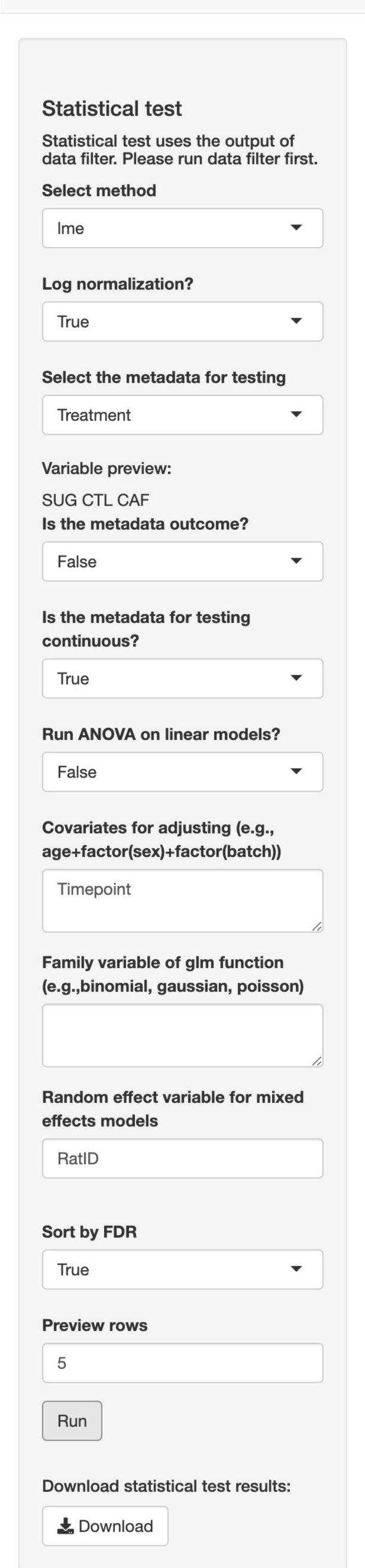
Data input Metadata input MDS plot Alpha diversity plot Taxa barplot Data filter Statistical test Tree plot Boxplot plot Correlation plot P vs P plot



|  | stats            | P                    | FDR                 | Enriched |
|--|------------------|----------------------|---------------------|----------|
| d_Bacteria p_Actinobacteria c_Coriobacteriia o_Coriobacteriales f_uncultured   | 4.17454318101241 | 0.000116516323425059 | 0.00132745382759264 | positive |
| d_Bacteria p_Actinobacteria c_Coriobacteriia o_Coriobacteriales f_uncultured g_uncultured_bacterium                        | 4.17454318101241 | 0.000116516323425059 | 0.00132745382759264 | positive |
| d_Bacteria p_Actinobacteria c_Coriobacteriia o_Coriobacteriales f_uncultured g_uncultured_bacterium s_uncultured_bacterium | 4.17454318101241 | 0.000116516323425059 | 0.00132745382759264 | positive |
| d_Bacteria p_Bacteroidetes c_Bacteroidia o_Bacteroidales f_Prevotellaceae g_Paraprevotella                                 | 4.13807906933272 | 0.000131237481745728 | 0.00138894465740487 | positive |
| d_Bacteria p_Proteobacteria c_Gammaproteobacteria o_Betaproteobacteriales f_Burkholderiaceae                               | 4.12945192888092 | 0.000134975813102041 | 0.00138894465740487 | positive |