## Reading 12 Fingerprint from Padel Descriptor Calculator

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## Function for Correlation Cut of at 0.7

```
### Regression Acetylachlineesterase
read_file <- function(x){</pre>
  library(caret)
  library(data.table)
  data <- fread(x)
  IC50 nm <- data$IC50
  IC50 <- as.numeric(IC50_nm)*10^-9</pre>
  pIC50 \leftarrow -log10(IC50)
  data <- as.data.frame(data)</pre>
  descriptors <- data[, 2:ncol(data)]</pre>
  set.seed(1)
  yes <- descriptors[, -nearZeroVar(descriptors)]</pre>
  raw <- cor(yes)
  raw_2 <- raw[1: ncol(raw), 1:ncol(raw)]</pre>
  high <- findCorrelation(raw_2, cutoff = 0.7)
  filtered_descriptors <- yes[, -high]</pre>
  filtered_data <- cbind(pIC50, filtered_descriptors)</pre>
  return(filtered_data)
```

## Reading each data frame and printing out of the dimension of each data Frame

```
AtomPairs2D_fingerPrintCount <- read_file("data/2D_Atom_Pairs_Count.csv")
## Warning: package 'caret' was built under R version 3.2.4
## Loading required package: lattice
## Loading required package: ggplot2
## Warning: package 'ggplot2' was built under R version 3.2.4
AtomPairs2D_fingerPrinter <- read_file("data/2D_Atom_Pairs.csv")
Substructure_fingerPrintCount <- read_file("data/Substructure_Count.csv")
Substructure_fingerPrinter <- read_file("data/Substructure.csv")</pre>
Extended_finterPrinter <- read_file("data/CDK_Extended.csv")</pre>
FingerPrinter <- read_file("data/CDK.csv")</pre>
Estate_FingerPrinter <- read_file("data/E_State.csv")</pre>
GraphOnly_FingerPrinter <- read_file("data/CDK_Graph_Only.csv")</pre>
KlekotaRoth_FingerprintCount <- read_file("data/Klekota_Roth_Count.csv")</pre>
KlekotaRoth_FingerPrinter <- read_file("data/Klekota_Roth.csv")</pre>
MACCS FingerPrinter <- read file("data/MACCS.csv")
Pubchem_FingerPrinter <- read_file("data/PubChem.csv")</pre>
```

```
input <- list(AtomPairs2D_fingerPrintCount=AtomPairs2D_fingerPrintCount,</pre>
              AtomPairs2D fingerPrinter = AtomPairs2D fingerPrinter,
              Substructure fingerPrintCount = Substructure fingerPrintCount,
              Substructure_fingerPrinter = Substructure_fingerPrinter,
              Extended finterPrinter = Extended finterPrinter,
              FingerPrinter = FingerPrinter,
              Estate FingerPrinter = Estate FingerPrinter,
              GraphOnly_FingerPrinter = GraphOnly_FingerPrinter,
              KlekotaRoth_FingerprintCount = KlekotaRoth_FingerprintCount,
              KlekotaRoth_FingerPrinter = KlekotaRoth_FingerPrinter,
              MACCS_FingerPrinter = MACCS_FingerPrinter,
              Pubchem_FingerPrinter = Pubchem_FingerPrinter)
print(lapply(input, function(x) dim(x)))
## $AtomPairs2D_fingerPrintCount
## [1] 2571
             39
##
## $AtomPairs2D_fingerPrinter
## [1] 2571
             43
## $Substructure_fingerPrintCount
## [1] 2571
## $Substructure_fingerPrinter
## [1] 2571
             30
## $Extended_finterPrinter
## [1] 2571 940
##
## $FingerPrinter
## [1] 2571 941
## $Estate_FingerPrinter
## [1] 2571
             22
## $GraphOnly_FingerPrinter
## [1] 2571 201
##
## $KlekotaRoth_FingerprintCount
## [1] 2571
             72
## $KlekotaRoth_FingerPrinter
## [1] 2571 112
## $MACCS_FingerPrinter
## [1] 2571
             78
##
## $Pubchem_FingerPrinter
## [1] 2571 109
```