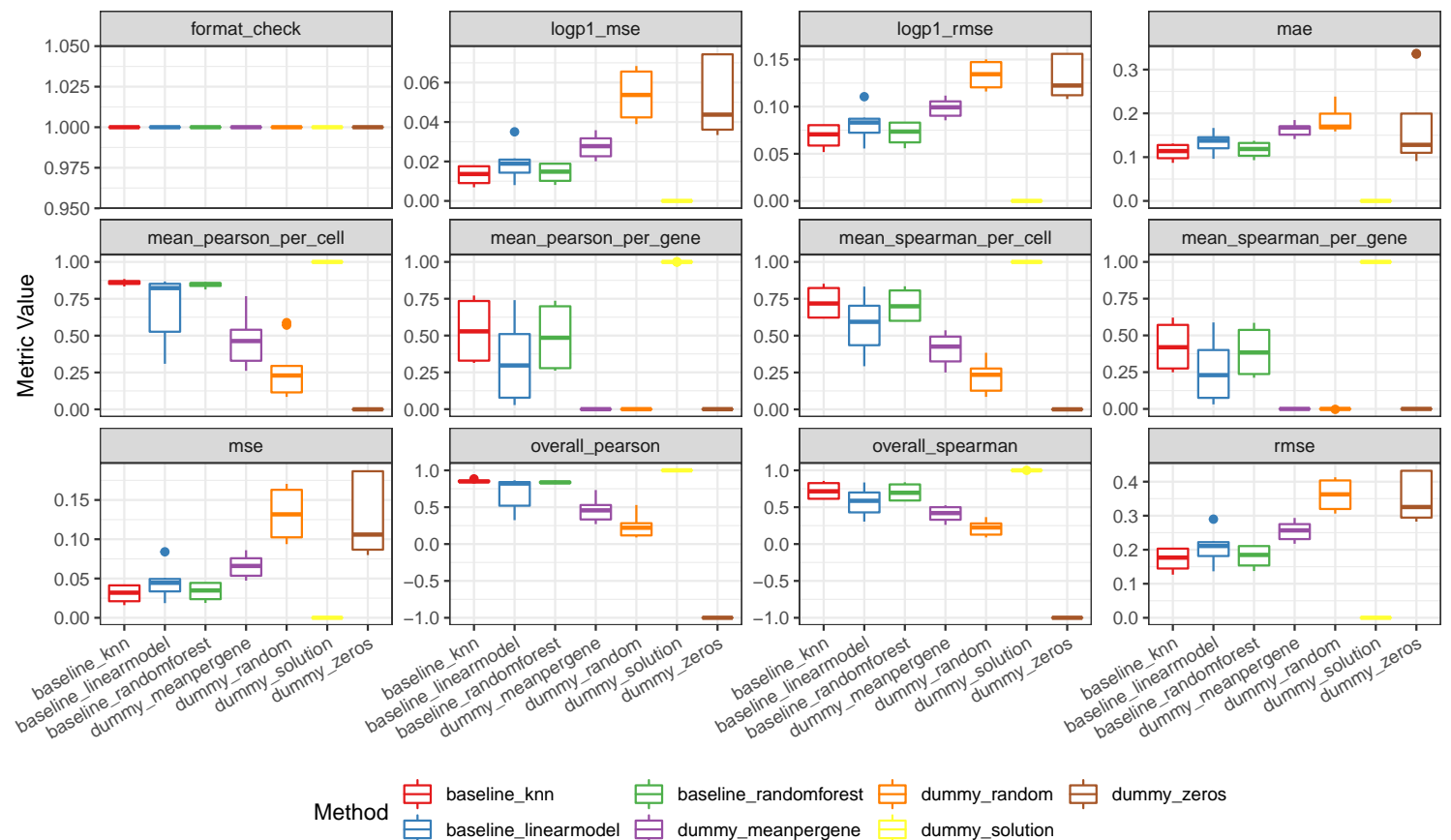
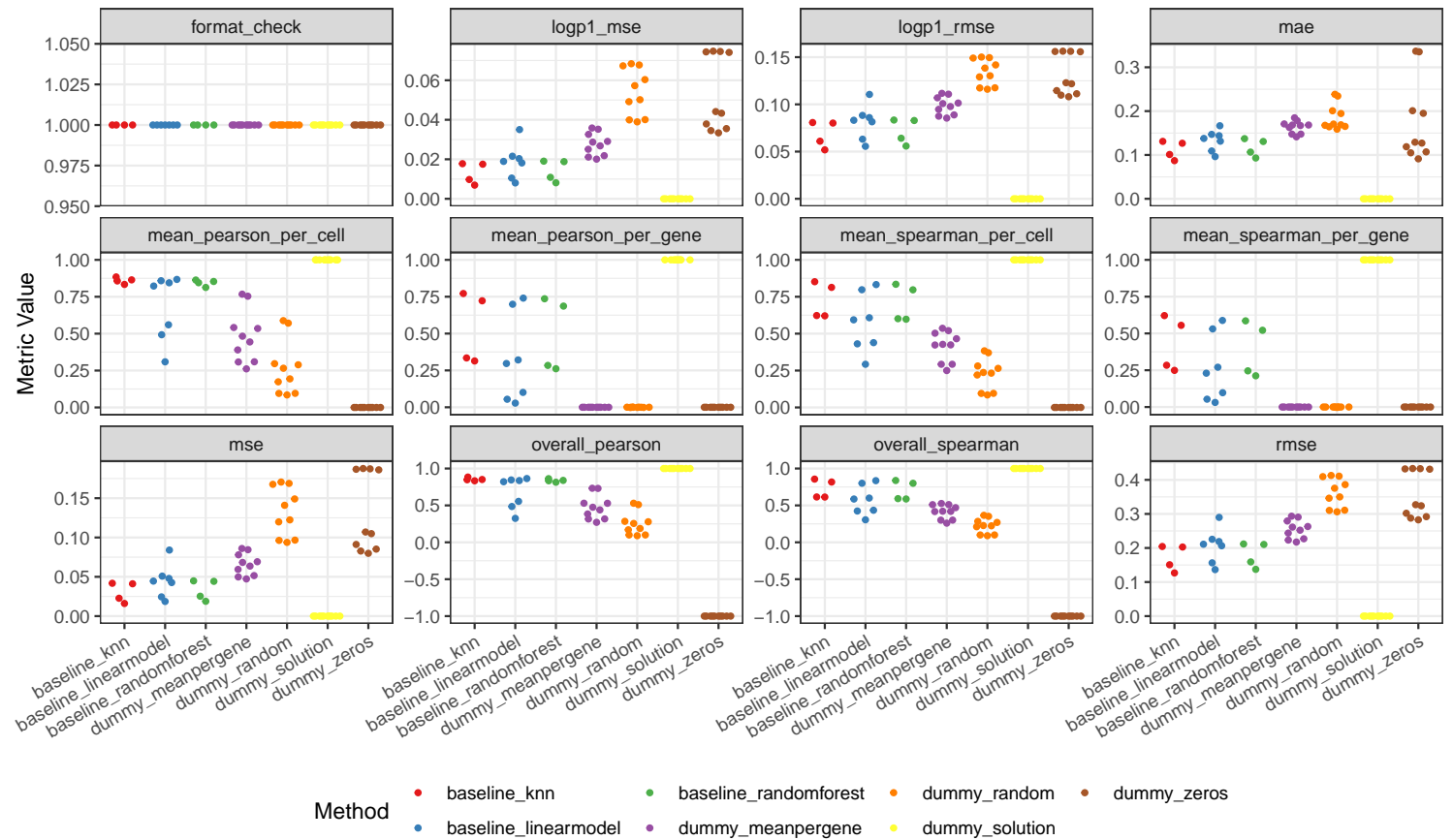


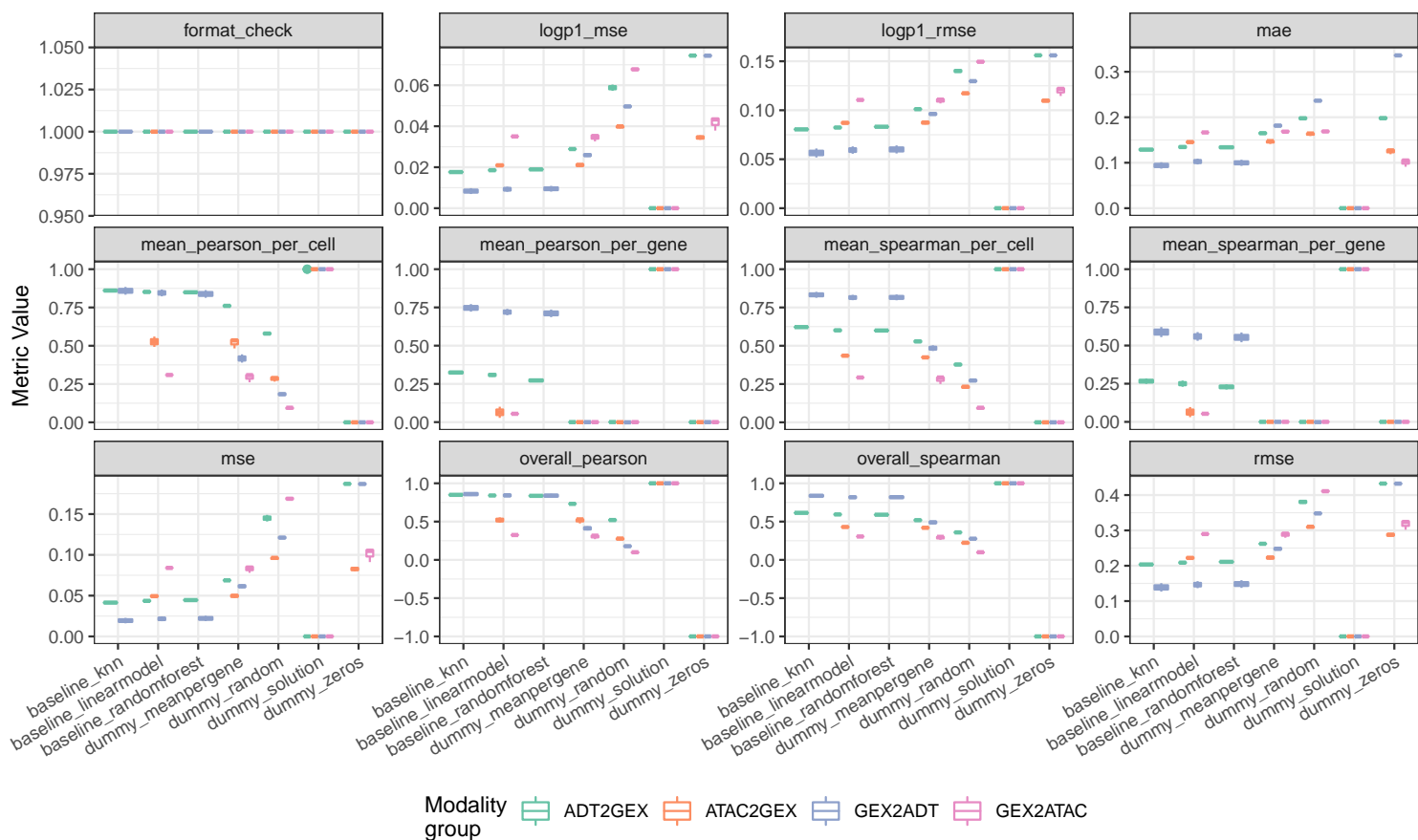
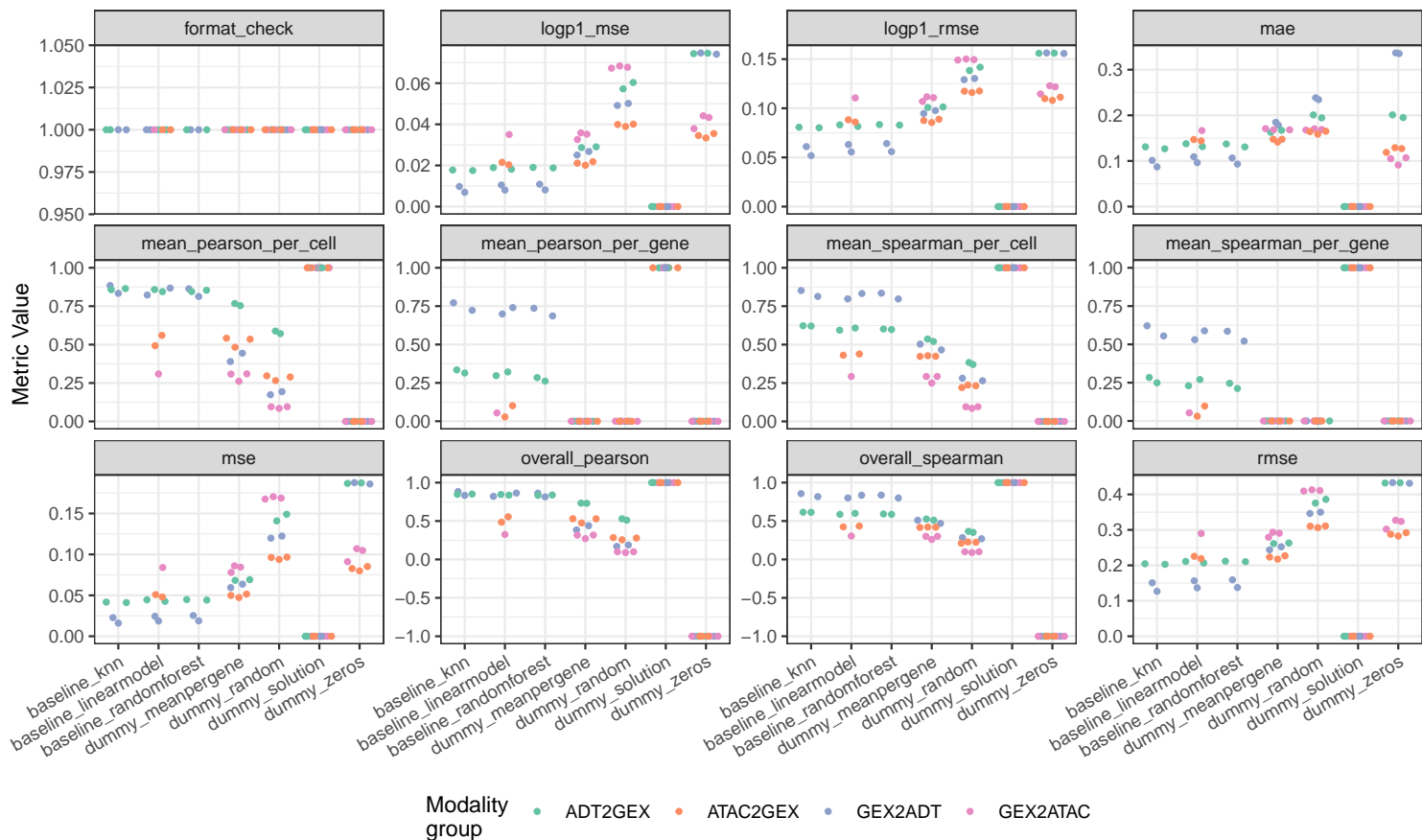
Predict Modality - Pilot Analysis

Visualise results

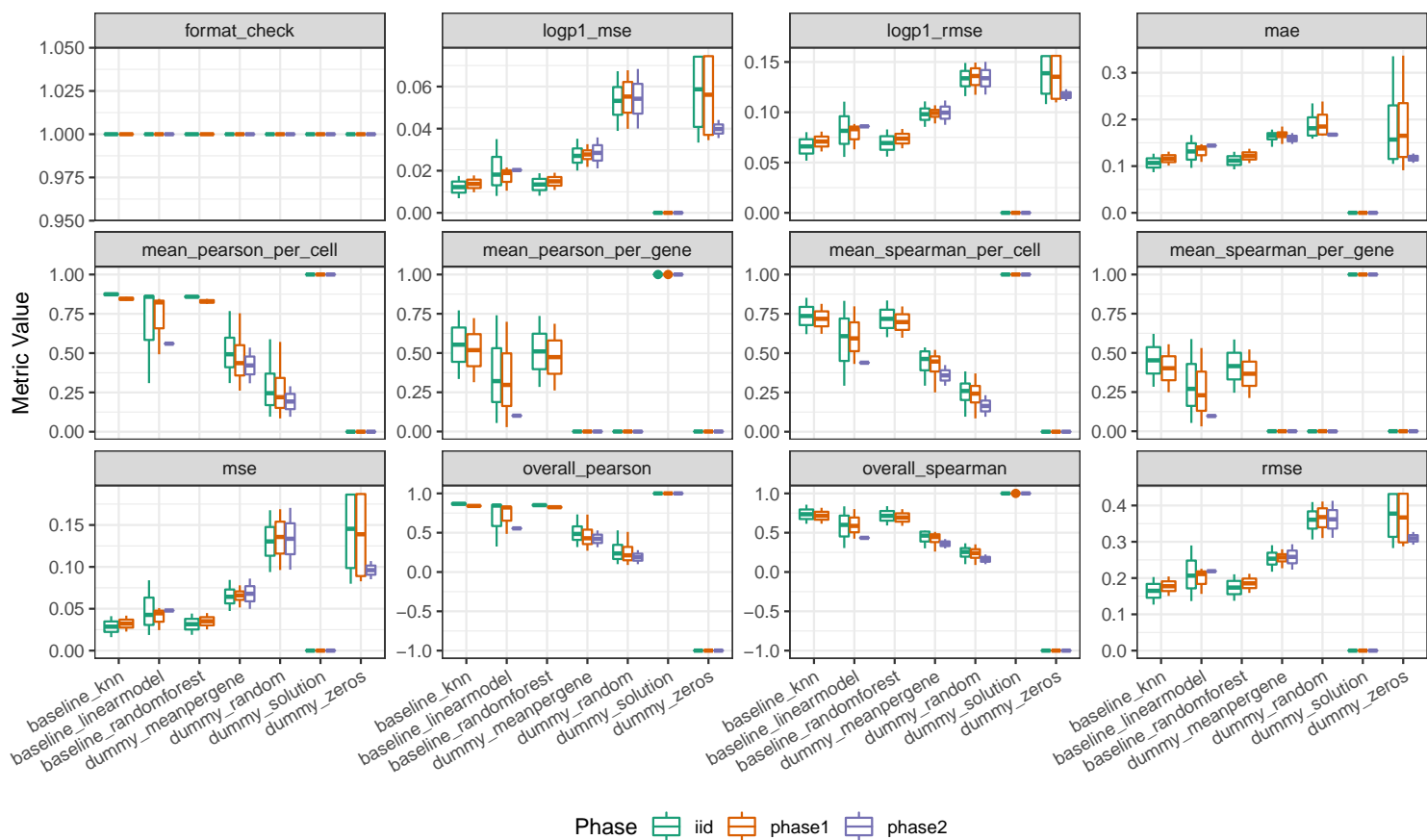
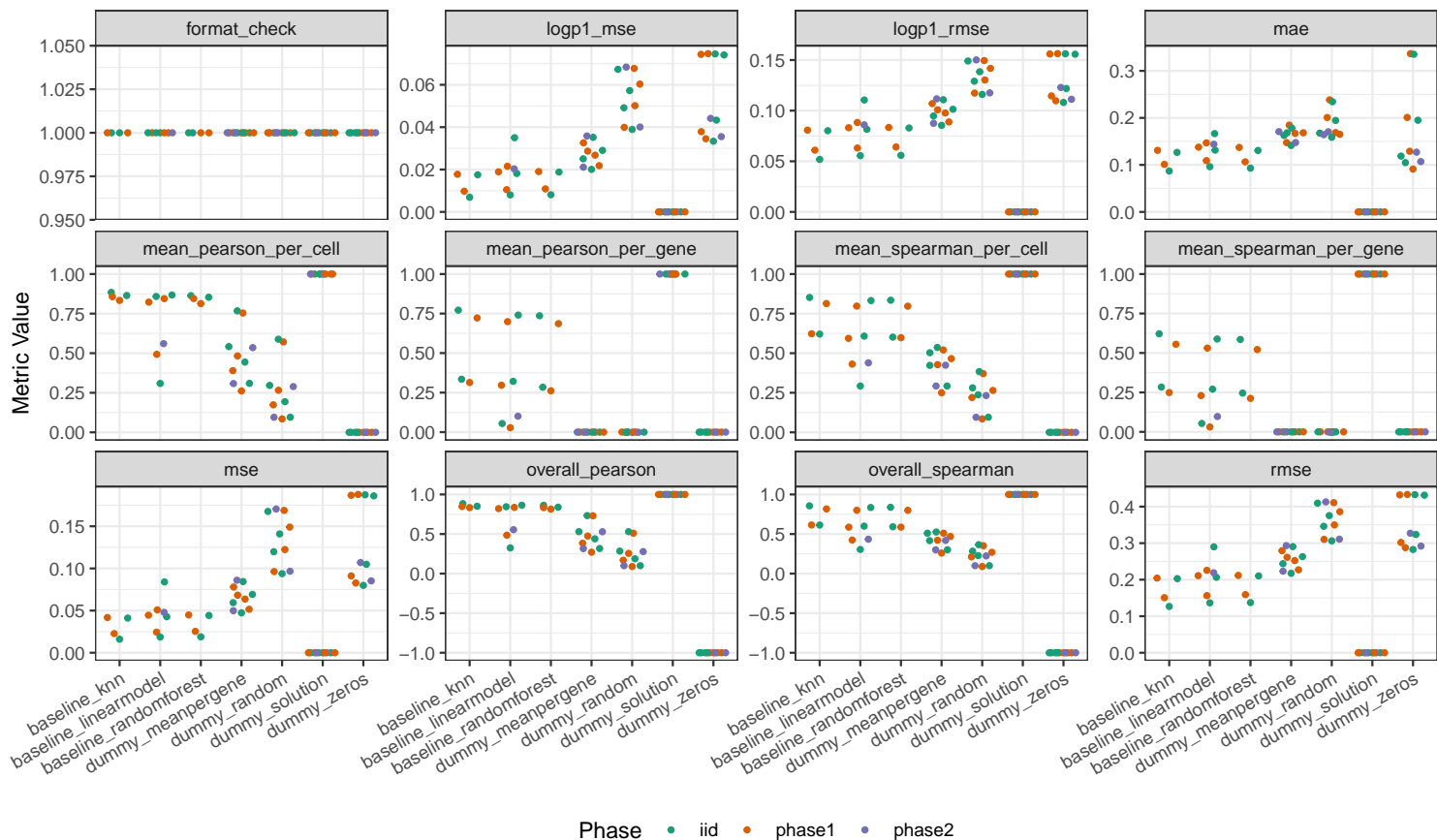
Colour by method.



Colour by modality group.



Colour by phase



Comparing most interesting metrics.

```
selected_metrics <- c("rmse", "mae", "mean_spearman_per_cell", "mean_spearman_per_gene", "overall_spearman")

GGally::ggpairs(
  df %>% na.omit,
  mapping = aes(colour = method_type, fill = method_type),
  columns = selected_metrics,
  diag = list(continuous = "barDiag", discrete = "barDiag", na = "naDiag")
) & theme_bw() &
  scale_colour_brewer(palette = "Set2") &
  scale_fill_brewer(palette = "Set2")

## Registered S3 method overwritten by 'GGally':
##   method from
##   +.gg      ggplot2
```

