Supplementary Material S5. List of Gene Ontology terms of the biological process category enriched for the set of upregulated transcripts in gill tissue.

UPREGULATED

term description	term number
biological_process	GO:0008150
positive regulation of cell growth	GO:0030307
carbohydrate metabolic process	GO:0005975
cellular catabolic process	GO:0044248
protein folding	GO:0006457
protein polyubiquitination	GO:0000209
lipid metabolic process	GO:0006629
intracellular transport	GO:0046907
nucleobase-containing compound catabolic process	GO:0034655
proteolysis	GO:0006508
single-organism developmental process	GO:0044767
cellular macromolecular complex assembly	GO:0034622
protein complex subunit organization	GO:0071822
generation of neurons	GO:0048699
vesicle-mediated transport	GO:0016192
response to oxygen-containing compound	GO:1901700
protein complex assembly	GO:0006461
cell cycle	GO:0007049
response to external stimulus	GO:0009605
positive regulation of transcription, DNA-templated	GO:0045893

DOWNREGULATED

term description	term number
microtubule-based process	GO:0007017
energy derivation by oxidation of organic compounds	GO:0015980
heterocycle biosynthetic process	GO:0018130
aromatic compound biosynthetic process	GO:0019438
cellular nitrogen compound biosynthetic process	GO:0044271
organic cyclic compound biosynthetic process	GO:1901362
biological_process	GO:0008150
regulation of signal transduction	GO:0009966
cellular protein modification process	GO:0006464
nucleotide metabolic process	GO:0009117
response to organic substance	GO:0010033
single-organism transport	GO:0044765
cell morphogenesis involved in differentiation	GO:0000904

regulation of multicellular organismal process	GO:0051239
purine-containing compound metabolic process	GO:0072521
single-organism biosynthetic process	GO:0044711
cell surface receptor signaling pathway	GO:0007166
regulation of biological quality	GO:0065008
anatomical structure morphogenesis	GO:0009653
protein modification by small protein conjugation	GO:0032446