

plots

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```
# column names
feat_names = colnames(features)[-1]

violin_plot <- function(features, y, fill_color = "darkgreen", logscale = FALSE) {
  # Remove NAs
  #df <- features %>% filter(!is.na(.data[[y]]))
  # Remove NAs and non-positive values for log scale
  df <- features %>% filter(!is.na(.data[[y]]) & (!logscale | .data[[y]] > 0))

  # Define scale transformation conditionally
  scale_y <- if (logscale) {
    scale_y_continuous(trans = "log",
                       breaks = scales::trans_breaks("log", function(x) exp(x)),
                       labels = scales::trans_format("log", scales::math_format(e^.x)))
  } else {
    scale_y_continuous() # Default linear scale
  }

  # Generate plot
  ggplot(df, aes(x = factor(1), y = .data[[y]], fill = fill_color)) + # Use factor(1) for a single x variable
    theme_bw() +
    geom_violin(color = "black") + # Violin plot with black border
    geom_hline(yintercept = median(df[[y]], na.rm = TRUE), linetype = "dashed", linewidth = 0.5) + # Dashed line at median
    stat_summary(fun = median, fun.min = median, fun.max = median,
                 geom = "crossbar", width = 0.25) + # Crossbar at median
    stat_summary(fun = mean, fun.min = mean, fun.max = mean,
                 geom = "point", shape = 5) + # Diamond shape for mean
    scale_y + # Apply the chosen scale (log or linear)
    scale_fill_manual(values = fill_color) + # Custom fill color
    theme(legend.position = "none") + # Remove legend
    labs(x = "All Loci", y = ifelse(logscale, paste0("ln(", y, ")"), y)) # Update y label to indicate log scale
}

log_list <- c("alignmentSPscore", "tree_rate", "tree_rate_var",
             "base_composition_variance", "alignment_length", "phyloinf")
counter <- 1
for (i in feat_names) {
  plot <- violin_plot(features, i) #plot the feature
  name <- paste0("p", counter) #generate a dynamic variable name
  assign(name, plot) #assign a plot to the dynamically named variable
  print(plot)
  if (i %in% log_list) {
```

```

plot <- violin_plot(features, i, fill_color = "green", logscale = TRUE) #plot feature with log scale
print(plot)
}
counter <- counter + 1 #increment counter
}

```





