

ANCOM-BC2: Log fold change in abundance. # Sig taxa = 20			
Taxon	lfc_Timepoint7mpf	diff_Timepoint7mpf	direct
Barnesiellaceae_Genus	3.743419	TRUE	Positive LFC
Cloacibacterium	3.179109	TRUE	Positive LFC
Glutamicibacter	2.139606	TRUE	Positive LFC
Mycobacterium	2.047308	TRUE	Positive LFC
Fluviicola	1.865743	TRUE	Positive LFC
Sutterellaceae_Genus	1.699197	TRUE	Positive LFC
Deinococcus	1.684537	TRUE	Positive LFC
Crenobacter	1.587856	TRUE	Positive LFC
Stenotrophomonas	1.387593	TRUE	Positive LFC
Brevibacterium	1.238637	TRUE	Positive LFC
Simkaniaceae_Genus	1.120585	TRUE	Positive LFC
Rhizobiales_Family_Genus	0.958539	TRUE	Positive LFC
Pseudorhodoplanes	0.957715	TRUE	Positive LFC
Smaragdicoccus	0.805331	TRUE	Positive LFC
Gordonia	0.775047	TRUE	Positive LFC
Candidatus_Proteochlamydia	0.746518	TRUE	Positive LFC
Paucibacter	-0.961683	TRUE	Negative LFC
Comamonadaceae_Genus	-1.492398	TRUE	Negative LFC
Plesiomonas	-2.108167	TRUE	Negative LFC
Shewanella	-2.303035	TRUE	Negative LFC