New insights into culture negative endophthalmitis by unbiased next generation sequencing

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Supplementary Table 3: Statistical comparison of bacterial genera abundance between culture positive and culture negative endophthalmitis cases

Bacterial genera exhibiting significant (BH corrected $P < 0.05$) differential abundance across culture positive and culture negative endophthalmitis cases

Genus	p value (Wilcoxon test)	BH corrected p value
Acinetobacter	0.353	0.53
Actinomycetales	0.353	0.53
Bacillus	0.165	0.53
Corynebacterium	0.353	0.53
Brevibacterium	0.559	0.789
Enterobacteriaceae	0.203	0.53
Erwinia	0.165	0.53
Gemella	0.638	0.851
Haemophilus	1	1
Klebsiella	0.038	0.344
Microbacterium	1	1
Micrococcus	0.353	0.53
Moraxellaceae	1	1
Morganella	1	1
Planococcaceae	0.97	1
Pseudomonas	1	1
Serratia	0.038	0.344
Sphingomonas	0.3	0.53
Staphylococcus	0.319	0.53
Streptococcus	0.043	0.344
Williamsia	0.096	0.53
Salmonella	0.353	0.53
Stenoptrophomonas	0.353	0.53
Unidentified	0.353	0.53