

(a)

# 1	978	41189	8.60%
	4.90E-29	GO:BP	biopolymer metabolic process ...
	1.03E-71	GO:CC	intracellular part
	5.08E-21	GO:MF	protein binding
	6.44E-12	KEGG	Proteasome
	3.86E-10	REACTOME	Formation of Exon Junction ...
# 2	291	4512	10.70%
	1.68E-19	GO:BP	cofactor metabolic process ...
	2.72E-147	GO:CC	mitochondrion
	1.60E-22	GO:MF	oxidoreductase activity
	9.57E-30	KEGG	Oxidative phosphorylation
	1.46E-16	REACTOME	Electron Transport Chain
# 3	246	3052	10.10%
	3.30E-19	GO:BP	biological adhesion
	4.94E-30	GO:CC	proteinaceous extracellular ...
	1.11E-18	GO:MF	extracellular matrix structural ...
	1.92E-14	KEGG	Focal adhesion
	4.02E-08	REACTOME	Hemostasis
# 4	145	3008	28.80%
	6.53E-51	GO:BP	cell cycle
	1.38E-42	GO:CC	nucleus
	8.46E-13	GO:MF	DNA binding
	4.69E-25	KEGG	Cell cycle
	7.76E-28	REACTOME	Cell Cycle, Mitotic
# 5	114	1437	22.30%
	1.29E-22	GO:BP	immune system process
	1.76E-08	GO:CC	lytic vacuole
	6.78E-07	GO:MF	cytokine binding
	1.07E-05	KEGG	B cell receptor signaling pathway ...
	1.46E-08	REACTOME	Signaling in Immune system
# 6	97	511	11%
	3.07E-06	GO:BP	protein transport
	3.32E-08	GO:CC	cytoplasm
	3.71E-09	GO:MF	binding

