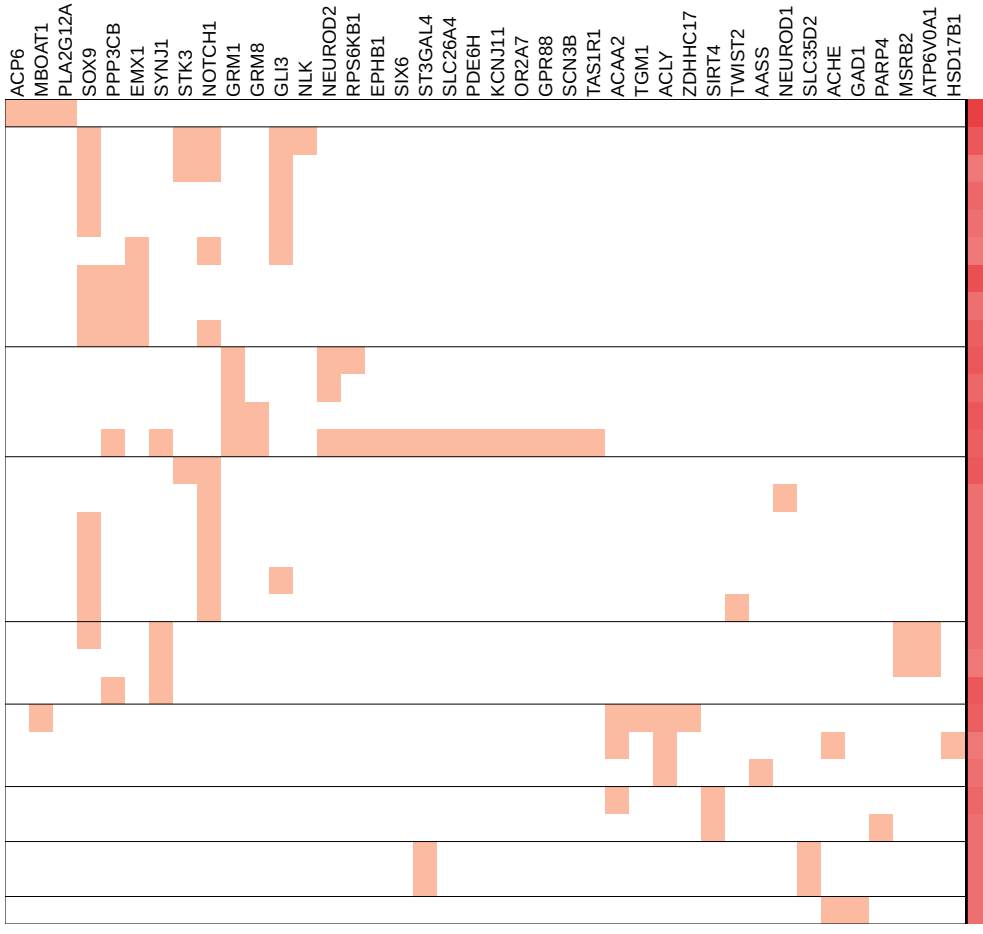

5.44e−03	GO_ALDITOL_PHOSPHATE_METABOLIC_PROCESS
1.39e−02	GO_NEGATIVE_REGULATION_OF_WNT_SIGNALING_PATHWAY
3.24e−02	GO_NEGATIVE_REGULATION_OF_CANONICAL_WNT_SIGNALING_PATHWAY
1.99e−02	GO_POSITIVE_REGULATION_OF_CHONDROCYTE_DIFFERENTIATION
2.72e−02	GO_POSITIVE_REGULATION_OF_CARTILAGE_DEVELOPMENT
3.42e−02	GO_NEUROEPITHELIAL_CELL_DIFFERENTIATION
9.44e−03	GO_DEVELOPMENTAL_CELL_GROWTH
2.96e−02	GO_CELL_GROWTH
1.60e−02	GO_DEVELOPMENTAL_GROWTH_INVOLVED_IN_MORPHOGENESIS
1.48e−02	GO_RESPONSE_TO_ELECTRICAL_STIMULUS
1.99e−02	GO_CELLULAR_RESPONSE_TO_ELECTRICAL_STIMULUS
1.36e−02	GO_G_PROTEIN_COUPLED_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY
1.67e−02	GO_NEUROLOGICAL_SYSTEM_PROCESS
1.36e−02	GO_ENDOCARDIUM_DEVELOPMENT
2.72e−02	GO_EPITHELIAL_CELL_FATE_COMMITMENT
2.72e−02	GO_REGULATION_OF_CELL_PROLIFERATION_INVOLVED_IN_HEART_MORPHOGENESIS
2.72e−02	GO_REGULATION_OF_EXTRACELLULAR_MATRIX_ORGANIZATION
2.96e−02	GO_OLIGODENDROCYTE_DIFFERENTIATION
2.54e−02	GO_NEGATIVE_REGULATION_OF_OSSIFICATION
2.92e−02	GO_CELLULAR_MACROMOLECULAR_COMPLEX_ASSEMBLY
3.42e−02	GO_CELLULAR_PROTEIN_COMPLEX_ASSEMBLY
1.36e−02	GO_CALCIUM_ION_REGULATED_EXOCYTOSIS
1.71e−02	GO_TRANSFERASE_ACTIVITY_TRANSFERRING_ACYL_GROUPS
3.24e−02	GO_ALCOHOL_BIOSYNTHETIC_PROCESS
2.72e−02	GO_ACETYL_COA_METABOLIC_PROCESS
1.99e−02	GO_NEGATIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION
2.72e−02	GO_PROTEIN_ADP_RIBOSYLATION
2.72e−02	GO_KERATAN_SULFATE_BIOSYNTHETIC_PROCESS
2.72e−02	GO_KERATAN_SULFATE_METABOLIC_PROCESS
2.72e−02	GO_NEUROTRANSMITTER_BIOSYNTHETIC_PROCESS