

ANCOM-BC2: Summary statistics of abundant taxa (4-7mpf, controls), sig taxa = 35				
Taxon	W	p_val	q_val	diff_abn
Acinetobacter	17.807	0.006	0.009	TRUE
Aeromonas	62.333	0.000	0.000	TRUE
Aurantisolimonas	70.133	0.000	0.000	TRUE
Bacteroides	98.973	0.000	0.000	TRUE
Barnesiellaceae_Genus	295.434	0.000	0.000	TRUE
Bosea	19.242	0.003	0.005	TRUE
Candidatus_Proteochlamydia	14.480	0.026	0.033	TRUE
Cellvibrio	47.047	0.000	0.000	TRUE
Cetobacterium	0.364	0.007	0.010	TRUE
Chitinibacter	742.399	0.000	0.000	TRUE
Chitinibacteraceae_Genus	102.132	0.000	0.000	TRUE
Cloacibacterium	64.301	0.000	0.000	TRUE
Comamonadaceae_Genus	32.364	0.000	0.000	TRUE
Crenobacter	171.541	0.000	0.000	TRUE
Exiguobacterium	14.737	0.023	0.031	TRUE
Flavobacterium	90.168	0.000	0.000	TRUE
Fluviicola	47.091	0.000	0.000	TRUE
Glutamicibacter	49.358	0.000	0.000	TRUE
Gordonia	28.607	0.000	0.000	TRUE
Luteimonas	19.843	0.003	0.004	TRUE
Macellibacteroides	187.360	0.000	0.000	TRUE
Microscillaceae_Genus	21.875	0.001	0.002	TRUE
Mycobacterium	40.130	0.000	0.000	TRUE
Paucibacter	27.598	0.000	0.000	TRUE
Peptostreptococcus	23.689	0.000	0.001	TRUE
Phreatobacter	24.733	0.000	0.001	TRUE
Plesiomonas	52.265	0.000	0.000	TRUE
Pseudomonas	22.598	0.001	0.001	TRUE
Rhizobiales_Family_Genus	64.211	0.000	0.000	TRUE
Rhodobacteraceae_Genus	0.361	0.007	0.010	TRUE
Shewanella	109.984	0.000	0.000	TRUE
Simkaniaceae_Genus	20.060	0.002	0.004	TRUE
Smaragdicroccus	23.591	0.001	0.001	TRUE
Sutterellaceae_Genus	43.357	0.000	0.000	TRUE
ZOR0006	26.897	0.000	0.000	TRUE