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Cell	Cell) ()	
മ	⊢'	Monocyte	⁻³ Enrichment score
			497960_GO_0019083_VIRAL_TRANSCRIPTION
			491334_GO_0006414_TRANSLATIONAL_ELONGATION 491508_GO_0006614_SRP_DEPENDENT_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE 491507_GO_0006613_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE
			507448_GO_0045047_PROTEIN_TARGETING_TO_ER 491335_GO_0006415_TRANSLATIONAL_TERMINATION
			517502_GO_0072599_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM 497957_GO_0019080_VIRAL_GENE_EXPRESSION 507049_GO_0044033_MULTI_ORGANISM_METABOLIC_PROCESS
			491333_GO_0006413_TRANSLATIONAL_INITIATION 515899_GO_0070972_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM
			486715_GO_0000184_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_NONSENSE_MEDIATED_DECAY 486555_GO_0042254_RIBOSOME_BIOGENESIS 491332_GO_0006412_TRANSLATION
			506655_GO_0043624_CELLULAR_PROTEIN_COMPLEX_DISASSEMBLY 487180_GO_0000956_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS 496166_GO_0016072_RRNA_METABOLIC_PROCESS
			499348_GO_0022613_RIBONUCLEOPROTEIN_COMPLEX_BIOGENESIS 505426_GO_0042274_RIBOSOMAL_SMALL_SUBUNIT_BIOGENESIS
			491301_GO_0006364_RRNA_PROCESSING 502206_GO_0032984_MACROMOLECULAR_COMPLEX_DISASSEMBLY 503672_GO_0034470_NCRNA_PROCESSING
			506308_GO_0043241_PROTEIN_COMPLEX_DISASSEMBLY 515063_GO_0070125_MITOCHONDRIAL_TRANSLATIONAL_ELONGATION
			503860_GO_0034660_NCRNA_METABOLIC_PROCESS 501773_GO_0032543_MITOCHONDRIAL_TRANSLATION 515064_GO_0070126_MITOCHONDRIAL_TRANSLATIONAL_TERMINATION
			491506_GO_0006612_PROTEIN_TARGETING_TO_MEMBRANE 515062_GO_0070124_MITOCHONDRIAL_TRANSLATIONAL_INITIATION
			499833_GO_0030490_MATURATION_OF_SSU_RRNA 505425_GO_0042273_RIBOSOMAL_LARGE_SUBUNIT_BIOGENESIS 491324_GO_0006402_MRNA_CATABOLIC_PROCESS
			492410_GO_0008033_TRNA_PROCESSING 486884_GO_0000387_SPLICEOSOMAL_SNRNP_ASSEMBLY
			491321_GO_0006399_TRNA_METABOLIC_PROCESS 516748_GO_0071826_RIBONUCLEOPROTEIN_COMPLEX_SUBUNIT_ORGANIZATION 491323_GO_0006401_RNA_CATABOLIC_PROCESS
			486955_GO_0000462_MATURATION_OF_SSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA_491322_GO_0006400_TRNA_MODIFICATION
			499353_GO_0022618_RIBONUCLEOPROTEIN_COMPLEX_ASSEMBLY 505414_GO_0042255_RIBOSOME_ASSEMBLY 486579_GO_0000028_RIBOSOMAL_SMALL_SUBUNIT_ASSEMBLY
			517497_GO_0072594_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_ORGANELLE 497937_GO_0019058_VIRAL_LIFE_CYCLE
			492649_GO_0008334_HISTONE_MRNA_METABOLIC_PROCESS 486953_GO_0000460_MATURATION_OF_5_8S_RRNA 506658_GO_0043628_NCRNA_3_END_PROCESSING
			518290_GO_0090150_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MEMBRANE 492576_GO_0008380_RNA_SPLICING
			486877_GO_0000375_RNA_SPLICING_VIA_TRANSESTERIFICATION_REACTIONS 493622_GO_0009451_RNA_MODIFICATION 512270_GO_0051386_REGULATION_OF_NEUROTROPHIN_TRK_RECEPTOR_SIGNALING_PATHWAY
			511847_GO_0050911_DETECTION_OF_CHEMICAL_STIMULUS_INVOLVED_IN_SENSORY_PERCEPTION_OF_SMELL 492350_GO_0007608_SENSORY_PERCEPTION_OF_SMELL
			511844_GO_0050907_DETECTION_OF_CHEMICAL_STIMULUS_INVOLVED_IN_SENSORY_PERCEPTION 493721_GO_0009593_DETECTION_OF_CHEMICAL_STIMULUS 492347_GO_0007606_SENSORY_PERCEPTION_OF_CHEMICAL_STIMULUS
			511843_GO_0050906_DETECTION_OF_STIMULUS_INVOLVED_IN_SENSORY_PERCEPTION 751912_GO_0052696_FLAVONOID_GLUCURONIDATION 493917_GO_0009813_FLAVONOID_BIOSYNTHETIC_PROCESS
			495806_GO_0015669_GAS_TRANSPORT
			491962_GO_0007156_HOMOPHILIC_CELL_ADHESION_VIA_PLASMA_MEMBRANE_ADHESION_MOLECULES 1053349_GO_0098742_CELL_CELL_ADHESION_VIA_PLASMA_MEMBRANE_ADHESION_MOLECULES 982314_GO_0098609_CELL_CELL_ADHESION
			489316_GO_0050832_DEFENSE_RESPONSE_TO_FUNGUS 513612_GO_0060054_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_PROLIFERATION_INVOLVED_IN_WOUND_HEALING
			504902_GO_0035725_SODIUM_ION_TRANSMEMBRANE_TRANSPORT 499181_GO_0021902_COMMITMENT_OF_NEURONAL_CELL_TO_SPECIFIC_NEURON_TYPE_IN_FOREBRAIN 1047426 GO 1903426 REGULATION OF REACTIVE OXYGEN SPECIES BIOSYNTHETIC PROCESS
			503923_GO_0034723_DNA_REPLICATION_DEPENDENT_NUCLEOSOME_ORGANIZATION 513567_GO_0060009_SERTOLI_CELL_DEVELOPMENT
			487918_GO_0002028_REGULATION_OF_SODIUM_ION_TRANSPORT 502002_GO_0032776_DNA_METHYLATION_ON_CYTOSINE 513511_GO_0055078_SODIUM_ION_HOMEOSTASIS
			513566_GO_0060008_SERTOLI_CELL_DIFFERENTIATION 812220_GO_0036376_SODIUM_ION_EXPORT_FROM_CELL
			754110_GO_1901142_INSULIN_METABOLIC_PROCESS 511686_GO_0050715_POSITIVE_REGULATION_OF_CYTOKINE_SECRETION 488581_GO_0002708_POSITIVE_REGULATION_OF_LYMPHOCYTE_MEDIATED_IMMUNITY
			507785_GO_0045428_REGULATION_OF_NITRIC_OXIDE_BIOSYNTHETIC_PROCESS 501961_GO_0032735_POSITIVE_REGULATION_OF_INTERLEUKIN_12_PRODUCTION
			488579_GO_0002706_REGULATION_OF_LYMPHOCYTE_MEDIATED_IMMUNITY 491279_GO_0006335_DNA_REPLICATION_DEPENDENT_NUCLEOSOME_ASSEMBLY 753031_GO_1900034_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT
			501959_GO_0032733_POSITIVE_REGULATION_OF_INTERLEUKIN_10_PRODUCTION 501951_GO_0032725_POSITIVE_REGULATION_OF_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION
			491744_GO_0006883_CELLULAR_SODIUM_ION_HOMEOSTASIS 501871_GO_0032645_REGULATION_OF_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION 502362_GO_0033141_POSITIVE_REGULATION_OF_PEPTIDYL_SERINE_PHOSPHORYLATION_OF_STAT_PROTEIN
			502360_GO_0033139_REGULATION_OF_PEPTIDYL_SERINE_PHOSPHORYLATION_OF_STAT_PROTEIN 516360_GO_0071436_SODIUM_ION_EXPORT
			491109_GO_0006749_GLUTATHIONE_METABOLIC_PROCESS 512175_GO_0051290_PROTEIN_HETEROTETRAMERIZATION 518223_GO_0090083_REGULATION_OF_INCLUSION_BODY_ASSEMBLY
			505339_GO_0042178_XENOBIOTIC_CATABOLIC_PROCESS 754636_GO_1901685_GLUTATHIONE_DERIVATIVE_METABOLIC_PROCESS 754638_GO_1901687_GLUTATHIONE_DERIVATIVE_BIOSYNTHETIC_PROCESS
			754638_GO_1901687_GLUTALHIONE_DERIVATIVE_BIOSYNTHETIC_PROCESS 518224_GO_0090084_NEGATIVE_REGULATION_OF_INCLUSION_BODY_ASSEMBLY 489292_GO_0042026_PROTEIN_REFOLDING
			403232_GO_0042020_FRO1EIN_REFOLDING