

B_Cell	T_Cell	Monocyte	<div> <div></div> <div>-3</div> <div>Enrichment score</div> <div>3</div> </div>		
					HSA_M00177_RIBOSOME_EUKARYOTES
					HSA_M00352_SPLICEOSOME_U2_SNRNP
					HSA_M00354_SPLICEOSOME_U4_U6_U5_TRI_SNRNP
					HSA_M00426_SURVIVAL_MOTOR_NEURON_SMN_COMPLEX
					HSA_M00427_NUCLEAR_PORE_COMPLEX
					HSA_M00341_PROTEASOME_19S_REGULATORY_PARTICLE_PA700_
					HSA_M00398_SM_CORE_COMPLEX
					HSA_M00351_SPLICEOSOME_U1_SNRNP
					HSA_M00391_EXOSOME_EUKARYOTES
					HSA_M00289_RF_C_COMPLEX
					HSA_M00143_NADH_DEHYDROGENASE_UBIQUINONE_FE_S_PROTEIN_FLAVOPROTEIN_COMPLEX_MITOCHONDRIA
					HSA_M00146_NADH_DEHYDROGENASE_UBIQUINONE_1_ALPHA_SUBCOMPLEX
					HSA_M00414_BLOOM_S_SYNDROME_COMPLEX
					HSA_M00396_LSM_2_8_COMPLEX
					HSA_M00383_ECV_COMPLEX
					HSA_M00154_CYTOCHROME_C_OXIDASE
					HSA_M00128_UBIQUINONE_BIOSYNTHESIS_EUKARYOTES_4_HYDROXYBENZOATE_UBIQUINONE
					HSA_M00098_ACYLGLYCEROL_DEGRADATION
					HSA_M00397_LSM_1_7_COMPLEX
					HSA_M00412_ESCRT_III_COMPLEX
					HSA_M00296_BER_COMPLEX
					HSA_M00389_APC_C_COMPLEX
					HSA_M00056_O_GLYCAN_BIOSYNTHESIS_MUCIN_TYPE_CORE
					HSA_M00403_HRD1_SEL1_ERAD_COMPLEX
					HSA_M00395_DECAPPING_COMPLEX
					HSA_M00048_INOSINE_MONOPHOSPHATE_BIOSYNTHESIS_PRPP_GLUTAMINE_IMP
					HSA_M00359_AMINOACYL_TRNA_BIOSYNTHESIS_EUKARYOTES
					HSA_M00014_GLUCURONATE_PATHWAY_URONATE_PATHWAY_
					HSA_M00340_PROTEASOME_20S_CORE_PARTICLE
					HSA_M00055_N_GLYCAN_PRECURSOR_BIOSYNTHESIS
					HSA_M00307_PYRUVATE_OXIDATION_PYRUVATE_ACETYL_COA
					HSA_M00694_CGMP_SIGNALING
					HSA_M00037_MELATONIN_BIOSYNTHESIS_TRYPTOPHAN_SEROTONIN_MELATONIN
					HSA_M00285_MCM_COMPLEX
					HSA_M00152_CYTOCHROME_BC1_COMPLEX
					HSA_M00181_RNA_POLYMERASE_III_EUKARYOTES
					HSA_M00413_FA_CORE_COMPLEX
					HSA_M00147_NADH_DEHYDROGENASE_UBIQUINONE_1_BETA_SUBCOMPLEX
					HSA_M00036_LEUCINE_DEGRADATION_LEUCINE_ACETOACETATE_ACETYL_COA
					HSA_M00011_CITRATE_CYCLE_SECOND_CARBON_OXIDATION_2_OXOGLUTARATE_OXALOACETATE
					HSA_M00160_V_TYPE_ATPASE_EUKARYOTES
					HSA_M00284_ORIGIN_RECOGNITION_COMPLEX
					HSA_M00295_BRCA1_ASSOCIATED_GENOME_SURVEILLANCE_COMPLEX_BASIC
					HSA_M00101_CHOLESTEROL_BIOSYNTHESIS_SQUALENE_2_3_EPOXIDE_CHOLESTEROL
					HSA_M00071_GLYCOSPHINGOLIPID_BIOSYNTHESIS_NEOLACTO_SERIES_LACER_NLC4CER
					HSA_M00052_PYRIMIDINE_RIBONUCLEOTIDE_BIOSYNTHESIS_UMP_UDP_UTP_CDP_CTP
					HSA_M00382_SCF_FBS_COMPLEX
					HSA_M00158_F_TYPE_ATPASE_EUKARYOTES
					HSA_M00388_ECS_COMPLEX
					HSA_M00007_PENTOSE_PHOSPHATE_PATHWAY_NON_OXIDATIVE_PHASE_FRUCTOSE_6P_RIBOSE_5P
					HSA_M00337_IMMUNOPROTEASOME
					HSA_M00679_BMP_SIGNALING
					HSA_M00009_CITRATE_CYCLE_TCA_CYCLE_KREBS_CYCLE_
					HSA_M00078_HEPARAN_SULFATE_DEGRADATION
					HSA_M00077_CHONDROITIN_SULFATE_DEGRADATION
					HSA_M00076_DERMATAN_SULFATE_DEGRADATION
					HSA_M00104_BILE_ACID_BIOSYNTHESIS_CHOLESTEROL_CHOLATE_CHENODEOXYCHOLATE
					HSA_M00044_TYROSINE_DEGRADATION_TYROSINE_HOMOGENTISATE
					HSA_M00085_FATTY_ACID_BIOSYNTHESIS_ELONGATION_MITOCHONDRIA
					HSA_M00677_WNT_SIGNALING
					HSA_M00010_CITRATE_CYCLE_FIRST_CARBON_OXIDATION_OXALOACETATE_2_OXOGLUTARATE
					HSA_M00355_SPLICEOSOME_35S_U5_SNRNP
					HSA_M00385_CUL4_DDB1_DDB2_COMPLEX
					HSA_M00386_CUL4_DDB1_CSA_COMPLEX
					HSA_M00428_EIF4F_COMPLEX
					HSA_M00034_METHIONINE_SALVAGE_PATHWAY
					HSA_M00004_PENTOSE_PHOSPHATE_PATHWAY_PENTOSE_PHOSPHATE_CYCLE_
					HSA_M00109_C21_STEROID_HORMONE_BIOSYNTHESIS_PROGESTERONE_CORTISOL_CORTISONE
					HSA_M00087_BETA_OXIDATION
					HSA_M00095_C5_ISOPRENOID_BIOSYNTHESIS_MEVALONATE_PATHWAY
					HSA_M00075_N_GLYCAN_BIOSYNTHESIS_COMPLEX_TYPE
					HSA_M00683_HIPPO_SIGNALING
					HSA_M00042_CATECHOLAMINE_BIOSYNTHESIS_TYROSINE_DOPAMINE_NORADRENALINE_ADRENALINE
					HSA_M00079 KERATAN_SULFATE_DEGRADATION
					HSA_M00131_INOSITOL_PHOSPHATE_METABOLISM_INS_1_3_4_5_P4_INS_1_3_4_P3_MYO_INOSITOL
					HSA_M00073_N_GLYCAN_PRECURSOR_TRIMMING
					HSA_M00692_CELL_CYCLE
					HSA_M00695_CAMP_SIGNALING
					HSA_M00013_MALONATE_SEMIALDEHYDE_PATHWAY_PROPYNOYL_COA_ACETYL_COA
					HSA_M00361_NUCLEOTIDE_SUGAR_BIOSYNTHESIS_EUKARYOTES
					HSA_M00049_ADENINE_RIBONUCLEOTIDE_BIOSYNTHESIS_IMP_ADP_ATP
					HSA_M00050_GUANINE_RIBONUCLEOTIDE_BIOSYNTHESIS_IMP_GDP_GTP
					HSA_M00135_GABA_BIOSYNTHESIS_EUKARYOTES_PUTRESCINE_GABA
					HSA_M00047_CREATINE_PATHWAY
					HSA_M00059_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_HEPARAN_SULFATE_BACKBONE
					HSA_M00003_GLUCONEOGENESIS_OXALOACETATE_FRUCTOSE_6P
					HSA_M00094_CERAMIDE_BIOSYNTHESIS
					HSA_M00032_LYSINE_DEGRADATION_LYSINE_SACCHAROPINE_ACETOACETYL_COA
					HSA_M00415_FATTY_ACID_BIOSYNTHESIS_ELONGATION_ENDOPLASMIC_RETICULUM
					HSA_M00130_INOSITOL_PHOSPHATE_METABOLISM_PI_PIP2_INS_1_4_5_P3_INS_1_3_4_5_P4
					HSA_M00035_METHIONINE_DEGRADATION
					HSA_M00678_HEDGEHOG_SIGNALING
					HSA_M00684_JAK_STAT_SIGNALING
					HSA_M00099_SPHINGOSINE_BIOSYNTHESIS
					HSA_M00002_GLYCOLYSIS_CORE_MODULE_INVOLVING_THREE_CARBON_COMPOUNDS
					HSA_M00549_NUCLEOTIDE_SUGAR_BIOSYNTHESIS_GLUCOSE_UDP_GLUCOSE
					HSA_M00688_MAPK_JNK_SIGNALING
					HSA_M00001_GLYCOLYSIS_EMBDEN_MEYERHOF_PATHWAY_GLUCOSE_PYRUVATE
					HSA_M00689_MAPK_P38_SIGNALING
					HSA_M00353_SPLICEOSOME_PRP19_CDC5L_COMPLEX