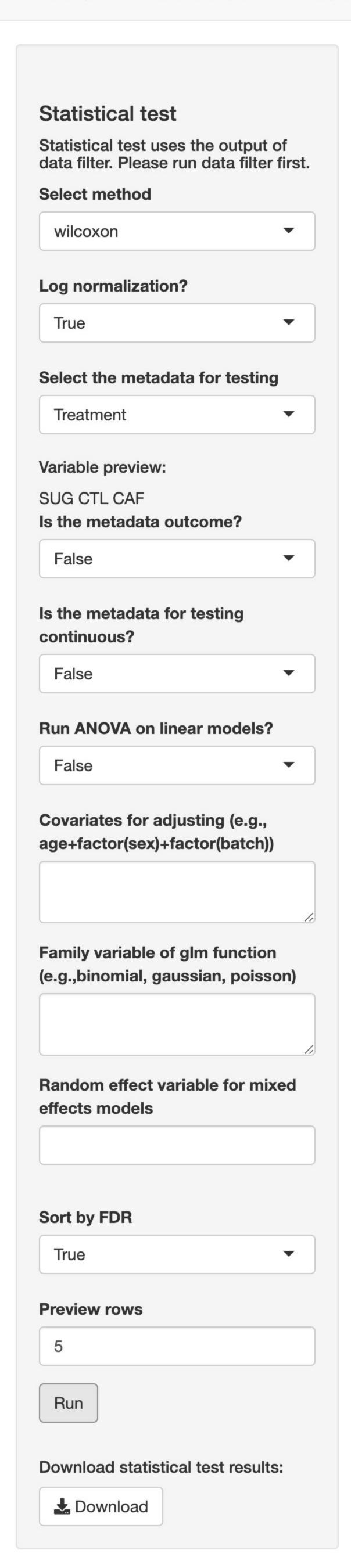
Data input Metadata input MDS plot Alpha diversity plot Taxa barplot Data filter Statistical test Tree plot Boxplot plot Correlation plot P vs P plot



	stats	P	FDR	Enriched
d_Bacteriap_Firmicutes c_Erysipelotrichiao_Erysipelotrichales f_Erysipelotrichaceaeg_Allobaculum s_uncultured_bacterium	571	0.000106041070691981	0.00281892512922849	CAF
d_Bacteriap_Firmicutesc_Clostridia o_Clostridialesf_Clostridiaceae_1 g_Clostridium_sensu_stricto_1	533	0.000161890031297217	0.00368877999884373	CAF
d_Bacteriap_Firmicutesc_Clostridia o_Clostridialesf_Clostridiaceae_1 g_Clostridium_sensu_stricto_1	533	0.000161890031297217	0.00368877999884373	CAF
d_Bacteriap_Firmicutesc_Clostridia o_Clostridialesf_Lachnospiraceae	155	0.000187862783445481	0.00374551424494428	CTL
d_Bacteriap_Firmicutesc_Clostridia o_Clostridialesf_Lachnospiraceae	155	0.000187862783445481	0.00374551424494428	CTL