

GO-Term	p-value
biopolymer metabolic process	1,14E-06
regulation of gene expression	1,33E-06
macromolecule metabolic process	2,59E-06
binding	5,57E-06
regulation of macromolecule biosynthetic process	5,57E-06
regulation of cellular metabolic process	5,57E-06
cellular metabolic process	5,57E-06
primary metabolic process	5,57E-06
regulation of biosynthetic process	6,96E-06
RNA metabolic process	7,35E-06
regulation of metabolic process	8,38E-06
nucleic acid binding	8,78E-06
regulation of nucleobase, nucleoside, nucleotide and nucleic acid metabolic process	1,12E-05
regulation of macromolecule metabolic process	1,20E-05
gene expression	1,26E-05
cellular macromolecule biosynthetic process	1,51E-05
macromolecule biosynthetic process	1,51E-05
regulation of transcription	2,82E-05
transcription	2,82E-05
transcription factor activity	3,58E-05
metabolic process	3,58E-05
regulation of RNA metabolic process	3,58E-05
transcription, DNA-dependent	4,58E-05
transcription regulator activity	4,58E-05
cellular biopolymer biosynthetic process	4,58E-05
biopolymer biosynthetic process	4,58E-05
DNA binding	4,73E-05
nucleobase, nucleoside, nucleotide and nucleic acid metabolic process	5,16E-05
regulation of transcription, DNA-dependent	5,51E-05
regulation of cellular process	6,34E-05
nucleus	7,86E-05
transmembrane receptor protein serinethreonine kinase activity 54	8,05E-05
transforming growth factor beta receptor activity	8,05E-05
cellular biopolymer metabolic process	8,05E-05
regulation of biological process	8,48E-05
transmembrane receptor protein kinase activity	0,00011
cellular biosynthetic process	0,00014