

Genus	<i>df</i>	logLik	AIC	BIC	deviance	<i>df</i> (residual)	<i>p</i>	<i>R</i> ²
<i>Pelagibacter</i>	2	171.5	-337.1	-325.2	9.1	380	<0.0001	0.126
SAR86A	2	15.2	-24.3	-21.0	0.3	20	0.052	0.135
<i>Litoricola</i>	2	14.3	-22.6	-19.9	0.2	16	0.683	-0.051
Pelagibacter_A	2	18.7	-31.5	-25.2	1.8	57	0.003	0.130
Synechococcus_C	2	2.6	0.8	2.7	0.6	12	0.89	-0.082
<i>Luminiphilus</i>	2	18.2	-30.4	-26.6	0.4	25	0.13	0.053
AG-337-I02	2	11.3	-16.6	-13.4	0.5	20	0.19	0.038