		Ф	
Cell	Cell	Monocyte	
B_(lonc	-3 Francials as a set 3
		2	Enrichment score
			489276_GO_0003735_STRUCTURAL_CONSTITUENT_OF_RIBOSOME 498634_GO_0019843_RRNA_BINDING
			492521_GO_0008173_RNA_METHYLTRANSFERASE_ACTIVITY 503685_GO_0034483_HEPARAN_SULFATE_SULFOTRANSFERASE_ACTIVITY
			486597_GO_0000049_TRNA_BINDING 489886_GO_0004549_TRNA_SPECIFIC_RIBONUCLEASE_ACTIVITY
			492523_GO_0008175_TRNA_METHYLTRANSFERASE_ACTIVITY 496778_GO_0016891_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS
			489277_GO_0003743_TRANSLATION_INITIATION_FACTOR_ACTIVITY 490271_GO_0004984_OLFACTORY_RECEPTOR_ACTIVITY
			492632_GO_0008312_7S_RNA_BINDING 489504_GO_0004035_ALKALINE_PHOSPHATASE_ACTIVITY
			489320_GO_0003810_PROTEIN_GLUTAMINE_GAMMA_GLUTAMYLTRANSFERASE_ACTIVITY 490644_GO_0005549_ODORANT_BINDING
			498722_GO_0019956_CHEMOKINE_BINDING 492807_GO_0008528_G_PROTEIN_COUPLED_PEPTIDE_RECEPTOR_ACTIVITY
			487463_GO_0005344_OXYGEN_TRANSPORTER_ACTIVITY 487566_GO_0001653_PEPTIDE_RECEPTOR_ACTIVITY
			492439_GO_0008066_GLUTAMATE_RECEPTOR_ACTIVITY 496691_GO_0016782_TRANSFERASE_ACTIVITY_TRANSFERRING_SULFUR_CONTAINING_GROUPS
			492498_GO_0008146_SULFOTRANSFERASE_ACTIVITY 490203_GO_0004908_INTERLEUKIN_1_RECEPTOR_ACTIVITY
			487554_GO_0001637_G_PROTEIN_COUPLED_CHEMOATTRACTANT_RECEPTOR_ACTIVITY 511517_GO_0050544_ARACHIDONIC_ACID_BINDING
			511516_GO_0050543_ICOSATETRAENOIC_ACID_BINDING 490241_GO_0004950_CHEMOKINE_RECEPTOR_ACTIVITY
			490613_GO_0005504_FATTY_ACID_BINDING 498721_GO_0019955_CYTOKINE_BINDING
			754527_GO_1901567_FATTY_ACID_DERIVATIVE_BINDING 511515_GO_0050542_ICOSANOID_BINDING
			490246_GO_0004955_PROSTAGLANDIN_RECEPTOR_ACTIVITY 492384_GO_0008009_CHEMOKINE_ACTIVITY
			490470_GO_0005229_INTRACELLULAR_CALCIUM_ACTIVATED_CHLORIDE_CHANNEL_ACTIVITY 490192_GO_0004896_CYTOKINE_RECEPTOR_ACTIVITY
			490187_GO_0004890_GABA_A_RECEPTOR_ACTIVITY 497601_GO_0018685_ALKANE_1_MONOOXYGENASE_ACTIVITY
			489513_GO_0004046_AMINOACYLASE_ACTIVITY 501782_GO_0032552_DEOXYRIBONUCLEOTIDE_BINDING
			510363_GO_0048248_CXCR3_CHEMOKINE_RECEPTOR_BINDING 499382_GO_0022836_GATED_CHANNEL_ACTIVITY
			499388_GO_0022843_VOLTAGE_GATED_CATION_CHANNEL_ACTIVITY 751289_GO_0038024_CARGO_RECEPTOR_ACTIVITY
			501784_GO_0032554_PURINE_DEOXYRIBONUCLEOTIDE_BINDING 499363_GO_0022803_PASSIVE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
			490168_GO_0004867_SERINE_TYPE_ENDOPEPTIDASE_INHIBITOR_ACTIVITY 490167_GO_0004866_ENDOPEPTIDASE_INHIBITOR_ACTIVITY
			490458_GO_0005216_ION_CHANNEL_ACTIVITY 495521_GO_0015267_CHANNEL_ACTIVITY
			499383_GO_0022838_SUBSTRATE_SPECIFIC_CHANNEL_ACTIVITY 496916_GO_0030414_PEPTIDASE_INHIBITOR_ACTIVITY
			492692_GO_0008392_ARACHIDONIC_ACID_EPOXYGENASE_ACTIVITY 492691_GO_0008391_ARACHIDONIC_ACID_MONOOXYGENASE_ACTIVITY
			490479_GO_0005245_VOLTAGE_GATED_CALCIUM_CHANNEL_ACTIVITY 496567_GO_0016903_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_ALDEHYDE_OR_OXO_GROUP_OF_DONORS
			489493_GO_0004024_ALCOHOL_DEHYDROGENASE_ACTIVITY_ZINC_DEPENDENT 490475_GO_0005237_INHIBITORY_EXTRACELLULAR_LIGAND_GATED_ION_CHANNEL_ACTIVITY
			496495_GO_0016538_CYCLIN_DEPENDENT_PROTEIN_SERINE_THREONINE_KINASE_REGULATOR_ACTIVITY 496869_GO_0017048_RHO_GTPASE_BINDING
			493634_GO_0015038_GLUTATHIONE_DISULFIDE_OXIDOREDUCTASE_ACTIVITY 500653_GO_0031402_SODIUM_ION_BINDING
			499444_GO_0023023_MHC_PROTEIN_COMPLEX_BINDING 490372_GO_0046875_EPHRIN_RECEPTOR_BINDING
			500248_GO_0030955_POTASSIUM_ION_BINDING 487582_GO_0001671_ATPASE_ACTIVATOR_ACTIVITY
			504171_GO_0034979_NAD_DEPENDENT_PROTEIN_DEACETYLASE_ACTIVITY 500177_GO_0030882_LIPID_ANTIGEN_BINDING
			490518_GO_0005343_ORGANIC_ACID_SODIUM_SYMPORTER_ACTIVITY 500178_GO_0030883_ENDOGENOUS_LIPID_ANTIGEN_BINDING
			489481_GO_0004012_PHOSPHOLIPID_TRANSLOCATING_ATPASE_ACTIVITY 486557_GO_0051082_UNFOLDED_PROTEIN_BINDING
			489236_GO_0005521_LAMIN_BINDING 489324_GO_0003823_ANTIGEN_BINDING
			495537_GO_0015293_SYMPORTER_ACTIVITY 495538_GO_0015294_SOLUTE_CATION_SYMPORTER_ACTIVITY
			490163_GO_0004861_CYCLIN_DEPENDENT_PROTEIN_SERINE_THREONINE_KINASE_INHIBITOR_ACTIVITY 490191_GO_0030246_CARBOHYDRATE_BINDING
			500179_GO_0030884_EXOGENOUS_LIPID_ANTIGEN_BINDING 490029_GO_0004713_PROTEIN_TYROSINE_KINASE_ACTIVITY
			499447_GO_0023026_MHC_CLASS_II_PROTEIN_COMPLEX_BINDING 490291_GO_0005005_TRANSMEMBRANE_EPHRIN_RECEPTOR_ACTIVITY
			507653_GO_0045294_ALPHA_CATENIN_BINDING 500176_GO_0030881_BETA_2_MICROGLOBULIN_BINDING
			515609_GO_0070679_INOSITOL_1_4_5_TRISPHOSPHATE_BINDING 495540_GO_0015296_ANION_CATION_SYMPORTER_ACTIVITY
			753721_GO_1900750_OLIGOPEPTIDE_BINDING 506356_GO_0043295_GLUTATHIONE_BINDING
			492421_GO_0008046_AXON_GUIDANCE_RECEPTOR_ACTIVITY 498574_GO_0019763_IMMUNOGLOBULIN_RECEPTOR_ACTIVITY
			499996_GO_0030695_GTPASE_REGULATOR_ACTIVITY 490567_GO_0005391_SODIUM_POTASSIUM_EXCHANGING_ATPASE_ACTIVITY
			490634_GO_0005536_GLUCOSE_BINDING 492830_GO_0008556_POTASSIUM_TRANSPORTING_ATPASE_ACTIVITY
			490289_GO_0005003_EPHRIN_RECEPTOR_ACTIVITY 490364_GO_0005096_GTPASE_ACTIVATOR_ACTIVITY
			496705_GO_0016801_HYDROLASE_ACTIVITY_ACTING_ON_ETHER_BONDS 500671_GO_0031420_ALKALI_METAL_ION_BINDING
			495364_GO_0015081_SODIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY 496892_GO_0031072_HEAT_SHOCK_PROTEIN_BINDING
			489730_GO_0004364_GLUTATHIONE_TRANSFERASE_ACTIVITY 513558_GO_0055131_C3HC4_TYPE_RING_FINGER_DOMAIN_BINDING