

HSD17B1	ACHE	GAD1	NEUROD1	AASS	TWIST2	SIRT4	PARP4	TGM1	ZDHC17	ACAA2	ACLY	ST3GAL4	SLC35D2	RPS6KB1	GRM1	NEUROD2	GRM8	TAS1R1	SCN3B	GPR88	OR2A7	KCNJ11	PDE6H	SLC26A4	EPHB1	SIX6	SYNJ1	MSRB2	ATP6V0A1	PPP3CB	EMX1	MBOAT1	ACP6	PLA2G12A	GLI3	SOX9	NOTCH1	STK3	NLK			
																																									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_CALCIIUM_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_ADG_RIBOSYLATION
																																									2.72e-02	GO_KERATAN_SULFATE_BIOSYNTHETIC_PROCESS
																																									2.72e-02	GO_KERATAN_SULFATE_METABOLIC_PROCESS
																																									2.72e-02	GO_NEUROTRANSMITTER_BIOSYNTHETIC_PROCESS