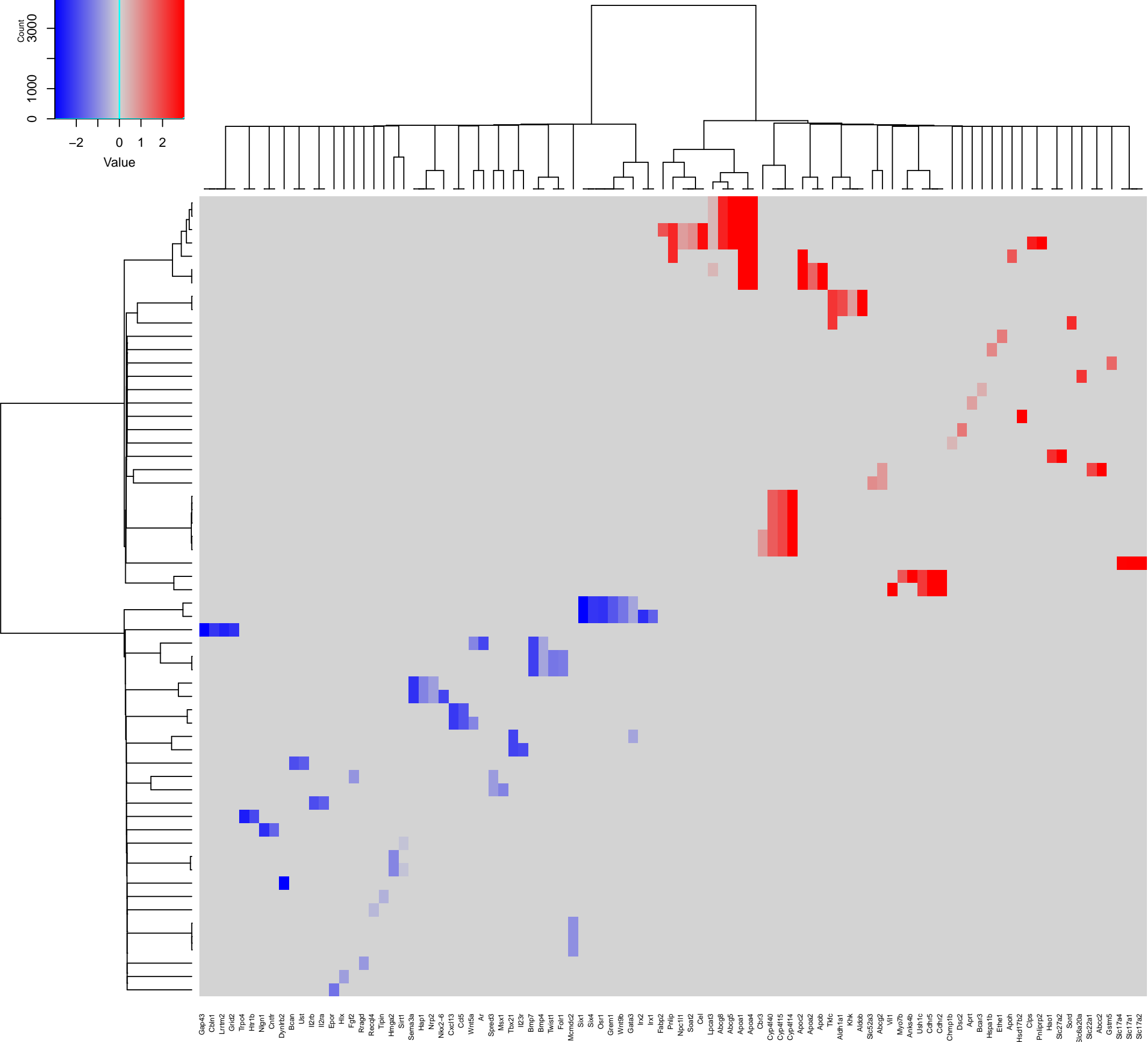


Snyder_All_C5.gmt



- GOBP_REGULATION_OF_INTESTINAL_CHOLESTEROL_ABSORPTION
- GOBP_REGULATION_OF_INTESTINAL_ABSORPTION
- GOBP_INTESTINAL_LIPID_ABSORPTION
- GOBP_LIPID_DIGESTION
- GOBP_POSITIVE_REGULATION_OF_TRIGLYCERIDE_LIPASE_ACTIVATION
- GOBP_CHYLOMICRON_ASSEMBLY
- GOBP_CHYLOMICRON_REMODELING
- GOBP_GLYCERALDEHYDE_3_PHOSPHATE_METABOLIC_PROCESS
- GOBP_FRUCTOSE_CATABOLIC_PROCESS_TO_HYDROXYACETONE
- GOBP_ALDITOL_CATABOLIC_PROCESS
- GOBP_SULFIDE_OXIDATION
- GOBP_HEAT_ACCLIMATION
- GOBP_DETOXIFICATION_OF_NITROGEN_COMPOUND
- GOBP_AMINO_ACID_BETAINE_TRANSPORT
- GOBP_ENDOTHELIN_RECEPTOR_SIGNALING_PATHWAY
- GOBP_PURINE_NUCLEOTIDE_SALVAGE
- GOBP_ESTROGEN_BIOSYNTHETIC_PROCESS
- GOBP_CARDIAC_MUSCLE_CELL_CARDIAC_MUSCLE_CELL_ADHESION
- GOBP_ESCRT_COMPLEX_DISASSEMBLY
- GOBP_FATTY_ACID_ALPHA_OXIDATION
- GOBP_XENOBIOTIC_TRANSPORT_ACROSS_BLOOD_BRAIN_BARRIER
- GOBP_RIBOFLAVIN_TRANSPORT
- GOBP_LEUKOTRIENE_B4_METABOLIC_PROCESS
- GOBP_LEUKOTRIENE_B4_CATABOLIC_PROCESS
- GOBP_ICOSANOID_CATABOLIC_PROCESS
- GOBP_KETONE_CATABOLIC_PROCESS
- GOBP_PHYLLOQUINONE_METABOLIC_PROCESS
- GOBP_SIALIC_ACID_TRANSPORT
- GOBP_BRUSH_BORDER_ASSEMBLY
- GOBP_REGULATION_OF_MICROVILLUS_LENGTH
- GOBP_MESONEPHRIC_TUBULE_FORMATION
- GOBP_NEPHRON_TUBULE_FORMATION
- GOBP_REGULATION_OF_POSTSYNAPTIC_SPECIALIZATION_ASSEMBLY
- GOBP_LATERAL_SPROUTING_FROM_AN_EPITHELIUM
- GOBP_CELL_MIGRATION_INVOLVED_IN_HEART_DEVELOPMENT
- GOBP_CARDIAC_NEURAL_CREST_CELL_MIGRATION_INVOLVED_IN
- GOBP_HYPOTHALAMUS_CELL_DIFFERENTIATION
- GOBP_HYPOTHALAMUS_DEVELOPMENT
- GOBP_POSITIVE_REGULATION_OF_CELL_CELL_ADHESION_MEDIATED
- GOBP_REGULATION_OF_T_CELL_CHEMOTAXIS
- GOBP_T_HELPER_2_CELL_CYTOKINE_PRODUCTION
- GOBP_REGULATION_OF_T_HELPER_17_CELL_LINEAGE_COMMITMENT
- GOBP_DERMATAN_SULFATE_METABOLIC_PROCESS
- GOBP_REGULATION_OF_LENS_FIBER_CELL_DIFFERENTIATION
- GOBP_POSITIVE_REGULATION_OF_DNA_DAMAGE_RESPONSE_SIGNALING
- GOBP_RESPONSE_TO_INTERLEUKIN_2
- GOBP_GAMMA_AMINO BUTYRIC ACID SECRETION
- GOBP_BRAINSTEM_DEVELOPMENT
- GOBP_HETEROCHROMATIN_ORGANIZATION_INVOLVED_IN_CHROMATIN
- GOBP_POSITIVE_REGULATION_OF_CELL_PROLIFERATION_IN_BONE
- GOBP_POSITIVE_REGULATION_OF_CELL_AGING
- GOBP_INTRACILIARY_TRANSPORT_INVOLVED_IN_CILIUM_ASSEMBLY
- GOBP_REPLICATION_FORK_PROTECTION
- GOBP_TELOMERIC_LOOP_DISASSEMBLY
- GOBP_DNA_STRAND_ELONGATION_INVOLVED_IN_DNA_REPLICATION
- GOBP_DOUBLE_STRAND_BREAK_REPAIR_VIA_BREAK_INDUCED
- GOBP_PRE_REPLICATIVE_COMPLEX_ASSEMBLY_INVOLVED_IN_CELL
- GOBP_CELLULAR_RESPONSE_TO_LEUCINE
- GOBP_POSITIVE_REGULATION_OF_T_HELPER_1_CELL_DIFFERENTIATION
- GOBP_RESPONSE_TO_ERYTHROPOIETIN