## Tukey HSD tests

## 2024-10-04

```
df <- read.csv("scaffold split metrics.csv")</pre>
metric name <- "prec"</pre>
df[[metric_name]] <- as.numeric(df[[metric_name]])</pre>
df$method <- factor(df$method)</pre>
formula <- as.formula(paste(metric_name, "~ method + (1 | cv_cycle)"))</pre>
m <- lmer(formula, data = df)
m_sum <- summary(m)$coefficients</pre>
df_residual <- as.integer(m_sum[nrow(m_sum), "df"])</pre>
test_out <- multcomp::glht(m, linfct = mcp(method = "Tukey"), df = df_residual)
summary(test_out)
##
##
     Simultaneous Tests for General Linear Hypotheses
## Multiple Comparisons of Means: Tukey Contrasts
##
##
## Fit: lmer(formula = formula, data = df)
## Linear Hypotheses:
                                    Estimate Std. Error t value Pr(>|t|)
## chemprop_st - chemprop_mt == 0 -0.016123  0.006302 -2.558
                                                                    0.036 *
## lgbm_morgan - chemprop_mt == 0 - 0.007599 0.006302 - 1.206
                                                                     0.456
## lgbm_morgan - chemprop_st == 0 0.008524 0.006302
                                                           1.353
                                                                     0.374
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## (Adjusted p values reported -- single-step method)
confint(test_out)
##
##
     Simultaneous Confidence Intervals
## Multiple Comparisons of Means: Tukey Contrasts
##
##
## Fit: lmer(formula = formula, data = df)
##
## Quantile = 2.4187
## 95% family-wise confidence level
```

## 95% family-wise confidence level

