

ANCOM-BC2: Log fold change in abundance. # Sig taxa = 20			
Taxon	lfc_Timepoint7mpf	diff_Timepoint7mpf	direct
Barnesiellaceae_Genus	6.931000	TRUE	Positive LFC
Fluvicola	4.114707	TRUE	Positive LFC
Macellibacteroides	3.981390	TRUE	Positive LFC
Bacteroides	3.457592	TRUE	Positive LFC
Rhizobiales_Family_Genus	1.971942	TRUE	Positive LFC
Mycobacterium	1.896726	TRUE	Positive LFC
Cellvibrio	1.799895	TRUE	Positive LFC
Cloacibacterium	1.488282	TRUE	Positive LFC
Chitinibacteraceae_Genus	1.479288	TRUE	Positive LFC
Sutterellaceae_Genus	1.324464	TRUE	Positive LFC
Simkaniaceae_Genus	1.310535	TRUE	Positive LFC
Bosea	1.259080	TRUE	Positive LFC
Halscomenobacter	1.249192	TRUE	Positive LFC
Smaragdicoccus	1.186623	TRUE	Positive LFC
Bdellovibrio	0.912548	TRUE	Positive LFC
Microscillaceae_Genus	0.912548	TRUE	Positive LFC
Crenobacter	0.737747	TRUE	Positive LFC
Candidatus_Proteochlamydia	0.678023	TRUE	Positive LFC
Phreatobacter	-1.336283	TRUE	Negative LFC
Flavobacterium	-2.000803	TRUE	Negative LFC