						protein deubiquitination (GO:0016579)	
						protein modification by small protein removal (GO:0070646)	14
						proteolysis involved in cellular protein catabolic process (GO:0051603)	10
						ubiquitin-dependent protein catabolic process (GO:0006511)	12
						modification-dependent macromolecule catabolic process (GO:0043632)	10
						modification-dependent protein catabolic process (GO:0019941)	10
						forebrain morphogenesis (GO:0048853)	8
						lateral ventricle development (GO:0021670)	
						negative regulation of peptidase activity (GO:0010466)	6
						cellular response to transforming growth factor beta stimulus (GO:0071560)	
						response to transforming growth factor beta (GO:0071559)	4
						DNA recombination (GO:0006310)	
						transmembrane receptor protein serine/threonine kinase signaling pathway (GO:0007178)	2
						regulation of cellular response to growth factor stimulus (GO:0090287)	_
						regulation of transmembrane receptor protein serine/threonine kinase signaling pathway (GO:0090092)	0
						regulation of proteasomal protein catabolic process (GO:0061136)	
						transforming growth factor beta receptor signaling pathway (GO:0007179)	
						regulation of cellular protein catabolic process (GO:1903362)	
						regulation of proteolysis involved in cellular protein catabolic process (GO:1903050)	
						midbrain development (GO:0030901)	
						negative regulation of transforming growth factor beta receptor signaling pathway (GO:0030512)	
						negative regulation of transmembrane receptor protein serine/threonine kinase signaling pathway (GO:0090101)	
						negative regulation of cellular response to growth factor stimulus (GO:0090288)	
						regulation of transforming growth factor beta receptor signaling pathway (GO:0017015)	
						regulation of mitochondrial membrane permeability involved in apoptotic process (GO:1902108)	
						regulation of membrane permeability (GO:0090559)	
						regulation of mitochondrial membrane permeability (GO:0046902)	
						mitochondrion organization (GO:0007005)	
						protein localization to mitochondrion (GO:0070585)	
						apoptotic mitochondrial changes (GO:0008637)	
						mitochondrial membrane organization (GO:0007006)	
						aminoglycan biosynthetic process (GO:0006023)	
						glycosaminoglycan biosynthetic process (GO:0006024)	
						regulation of myoblast differentiation (GO:0045661)	
						protein K11-linked ubiquitination (GO:0070979)	
						cytoplasmic pattern recognition receptor signaling pathway (GO:0002753)	
						nucleotide-binding oligomerization domain containing signaling pathway (GO:0070423)	
						aminoglycan metabolic process (GO:0006022)	
						glycosaminoglycan metabolic process (GO:0030203)	
						negative regulation of myoblast differentiation (GO:0045662)	
						antigen processing and presentation (GO:0019882)	
						histone H2A ubiquitination (GO:0033522)	
IsM	ls10ppm	IsLHet	S	Nsl	Nsl		
IsMale	Орр	Het	Het	lale	lale		
	3		lsLHet10ppm	lsMale10ppm	lsMaleLHet		
			рm	ρm	¥		