Pathway	Gene ranks	NES	pval	padj
GOBP_CILIUM_ORGANIZATION		1.76	1.2·10 ⁻⁸	1.8·10 ⁻⁶
GOBP_CELL_PROJECTION_ASSEMBLY		1.50	7.2·10 ⁻⁷	6.9·10 ⁻⁵
GOBP_INTRACILIARY_TRANSPORT		2.21	6.7·10 ⁻⁷	6.5·10 ⁻⁵
GOBP_MICROTUBULE_BASED_TRANSPORT		1.76	1.4·10 ⁻⁶	1.2-10 ⁻⁴
GOBP_AXON_DEVELOPMENT		1.50	8.5·10 ⁻⁶	6.0-10 ⁻⁴
GOBP_PROCESS_UTILIZING_AUTOPHAGIC_MECHANISM		1.46	1.6·10 ⁻⁵	1.0·10 ⁻³
GOBP_VESICLE_ORGANIZATION		1.55	1.7·10 ⁻⁵	1.0·10 ⁻³
GOBP_STEROL_BIOSYNTHETIC_PROCESS		2.13	5.3·10 ⁻⁶	3.9·10 ⁻⁴
GOBP_CILIUM_MOVEMENT		1.77	2.9·10 ⁻⁵	1.6·10 ⁻³
GOBP_MEMBRANE_FUSION		1.77	2.7·10 ⁻⁵	1.6·10 ⁻³
GOBP_RIBOSOMAL_LARGE_SUBUNIT_BIOGENESIS		-2.53	6.0·10 ⁻¹³	2.6·10 ⁻¹⁰
GOBP_MITOTIC_NUCLEAR_DIVISION		-2.17	3.0·10 ⁻¹³	1.7·10 ⁻¹⁰
GOBP_MITOTIC_SISTER_CHROMATID_SEGREGATION		-2.28	3.2·10 ⁻¹³	1.7·10 ⁻¹⁰
GOBP_TRANSLATION_AT_SYNAPSE	' ' '	-2.75	3.4·10 ⁻¹⁶	2.3·10 ⁻¹³
GOBP_RIBOSOMAL_SMALL_SUBUNIT_BIOGENESIS		-2.76	1.3·10 ⁻²¹	1.0·10 ⁻¹⁸
GOBP_CYTOPLASMIC_TRANSLATION		-2.86	1.2-10 ⁻²⁸	1.2·10 ⁻²⁵
GOBP_RRNA_METABOLIC_PROCESS		-2.69	7.8·10 ⁻³¹	9.2·10 ⁻²⁸
GOBP_RRNA_PROCESSING		-2.77	2.3·10 ⁻³²	3.7·10 ⁻²⁹
GOBP_RIBOSOME_BIOGENESIS		-2.82	1.3.10 ⁻⁴³	3.1·10 ⁻⁴⁰
GOBP_RIBONUCLEOPROTEIN_COMPLEX_BIOGENESIS		-2.70	2.2-10 ⁻⁴⁴	1.0·10 ⁻⁴⁰
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