

Analysis3_Maintenance_GO_Top10 (with values)

small GTPase-mediated signal transduction	7.18		
regulation of small GTPase mediated signal transduction	6.62		
clathrin-dependent endocytosis	6.6		
Golgi vesicle transport	6.33		
endosomal transport	6.04		
cellular component disassembly	5.86		
cytosolic transport	5.74		
actomyosin structure organization	5.64		
cell-substrate junction organization	5.41		
actin filament organization	4.69		
cellular response to interleukin-1		0.64	
response to interleukin-1		0.64	
DNA-templated DNA replication	0.93	0.64	
fatty acid derivative metabolic process		0.64	
DNA replication		0.64	
coenzyme A metabolic process		0.64	
pyridoxal phosphate catabolic process		0.64	
T-helper 1 cell extravasation		0.64	
T-helper 1 cell diapedesis		0.64	
chemokine (C-C motif) ligand 5 signaling pathway		0.64	
positive regulation of macrophage fusion			0.71
negative regulation of heterochromatin formation			0.71
regulation of macrophage fusion			0.71
eye pigmentation			0.71
negative regulation of extrinsic apoptotic signaling pathway via death domain receptors			0.68
macrophage fusion			0.65
regulation of heterochromatin formation			0.64
negative regulation of chromatin organization			0.64
positive regulation of myeloid leukocyte differentiation			0.63
toll-like receptor signaling pathway			0.63
	HFDvsSD	KDvsSD	KDvsHFD

−log10(Padj)

