fraction	size	typical GO term	p-value
1	5	N-acetylglucosamine metabolic process	1.98E-03
		ribosomal small subunit assembly	1.97E-02
2	4	glycine biosynthetic process from serine	1.98E-03
		cellular response to tetrahydrofolate	1.98E-03
		L-serine catabolic process	2.64E-03
		'de novo' UMP biosynthetic process	4.62E-03
		folic acid metabolic process	5.28E-03
		tetrahydrofolate interconversion	5.28E-03
		'de novo' pyrimidine nucleobase biosynthetic process	5.94E-03
3	5	Rab protein signal transduction	1.15E-02
		septin ring assembly	1.40E-02
		protein localization to phagophore assembly site	1.48E-02
		cellular response to salt stress	1.56E-02
		regulation of cell shape	1.97E-02
		budding cell bud growth	2.62E-02
4	10	xenobiotic transport	5.13E-05
		plasma membrane acetate transport	6.59E-03
		biotin biosynthetic process	1.15E-02
		ammonium transmembrane transport	2.77E-02
		drug export	3.57E-02
5	8	arsenate ion transmembrane transport	3.46E-03
		positive regulation of cellular response to drug	5.77E-03
		mitotic spindle assembly	1.04E-02
		zinc ion transmembrane transport	1.15E-02
		cellular zinc ion homeostasis	1.38E-02
		establishment of mitotic spindle orientation	1.84E-02
		regulation of cyclin-dependent protein serine/threonine kinase activity	7.89E-04
		drug export	2.52E-02
		positive regulation of macroautophagy	2.74E-02
6	4	age-dependent response to oxidative stress involved in chronological cell aging	4.62E-03
		amino acid catabolic process to alcohol via Ehrlich pathway	7.91E-03
		positive regulation of autophagy	2.03E-02
		positive regulation of mitotic cell cycle phase transition	2.16E-02
7	7	mannosyl-inositol phosphorylceramide biosynthetic process	2.97E-03
		putrescine transport	4.95E-03
		pentose transmembrane transport	4.95E-03
		spermidine transmembrane transport	4.95E-03