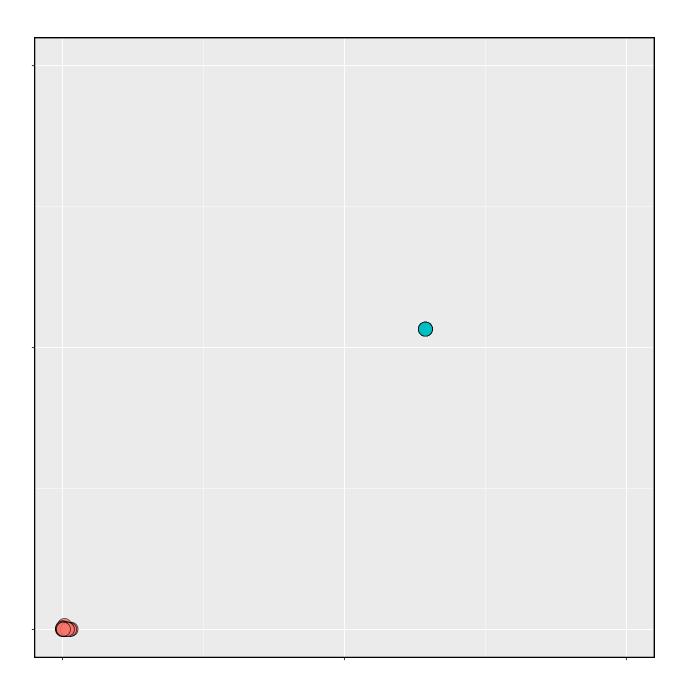
# CDK Graph Only

```
input <- readRDS("data.Rds")
Graph_CDK_Fingerprint <- input$GraphOnly_FingerPrinter
plot_scrambling(Graph_CDK_Fingerprint)</pre>
```



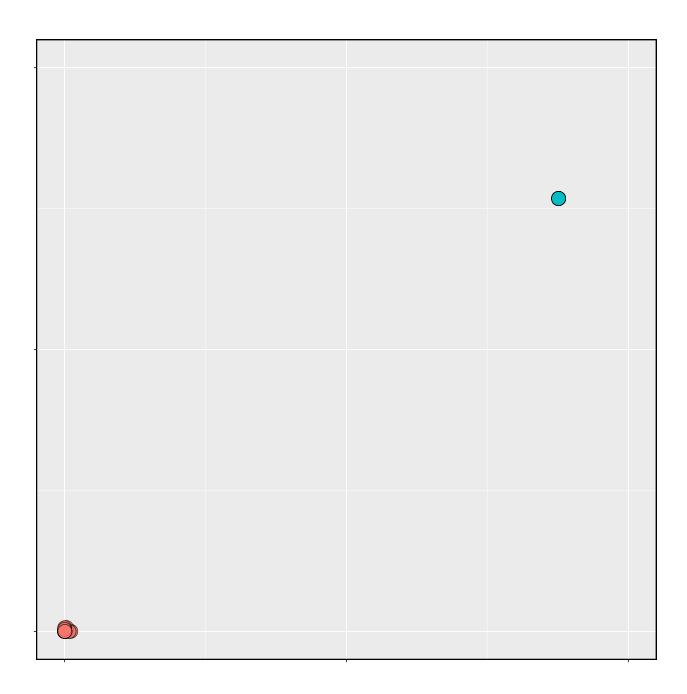
## E-State

```
input <- readRDS("data.Rds")
E_state_Fingerprint <- input$Estate_FingerPrinter
plot_scrambling(E_state_Fingerprint)</pre>
```



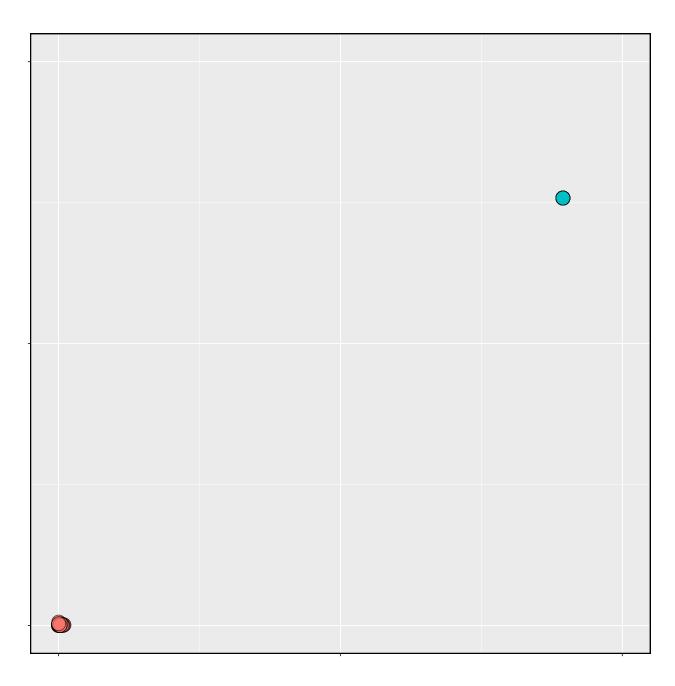
# MACCS

```
input <- readRDS("data.Rds")
MACCS_Fingerprint <- input$MACCS_FingerPrinter
plot_scrambling(MACCS_Fingerprint)</pre>
```



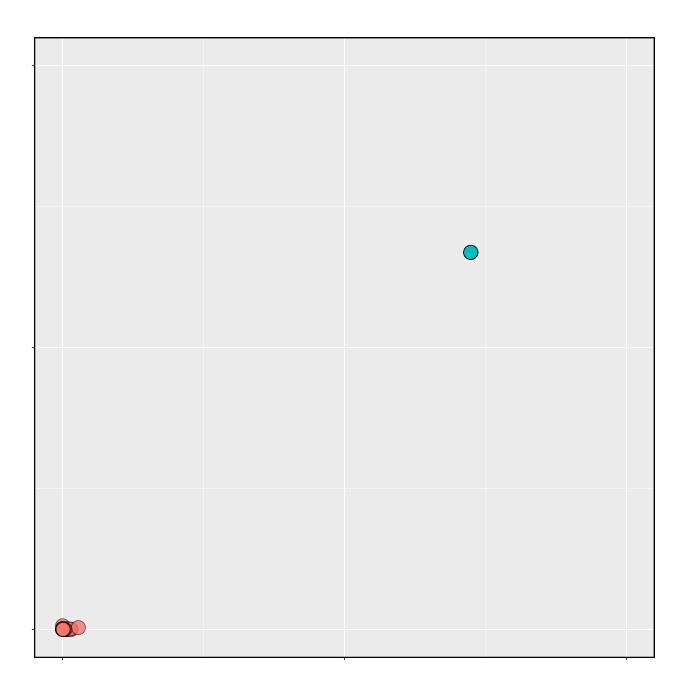
# PubChem

```
input <- readRDS("data.Rds")
PubChem_Fingerprint <- input$Pubchem_FingerPrinter
plot_scrambling(PubChem_Fingerprint)</pre>
```



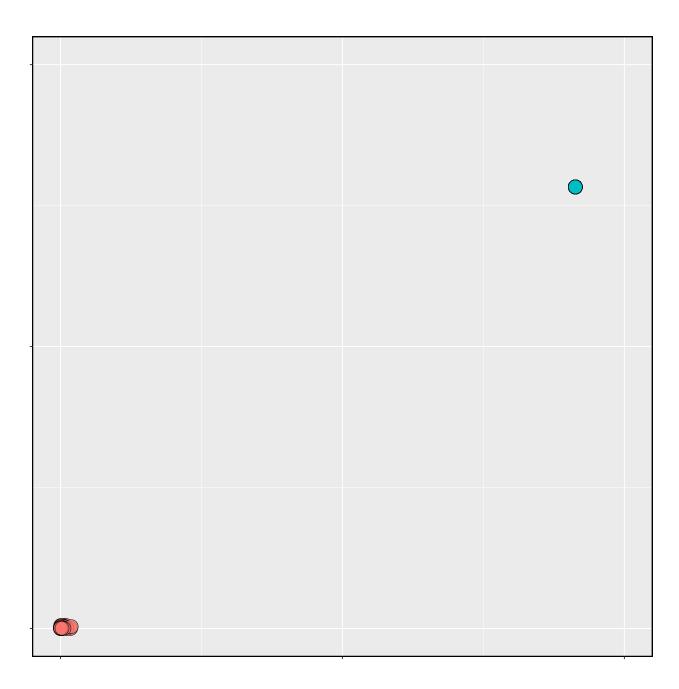
#### Substructure

```
input <- readRDS("data.Rds")
Substructure <- input$Substructure_fingerPrinter
plot_scrambling(Substructure)</pre>
```



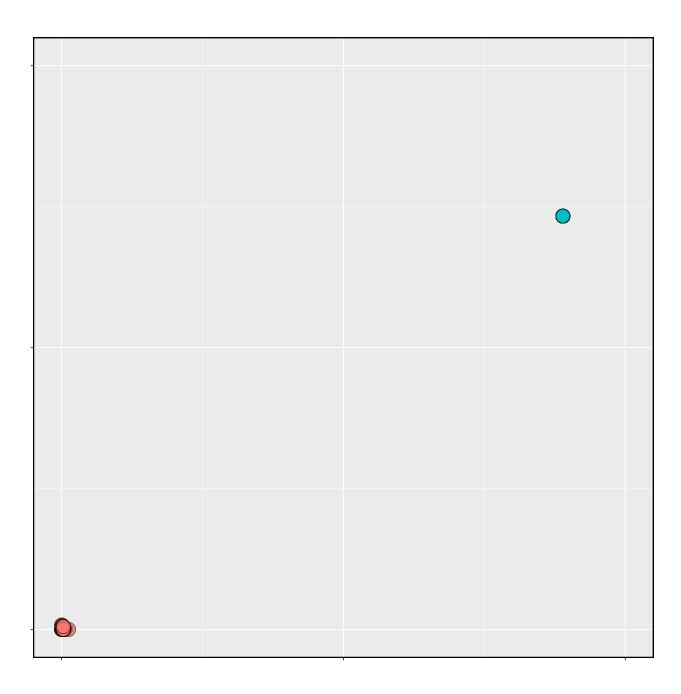
## Substructure Count

```
input <- readRDS("data.Rds")
Substructure_Count <- input$Substructure_fingerPrintCount
plot_scrambling(Substructure_Count)</pre>
```



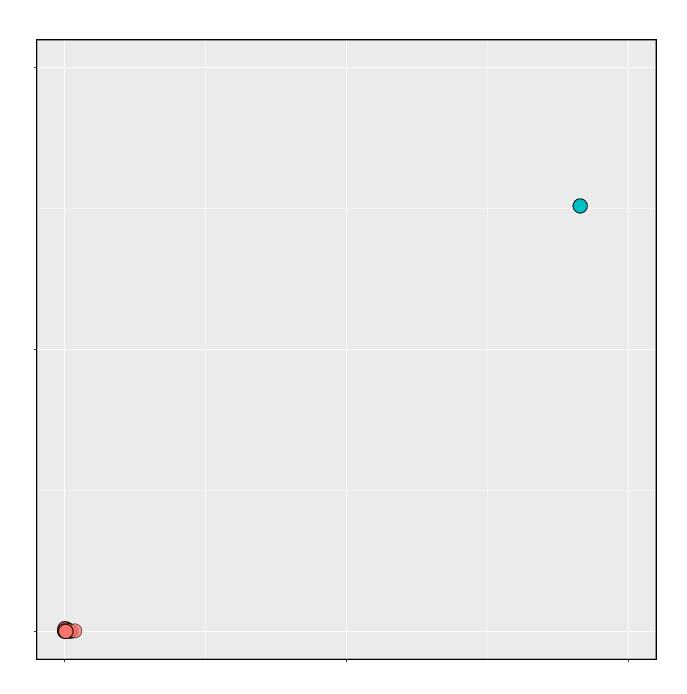
## Klekota-Roth

```
input <- readRDS("data.Rds")
Klekota_Roth <- input$KlekotaRoth_FingerPrinter
plot_scrambling(Klekota_Roth)</pre>
```



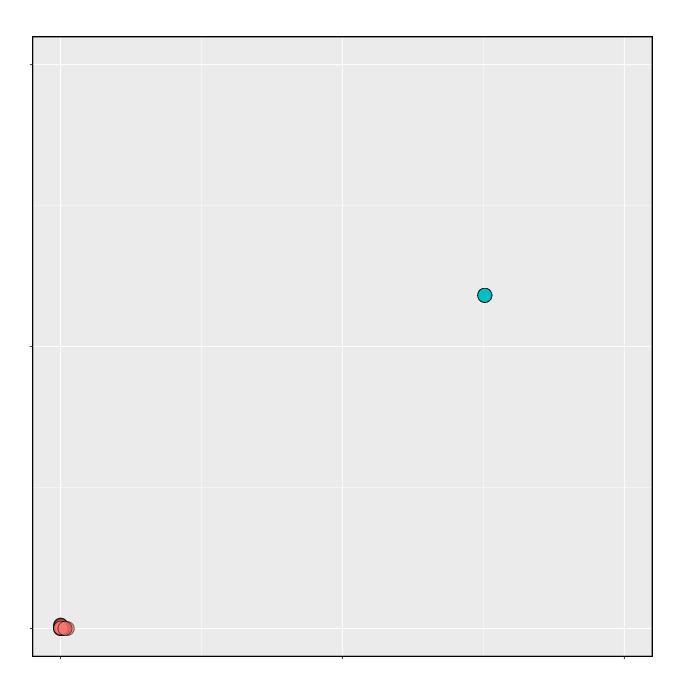
# Klekota-Roth Count

```
input <- readRDS("data.Rds")
Klekota_Roth_Count <- input$KlekotaRoth_FingerprintCount
plot_scrambling(Klekota_Roth_Count)</pre>
```



# 2D Atom Pairs

```
input <- readRDS("data.Rds")
Atom_Pairs <- input$AtomPairs2D_fingerPrinter
plot_scrambling(Atom_Pairs)</pre>
```



# 2D Atom Pairs Count

```
input <- readRDS("data.Rds")
Atom_Pairs_Count <- input$AtomPairs2D_fingerPrintCount
plot_scrambling(Atom_Pairs_Count)</pre>
```

