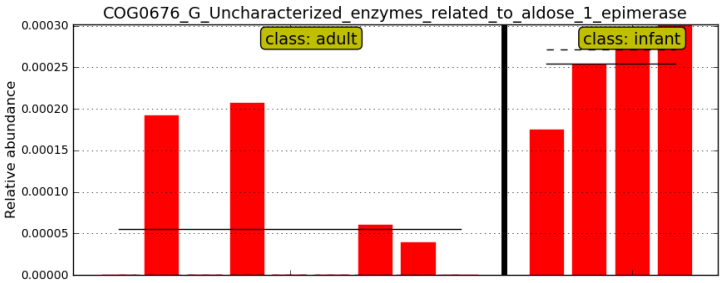
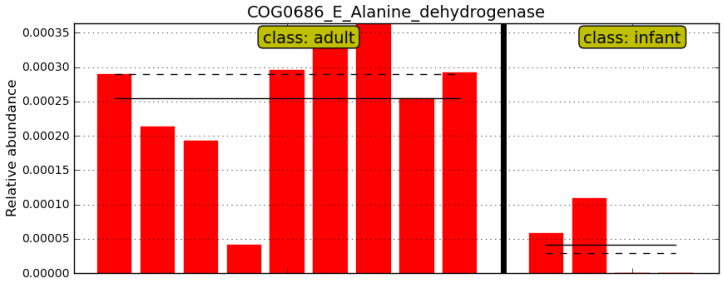
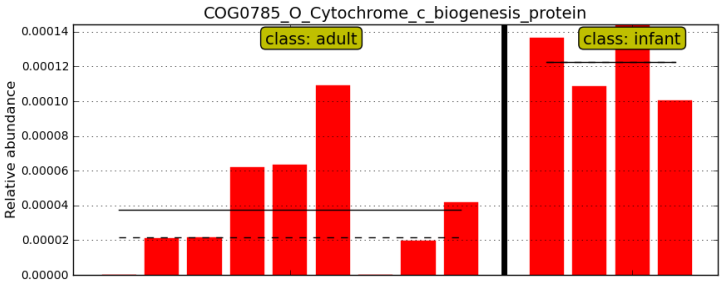
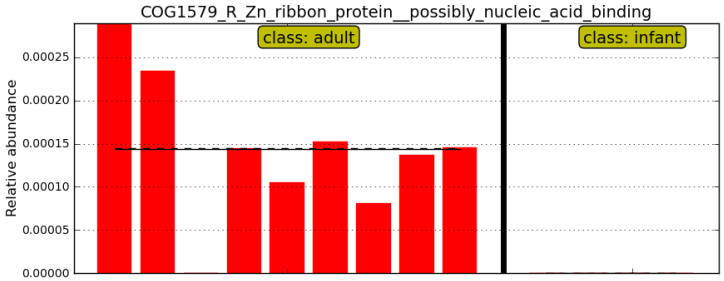
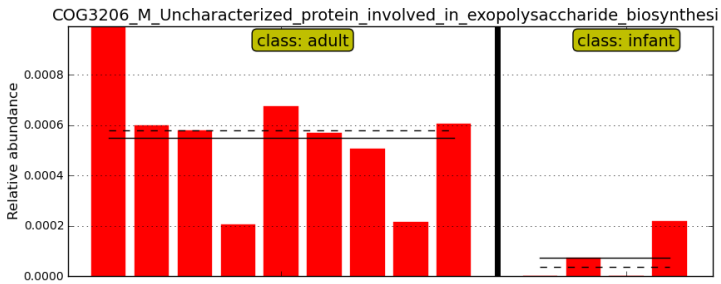
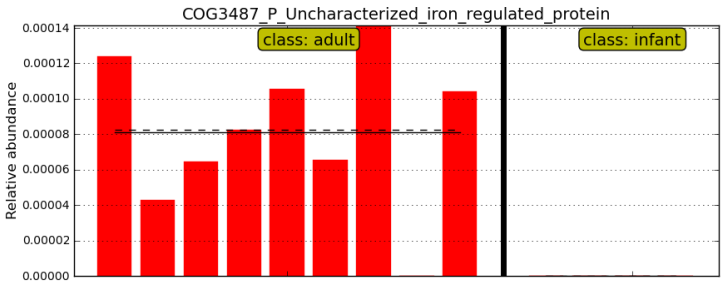

COG0676_G_Uncharacterized_enzymes_related_to_aldose_1_epimerase
class: adult
class: infant
Relative abundance
0.00000
0.00005
0.00010
0.00015
0.00020
0.00025
0.00030

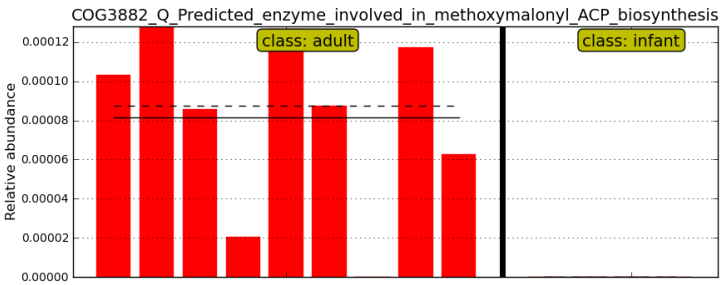

COG0686_E_Alanine_dehydrogenase
class: adult
class: infant
Relative abundance
0.00000
0.00005
0.00010
0.00015
0.00020
0.00025
0.00030
0.00035

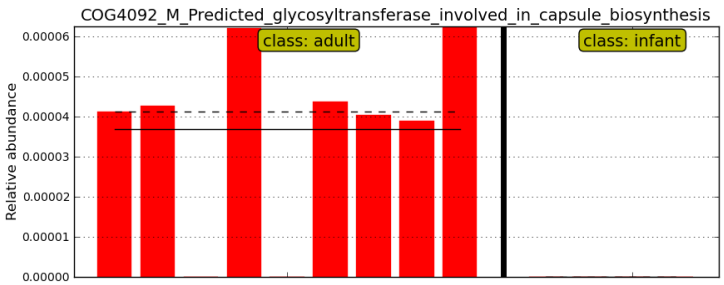

COG0785_O_Cytochrome_c_biogenesis_protein
class: adult
class: infant
Relative abundance
0.00000
0.00002
0.00004
0.00006
0.00008
0.00010
0.00012
0.00014

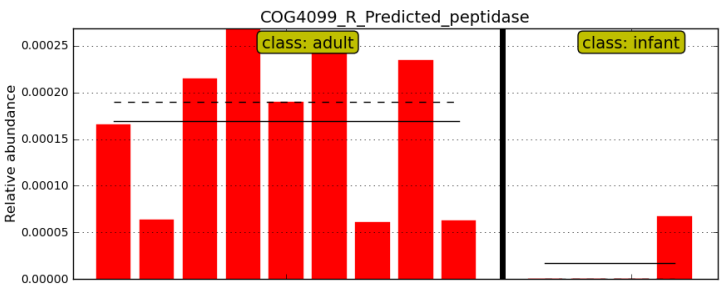

COG1579_R_Zn_ribbon_protein__possibly_nucleic_acid_binding
class: adult
class: infant
Relative abundance
0.00000
0.00005
0.00010
0.00015
0.00020
0.00025

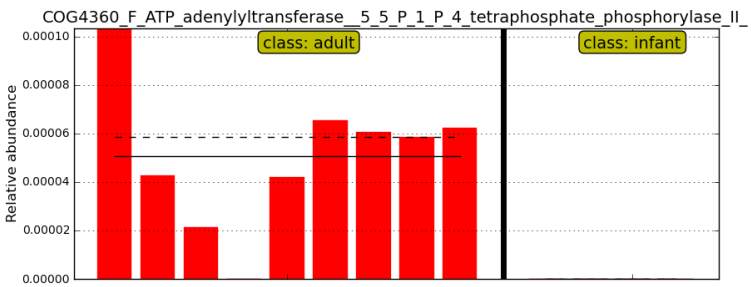

COG3206_M_Uncharacterized_protein_involved_in_exopolysaccharide_biosynthesi
class: adult
class: infant
Relative abundance
0.0000
0.0002
0.0004
0.0006
0.0008

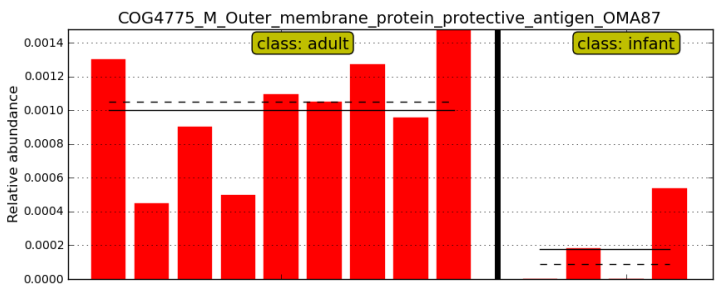

COG3487_P_Uncharacterized_iron_regulated_protein
class: adult
class: infant
Relative abundance
0.00000
0.00002
0.00004
0.00006
0.00008
0.00010
0.00012
0.00014


COG3882_Q_Predicted_enzyme_involved_in_methoxymalonyl_ACP_biosynthesis
class: adult
class: infant
Relative abundance
0.00000
0.00002
0.00004
0.00006
0.00008
0.00010
0.00012


COG4092_M_Predicted_glycosyltransferase_involved_in_capsule_biosynthesis
class: adult
class: infant
Relative abundance
0.00000
0.00001
0.00002
0.00003
0.00004
0.00005
0.00006


COG4099_R_Predicted_peptidase
class: adult
class: infant
Relative abundance
0.00000
0.00005
0.00010
0.00015
0.00020
0.00025


COG4360_F_ATP_adenylyltransferase__5_5_P_1_P_4_tetraphosphate_phosphorylase_II_
class: adult
class: infant
Relative abundance
0.00000
0.00002
0.00004
0.00006
0.00008
0.00010


COG4775_M_Outer_membrane_protein_protective_antigen_OMA87
class: adult
class: infant
Relative abundance
0.0000
0.0002
0.0004
0.0006
0.0008
0.0010
0.0012
0.0014