

Visual representation of Lipinski's descriptors

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Chemical space of AChE inhibitors

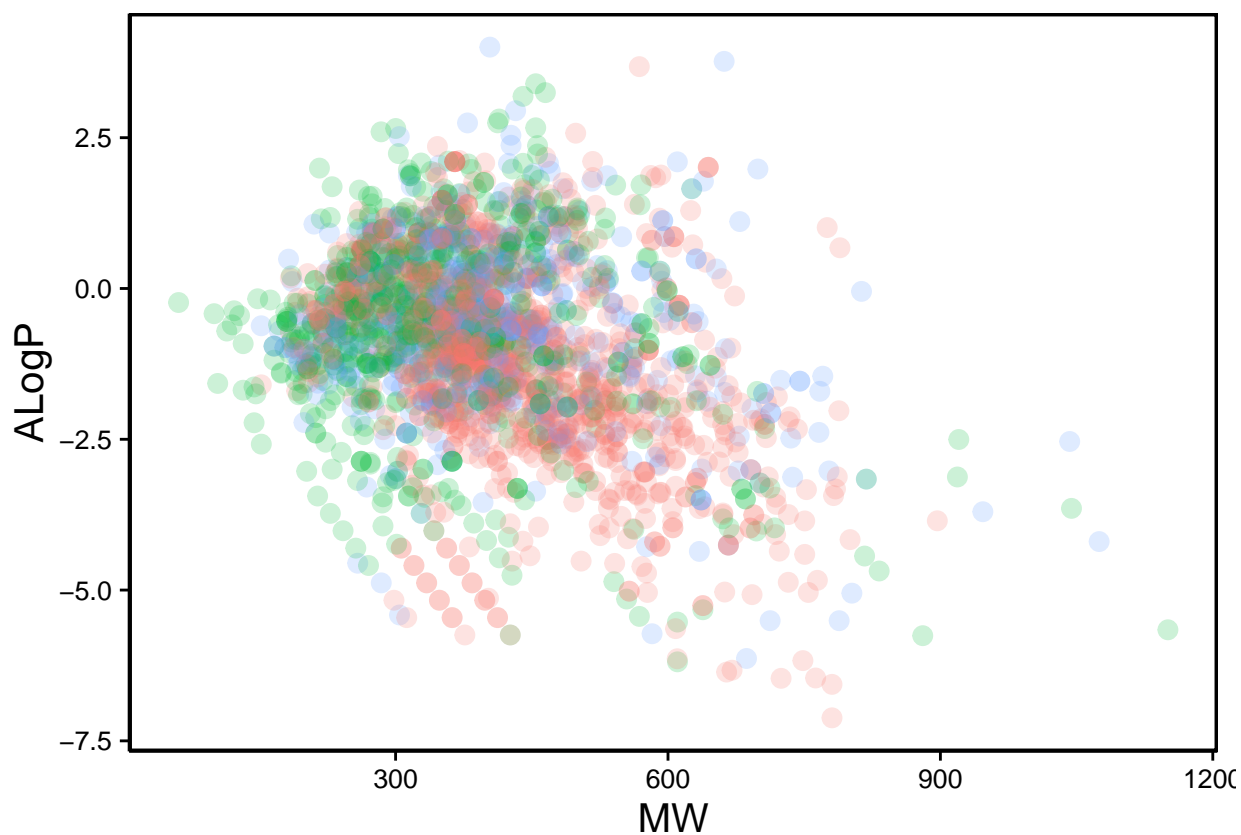
Convert IC50 to pIC50 as well as binning the activity (i.e., Active, Inactive and Intermediate)

```
data <- read.csv("Lipinski_descriptors.csv")
IC50_nm <- data$STANDARD_VALUE
IC50 <- as.numeric(IC50_nm)*10^-9
pIC50 <- -log10(IC50)
descriptors_name <- c("ALogP", "nHBAcc", "nHBDOn", "MW")
descriptors <- data[, descriptors_name]
#data_classification <- cbind(pIC50, descriptors)
Activity <- ifelse(pIC50 > 6, "Active",
                  ifelse((pIC50 <= 6) & (pIC50 < 5), "Inactive",
                        "Intermediate"))
Activity <- data.frame(Activity)
```

Visualization of Lipinski's descriptors

Chemical space of AChE inhibitors are shown as active (green), inactive (red) and intermediate (blue)

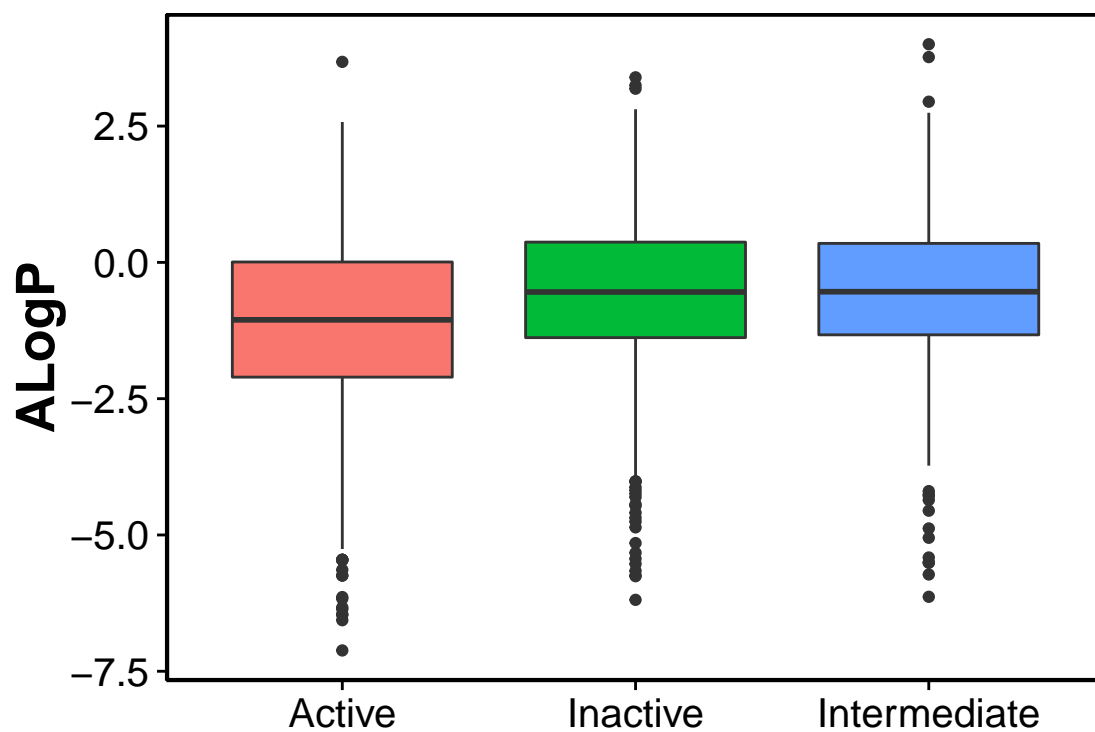
```
library(ggplot2)
library(cowplot)
data_2 <- cbind(Activity, descriptors)
data_2 <- na.omit(data_2)
data_plot <- data.frame(data_2)
p <- ggplot(data_plot, aes(MW, ALogP))
p <- p + geom_point(aes(colour = factor(Activity)), size = 3, alpha = 0.2)
p <- p + theme(legend.position = ("none"),
              panel.border = element_rect(linetype = "solid",
                                           colour = "black", fill = NA, size = 1),
              axis.text.x = element_text(colour = "black", size = 10),
              axis.text.y = element_text(colour = "black", size = 10),
              axis.title.x = element_text(colour = "black", size = 15),
              axis.title.y = element_text(colour = "black", size = 15)
              )
print(p)
```



Boxplot of Lipinski's rule-of-five descriptors (ALogP)

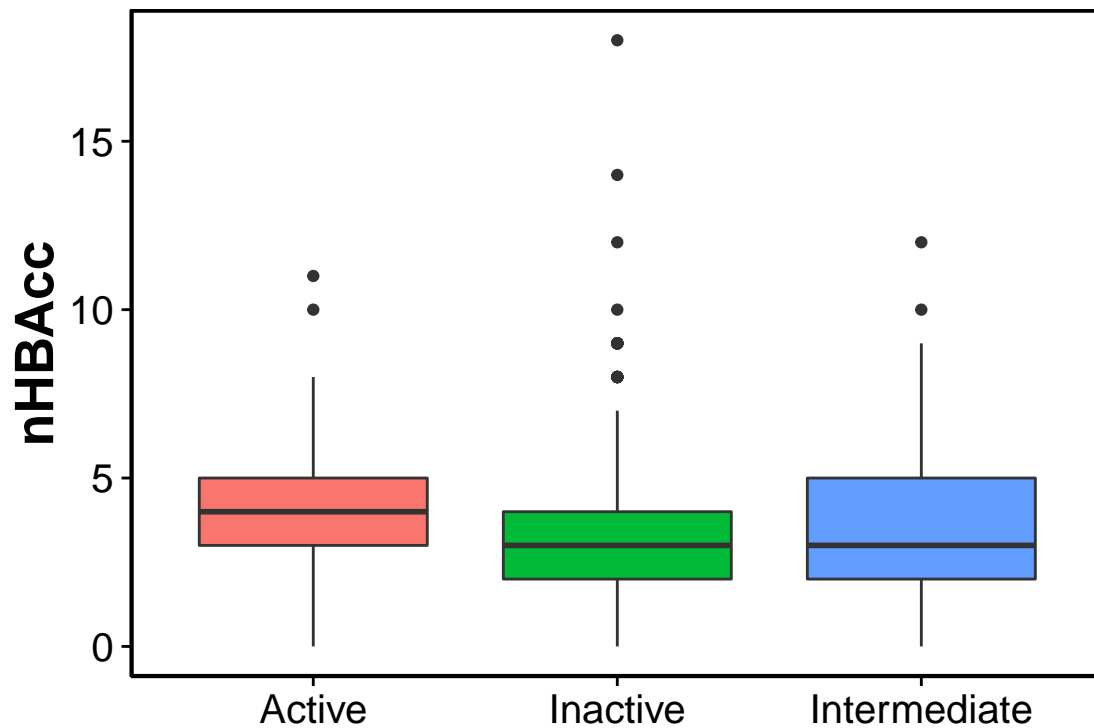
```
p_1 <- ggplot(data_plot, aes(factor(Activity), ALogP))
p_1 <- p_1 + geom_boxplot(aes(fill = factor(Activity)))
p_1 <- p_1 + theme(legend.position = ("none"),
  panel.border = element_rect(linetype = "solid",
    colour = "black", fill = NA, size = 1),
  axis.text.x = element_text(colour = "black", size = 15),
  axis.text.y = element_text(colour = "black", size = 15),
  plot.margin = unit(c(1, 1, 1, 1), "cm"),
  axis.title.y = element_text(size = 20, face = "bold"),
  axis.title.x = element_blank())

print(p_1)
```



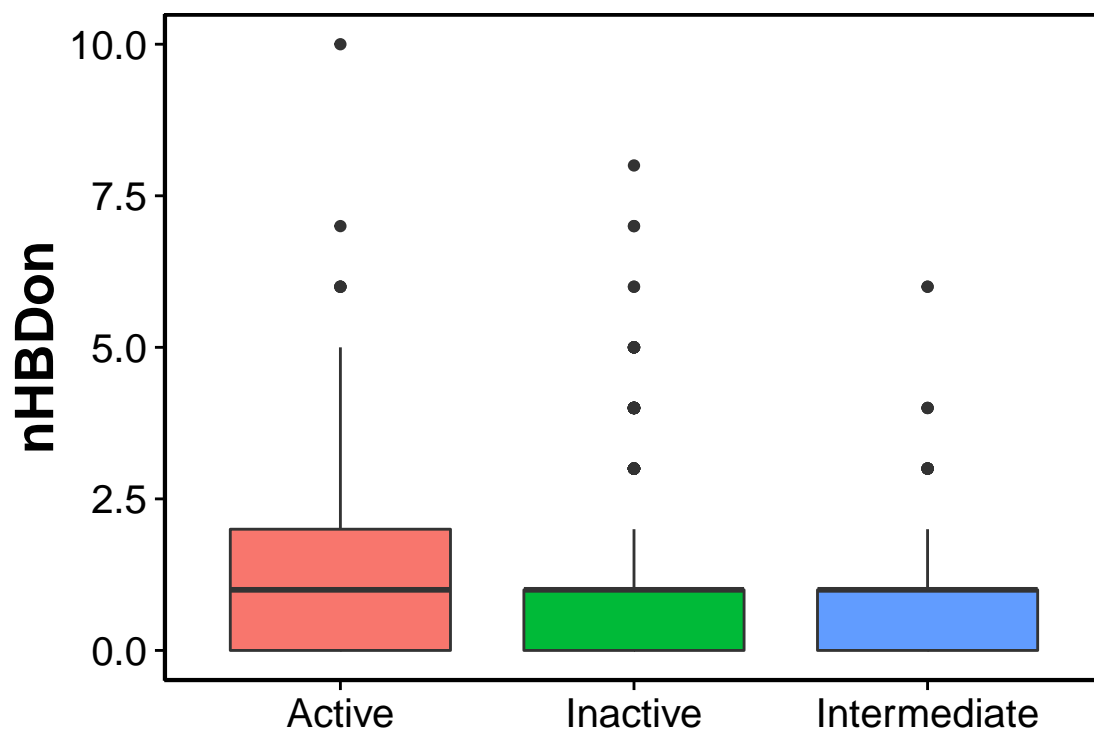
Boxplot of Lipinski's rule-of-five descriptors (nHBAcc)

```
p_2 <- ggplot(data_plot, aes(factor(Activity), nHBAcc))
p_2 <- p_2 + geom_boxplot(aes(fill = factor(Activity)))
p_2 <- p_2 + theme(legend.position = ("none"),
  panel.border = element_rect(linetype = "solid",
    colour = "black", fill = NA, size = 1),
  axis.text.x = element_text(colour = "black", size = 15),
  axis.text.y = element_text(colour = "black", size = 15),
  plot.margin = unit(c(1, 1, 1, 1), "cm"),
  axis.title.y = element_text(size = 20, face = "bold"),
  axis.title.x = element_blank())
print(p_2)
```



Boxplot of Lipinski's rule-of-five descriptors (nHBDon)

```
p_3 <- ggplot(data_plot, aes(factor(Activity), nHBDon))
p_3 <- p_3 + geom_boxplot(aes(fill = factor(Activity)))
p_3 <- p_3 + theme(legend.position = ("none"),
  panel.border = element_rect(linetype = "solid",
    colour = "black", fill = NA, size = 1),
  axis.text.x = element_text(colour = "black", size = 15),
  axis.text.y = element_text(colour = "black", size = 15),
  plot.margin = unit(c(1, 1, 1, 1), "cm"),
  axis.title.y = element_text(size = 20, face = "bold"),
  axis.title.x = element_blank())
print(p_3)
```



Boxplot of Lipinski's rule-of-five descriptors (MW)

```
p_4 <- ggplot(data_plot, aes(factor(Activity), MW))
p_4 <- p_4 + geom_boxplot(aes(fill = factor(Activity)))
p_4 <- p_4 + theme(legend.position = ("none"),
  panel.border = element_rect(linetype = "solid",
    colour = "black", fill = NA, size = 1),
  axis.text.x = element_text(colour = "black", size = 15),
  axis.text.y = element_text(colour = "black", size = 15),
  plot.margin = unit(c(1, 1, 1, 0.4), "cm"),
  axis.title.y = element_text(size = 20, face = "bold"),
  axis.title.x = element_blank())

print(p_4)
```

