Cell	Cell	Monocyte	
BC		ouo	
		Š	⁻³ Enrichment score ³
			751604_GO_0044391_RIBOSOMAL_SUBUNIT 499360_GO_0022626_CYTOSOLIC_RIBOSOME
			490649_GO_0005840_RIBOSOME 496055_GO_0015934_LARGE_RIBOSOMAL_SUBUNIT
			486711_GO_0022625_CYTOSOLIC_LARGE_RIBOSOMAL_SUBUNIT 496056_GO_0015935_SMALL_RIBOSOMAL_SUBUNIT
			486712_GO_0022627_CYTOSOLIC_SMALL_RIBOSOMAL_SUBUNIT
			503919_GO_0034719_SMN_SM_PROTEIN_COMPLEX 499986_GO_0030684_PRERIBOSOME
			507393_GO_0044445_CYTOSOLIC_PART 489275_GO_0030532_SMALL_NUCLEAR_RIBONUCLEOPROTEIN_COMPLEX
			490794_GO_0005761_MITOCHONDRIAL_RIBOSOME 501277_GO_0032040_SMALL_SUBUNIT_PROCESSOME
			911352_GO_0097525_SPLICEOSOMAL_SNRNP_COMPLEX 486830_GO_0000313_ORGANELLAR_RIBOSOME
			495325_GO_0015030_CAJAL_BODY
			490735_GO_0005681_SPLICEOSOMAL_COMPLEX 515939_GO_0071013_CATALYTIC_STEP_2_SPLICEOSOME
			490796_GO_0005763_MITOCHONDRIAL_SMALL_RIBOSOMAL_SUBUNIT 490795_GO_0005762_MITOCHONDRIAL_LARGE_RIBOSOMAL_SUBUNIT
			486832_GO_0000315_ORGANELLAR_LARGE_RIBOSOMAL_SUBUNIT 486831_GO_0000314_ORGANELLAR_SMALL_RIBOSOMAL_SUBUNIT
			502020_GO_0032797_SMN_COMPLEX 499988_GO_0030686_90S_PRERIBOSOME
			507400_GO_0044452_NUCLEOLAR_PART
			911332_GO_0097504_GEMINI_OF_COILED_BODIES 499359_GO_0022624_PROTEASOME_ACCESSORY_COMPLEX
			490739_GO_0005685_U1_SNRNP 490728_GO_0005672_TRANSCRIPTION_FACTOR_TFIIA_COMPLEX
			500369_GO_0031082_BLOC_COMPLEX 490858_GO_0005838_PROTEASOME_REGULATORY_PARTICLE
			486709_GO_0000178_EXOSOME_RNASE_COMPLEX_ 911353_GO_0097526_SPLICEOSOMAL_TRI_SNRNP_COMPLEX
			490743_GO_0005689_U12_TYPE_SPLICEOSOMAL_COMPLEX 503909_GO_0034709_METHYLOSOME
			490714_GO_0005655_NUCLEOLAR_RIBONUCLEASE_P_COMPLEX 499979_GO_0030677_RIBONUCLEASE_P_COMPLEX
			508797_GO_0046540_U4_U6_X_U5_TRI_SNRNP_COMPLEX
			499983_GO_0030681_MULTIMERIC_RIBONUCLEASE_P_COMPLEX 490777_GO_0005742_MITOCHONDRIAL_OUTER_MEMBRANE_TRANSLOCASE_COMPLEX
			517572_GO_0072669_TRNA_SPLICING_LIGASE_COMPLEX 503915_GO_0034715_PICLN_SM_PROTEIN_COMPLEX
			486707_GO_0000176_NUCLEAR_EXOSOME_RNASE_COMPLEX_ 490740_GO_0005686_U2_SNRNP
			500370_GO_0031083_BLOC_1_COMPLEX 500257_GO_0030964_NADH_DEHYDROGENASE_COMPLEX
			490782_GO_0005747_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_I 490814_GO_0005786_SIGNAL_RECOGNITION_PARTICLE_ENDOPLASMIC_RETICULUM_TARGETING
			510573_GO_0048500_SIGNAL_RECOGNITION_PARTICLE 500367_GO_0031080_NUCLEAR_PORE_OUTER_RING
			490737_GO_0005683_U7_SNRNP 490741_GO_0005687_U4_SNRNP
			490736_GO_0005682_U5_SNRNP 507636_GO_0045271_RESPIRATORY_CHAIN_COMPLEX_I
			490855_GO_0005833_HEMOGLOBIN_COMPLEX
			490778_GO_0005743_MITOCHONDRIAL_INNER_MEMBRANE 490824_GO_0005796_GOLGI_LUMEN
			490792_GO_0005759_MITOCHONDRIAL_MATRIX 500378_GO_0031091_PLATELET_ALPHA_GRANULE
			516606_GO_0071682_ENDOCYTIC_VESICLE_LUMEN 500380_GO_0031093_PLATELET_ALPHA_GRANULE_LUMEN
			503973_GO_0034774_SECRETORY_GRANULE_LUMEN 487622_GO_0005882_INTERMEDIATE_FILAMENT
			507495_GO_0045095_KERATIN_FILAMENT 495323_GO_0031225_ANCHORED_COMPONENT_OF_MEMBRANE
			492644_GO_0008328_IONOTROPIC_GLUTAMATE_RECEPTOR_COMPLEX 490653_GO_0005578_PROTEINACEOUS_EXTRACELLULAR_MATRIX
			500302_GO_0031012_EXTRACELLULAR_MATRIX
			490924_GO_0005925_FOCAL_ADHESION 490923_GO_0005924_CELL_SUBSTRATE_ADHERENS_JUNCTION
			499503_GO_0030055_CELL_SUBSTRATE_JUNCTION 490911_GO_0005912_ADHERENS_JUNCTION
			515098_GO_0070161_ANCHORING_JUNCTION 505281_GO_0042105_ALPHA_BETA_T_CELL_RECEPTOR_COMPLEX
			503902_GO_0034702_ION_CHANNEL_COMPLEX 982594_GO_1990351_TRANSPORTER_COMPLEX
			499432_GO_0042101_T_CELL_RECEPTOR_COMPLEX 911483_GO_1902495_TRANSMEMBRANE_TRANSPORTER_COMPLEX
			487787_GO_001891_PHAGOCYTIC_CUP 490903_GO_0005901_CAVEOLA
			752880_GO_0097381_PHOTORECEPTOR_DISC_MEMBRANE
			490893_GO_0005891_VOLTAGE_GATED_CALCIUM_CHANNEL_COMPLEX 492527_GO_0008180_COP9_SIGNALOSOME
			487083_GO_0000790_NUCLEAR_CHROMATIN 487822_GO_0001931_UROPOD
			500513_GO_0031254_CELL_TRAILING_EDGE 496363_GO_0016323_BASOLATERAL_PLASMA_MEMBRANE
			751048_GO_0036128_CATSPER_COMPLEX 507512_GO_0045121_MEMBRANE_RAFT
			507297_GO_0044291_CELL_CELL_CONTACT_ZONE 505527_GO_0042383_SARCOLEMMA
			499715_GO_0030315_T_TUBULE 495152_GO_0014704_INTERCALATED_DISC
			911320_GO_0044815_DNA_PACKAGING_COMPLEX
			850015_GO_1990104_DNA_BENDING_COMPLEX 487079_GO_0000786_NUCLEOSOME
			487081_GO_0000788_NUCLEAR_NUCLEOSOME 490892_GO_0005890_SODIUM_POTASSIUM_EXCHANGING_ATPASE_COMPLEX
			911391_GO_0098533_ATPASE_DEPENDENT_TRANSMEMBRANE_TRANSPORT_COMPLEX 752590_GO_0090533_CATION_TRANSPORTING_ATPASE_COMPLEX
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