		Significant		
	0	20	40	60
multi-organism reproductive process (GO:0044703) positive regulation of apoptotic process (GO:0043065) Ras protein signal transduction (GO:0007265) dorsal closure (GO:0007265) JNK cascade (GO:0007264) regulation of Notch signaling pathway (GO:0008593) positive regulation of fliopodium assembly (GO:0005491) actin cytoskeleton reorganization (GO:0031532) negative regulation of ubiquitin-dependent protein catabolic process (GO:000069) positive regulation of protein binding (GO:0031532) negative regulation of dendritic spine development (GO:00302092) gonad morphogenesis (GO:0035262) positive regulation of dendritic spine development (GO:0060999) sperim individualization (GO:0007291) T cell receptor signaling pathway (GO:0050852) head involution (GO:0008268) regulation of hemocyte differentiation (GO:0003563) melanotic encapsulation of foreign target (GO:003563) melanotic encapsulation of foreign target (GO:0035631) adherens junction maintenance (GO:0034334) endodermal cell differentiation (GO:0035987) germ-band shortening (GO:0007790) germarium-derived female germ-line cyst formation (GO:003579) hemidesmosome assembly (GO:0031581) hemocyte migration (GO:0035099) ovarian fusome organization (GO:0038714) regulation of protein localization to cell surface (GO:0030723) positive regulation of oligodendrocyte differentiation (GO:0048714) regulation of protein localization to cell surface (GO:200008) sensory organ precursor cell fate determination (GO:0004714) regulation of protein localization to cell surface (GO:20000723) acrosome assembly (GO:0001725) dolichyl diphosphate biosynthetic process (GO:0006489) dorsal closure, spreading of leading edge cells (GO:000735) negative regulation of Toll signaling pathway (GO:0006175) regulation modulating pathway (GO:0006179) intermediate filament organization (GO:004519)		-		60