GO Biological Process

-log10(Binomial p value)

GO Term	Value
cellular response to topologically incorrect protein	66.59
nuclear-transcribed mRNA catabolic process	58.31
mRNA catabolic process	56.89
cellular response to unfolded protein	55.97
endoplasmic reticulum unfolded protein response	54.12
negative regulation of intrinsic apoptotic signaling pathway	52.75
positive regulation of release of cytochrome c from mitochondria	49.47
regulation of mitochondrial membrane permeability	47.22
intrinsic apoptotic signaling pathway in response to DNA damage	46.34
regulation of transcription from RNA polymerase II promoter in response to stress	44.78
regulation of DNA-templated transcription in response to stress	43.45
positive regulation of nuclear-transcribed mRNA catabolic process, deadenylation-dependent decay	43.10
positive regulation of mRNA catabolic process	42.36
regulation of membrane permeability	41.34
response to cadmium ion	41.01
positive regulation of transcription from RNA polymerase II promoter in response to stress	39.90
nuclear-transcribed mRNA catabolic process, deadenylation-dependent decay	38.71
retrograde transport, endosome to Golgi	38.70
regulation of toll-like receptor signaling pathway	37.96
regulation of hematopoietic progenitor cell differentiation	36.92