GO enrichment analysis dN/dS U12-type spliceosomal complex transcription factor TFIID complex synaptic vesicle Count stem cell population maintenance proteasome-mediated ubiquitin-dependent protein catabolic process proteasome complex P-body pvalue nucleotide binding 0.003 nuclear speck 0.002 0.001 mRNA processing intracellular anatomical structure helicase activity Cul3-RING ubiquitin ligase complex 14/22/55/22/58/14/6/22/55/22/55/14/66/17/46/22/57/22/58/17/4