

ProbeName	RefSeq	Species	Description	qPCR	Array	qPCR	Array	qPCR	Array
				0.5 mg/L	0.5 mg/L	2.0 mg/L	2.0 mg/L	8.0 mg/L	8.0 mg/L
Contig03325	XP_001624662	Coral	similar to cytochrome P450	1.83	1.56	0.86	0.91	1.23	1.21
Contig03406	XP_971267	Coral	UDP-glucose 6-dehydrogenase[EC11122]	0.33	-0.29	-0.95	-0.61	-1.51	-1.23
Contig03961	XP_001700637	Algal Symb	pyruvate kinase [EC27140]	0.00	-0.27	-0.88	-0.52	-1.40	-1.27
Contig04866	AAU87579	Algal Symb	major facilitator superfamily MFS_1	0.06	-0.19	-0.58	-0.79	-0.63	-1.32
Contig05643	BAE65110	Other	acyl-CoA synthetase [EC6213]	0.44	-0.22	0.34	-0.79	-0.34	-1.27
Contig06023	XP_001624662	Coral	hypothetical LOC556280	1.71	1.50	1.72	0.93	1.52	1.27
Contig07143	XP_780234	Coral	similar to ENSANGP00000017306	-0.18	-1.97	-0.78	-3.45	-0.16	-2.27
Contig07816	ZP_01750245	Algal Symb	chloroplast translation elongation factor G EF-G [EC3653]	-0.12	-0.22	0.70	-0.59	0.78	-1.18
Contig15230	XP_001625659	Coral	Cytochrome P450 family[EC114--]	0.89	0.5	0.37	0.41	1.14	0.87
Contig16339	XP_001179448	Coral	similar to Pck1 protein [EC41132]	1.44	1.18	0.41	0.46	0.00	-0.2
Contig17964	XP_647496	Other	3-methyl-2-oxobutanoyl-CoA dehydrogenase lipoamide [EC1241]	0.16	-0.17	0.15	-0.76	-2.48	-1.21
Contig25605	AAW79366	Algal Symb	chloroplast light harvesting complex protein	0.29	-0.07	-0.57	-0.65	-0.71	-1.06
Contig28198	XP_001638304	Coral	similar to UDP-glucuronosyltransferase 2 family polypeptideB4	0.78	0.57	0.90	0.71	1.99	1.81

