size		typical GO term N-acetylglucosamine metabolic process ribosomal small subunit assembly glycine biosynthetic process from serine cellular response to tetrahydrofolate L-serine catabolic process 'de novo' UMP biosynthetic process folic acid metabolic process	p-value 1.98E-03 1.97E-02 1.98E-03 1.98E-03 2.64E-03 4.62E-03 5.28E-03
	5	tetrahydrofolate interconversion 'de novo' pyrimidine nucleobase biosynthetic Rab protein signal transduction septin ring assembly protein localization to phagophore assembly cellular response to salt stress	1.15E-02 1.40E-02
	10	regulation of cell shape budding cell bud growth xenobiotic transport plasma membrane acetate transport biotin biosynthetic process ammonium transmembrane transport	1.97E-02 2.62E-02 5.13E-05 6.59E-03 1.15E-02 2.77E-02
	8	drug export arsenate ion transmembrane transport positive regulation of cellular response to dru mitotic spindle assembly zinc ion transmembrane transport cellular zinc ion homeostasis establishment of mitotic spindle orientation regulation of cyclin-dependent protein serine drug export positive regulation of macroautophagy	1.04E-02 1.15E-02 1.38E-02 1.84E-02
		positive regulation of transcription from RNA age-dependent response to oxidative stress amino acid catabolic process to alcohol via E positive regulation of autophagy positive regulation of mitotic cell cycle phase mannosyl-inositol phosphorylceramide biosy putrescine transport pentose transmembrane transport spermidine transmembrane transport	3.96E-03 4.62E-03 7.91E-03 2.03E-02 2.16E-02