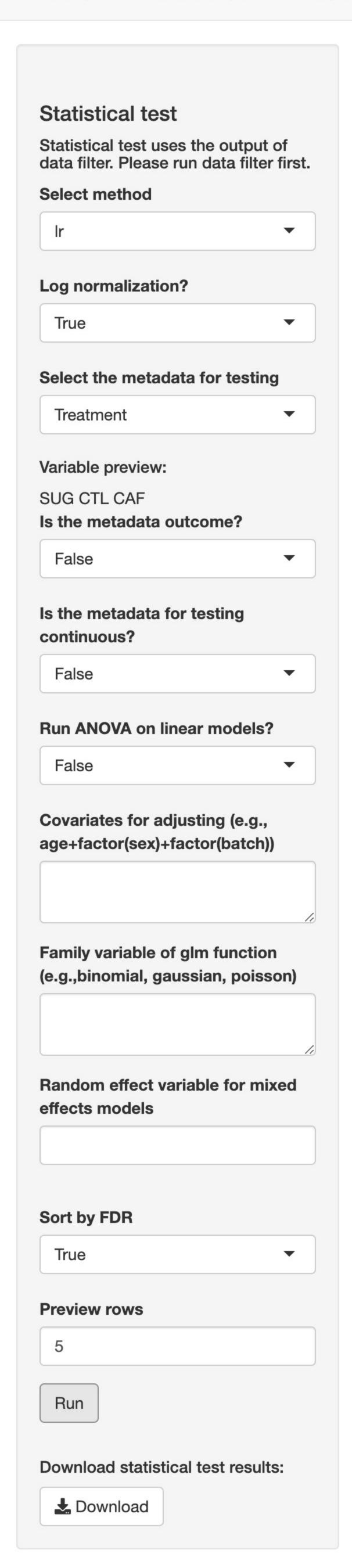
Data input Metadata input MDS plot Alpha diversity plot Taxa barplot Data filter Statistical test Tree plot Boxplot plot Correlation plot P vs P plot



	stats	P	FDR	Enriched
d_Bacteriap_Firmicutes c_Clostridiao_Clostridialesf_Ruminococcaceae g_Ruminiclostridium_6	3.63729726081992	0.000275513900126186	0.0215718924874148	CTL
d_Bacteriap_Firmicutes c_Clostridiao_Clostridialesf_Ruminococcaceae g_Ruminiclostridium_6	3.63729726081992	0.000275513900126186	0.0215718924874148	CTL
d_Bacteriap_Firmicutes c_Clostridiao_Clostridiales- -f_Clostridiaceae_1	-3.33711192440922	0.000846538504279101	0.0215718924874148	CAF
d_Bacteria p_Bacteroidetes c_Bacteroidia o_Bacteroidales f_Prevotellaceae g_Paraprevotella	3.32789568139099	0.000875046265264913	0.0215718924874148	CTL
d_Bacteriap_Firmicutes c_Clostridiao_Clostridialesf_Lachnospiraceae g_Sellimonas	3.32135404973416	0.000895818209184395	0.0215718924874148	CTL