

Typical GO terms	p-value
<a href="#">xenobiotic transport</a>	2.50E-05
<a href="#">glycine biosynthetic process from serine</a>	2.34E-02
<a href="#">mannosyl-inositol phosphorylceramide bio:</a>	2.34E-02
<a href="#">pre-miRNA processing</a>	2.34E-02
<a href="#">box H/ACA snoRNA processing</a>	2.34E-02
<a href="#">production of siRNA involved in RNA interf</a>	2.34E-02
<a href="#">cellular response to tetrahydrofolate</a>	2.34E-02
<a href="#">N-acetylglucosamine metabolic process</a>	2.34E-02
<a href="#">microtubule anchoring</a>	2.34E-02
<a href="#">arsenate ion transmembrane transport</a>	2.34E-02
<a href="#">plasma membrane acetate transport</a>	3.11E-02
<a href="#">positive regulation of G1 to G0 transition</a>	3.11E-02
<a href="#">L-serine catabolic process</a>	3.11E-02
<a href="#">primary miRNA processing</a>	3.11E-02
<a href="#">response to cycloheximide</a>	3.11E-02
<a href="#">water transport</a>	3.88E-02
<a href="#">morphogenesis checkpoint</a>	3.88E-02
<a href="#">post-chaperonin tubulin folding pathway</a>	3.88E-02
<a href="#">putrescine transport</a>	3.88E-02
<a href="#">pentose transmembrane transport</a>	3.88E-02
<a href="#">positive regulation of cellular response to d</a>	3.88E-02
<a href="#">cellular response to phosphate starvation</a>	3.88E-02
<a href="#">spermidine transmembrane transport</a>	3.88E-02
<a href="#">spermine transmembrane transport</a>	3.88E-02
<a href="#">positive regulation of transcription involved</a>	3.88E-02
<a href="#">sterol import</a>	3.19E-03
<a href="#">tubulin complex assembly</a>	3.81E-03
<a href="#">positive regulation of transcription from RN</a>	4.63E-02
<a href="#">drug export</a>	8.25E-04
<a href="#">regulation of cell size</a>	2.59E-02
<a href="#">positive regulation of autophagy</a>	2.59E-02
<a href="#">septin ring organization</a>	3.06E-02
<a href="#">regulation of cyclin-dependent protein seri</a>	3.56E-02
<a href="#">divalent inorganic cation transport</a>	3.92E-02
<a href="#">organelle assembly</a>	1.11E-02
<a href="#">cation homeostasis</a>	3.71E-02
<a href="#">cellular macromolecule metabolic process</a>	1.35E-02