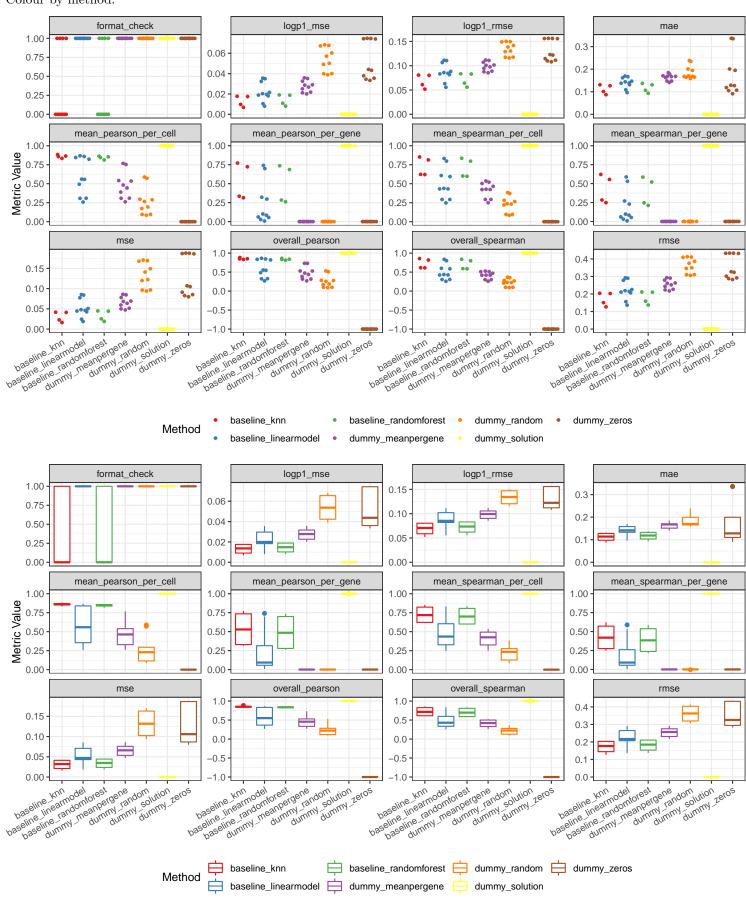
Predict Modality - Pilot Analysis

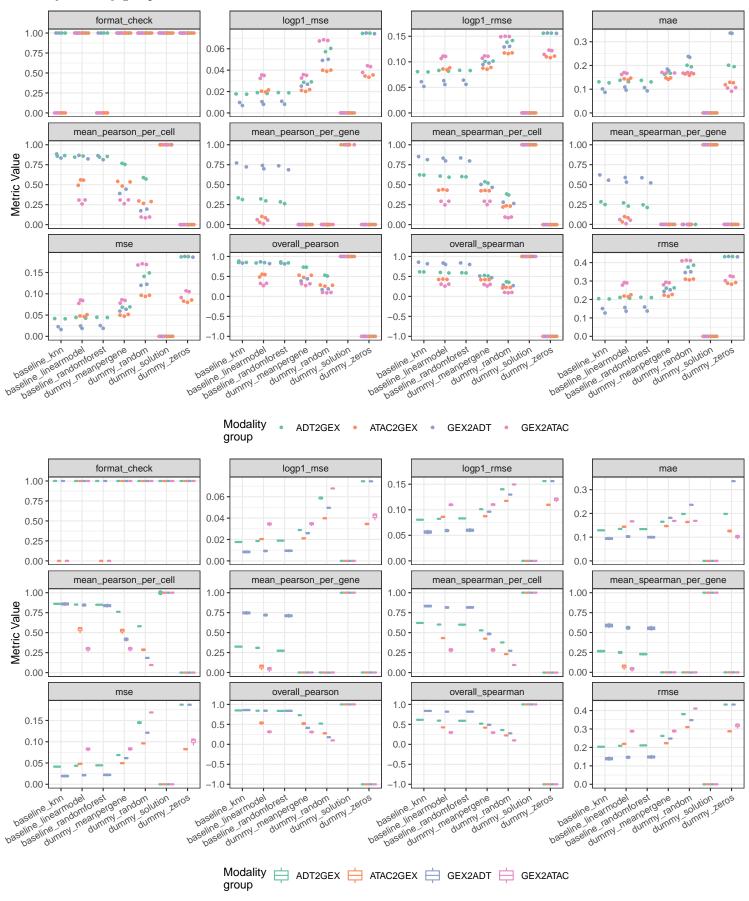
1

Visualise results

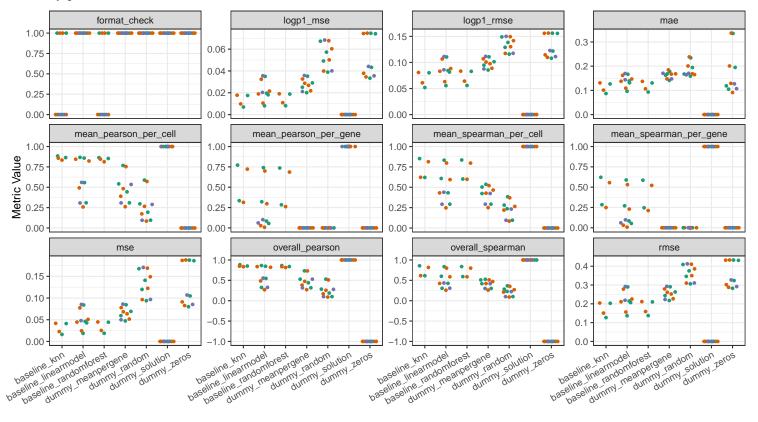
Colour by method.



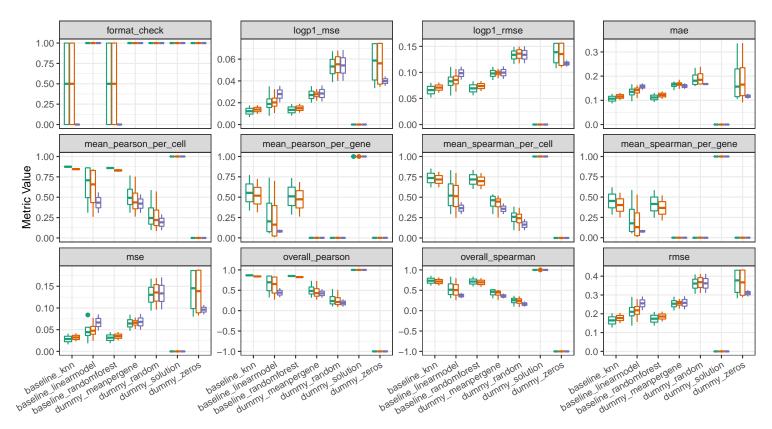
Colour by modality group.



Colour by phase







Phase iid phase1 phase2

Comparing most interesting metrics.

```
selected_metrics <- c("rmse", "mae", "mean_spearman_per_cell", "mean_spearman_per_gene", "overall_spearman")</pre>
GGally::ggpairs(
 df,
 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
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
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```

