Pathway	Gene ranks	NES	pval	padj
PROTEIN-DNA COMPLEX DISASSEMBLY%GOBP%GO:0032986		2.14	2.2e-04	2.7e-03
PRC2 METHYLATES HISTONES AND DNA%REACTOME DATABASE ID RELEASE 74%212300	MITTO 1 11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	2.26	2.2e-04	2.7e-03
G ALPHA (Z) SIGNALLING EVENTS%REACTOME%R-HSA-418597.4		2.10	2.2e-04	2.7e-03
CONDENSATION OF PROPHASE CHROMOSOMES%REACTOME%R-HSA-2299718.1		2.04	2.2e-04	2.7e-03
REGULATION OF MRNA POLYADENYLATION%GOBP%GO:1900363		2.13	2.2e-04	2.7e-03
PROCESSING OF INTRONLESS PRE-MRNAS%REACTOME%R-HSA-77595.2	The Hermitian control of the management of the second of t	2.38	2.2e-04	2.7e-03
TUMOR SUPPRESSOR ACTIVITY OF SMARCB1%WIKIPATHWAYS_20201110%WP4204%HOMO SAPIENS	THILLIE TO THE TOTAL TO THE TOTAL TO	2.17	2.2e-04	2.7e-03
MRNA CLEAVAGE%GOBP%GO:0006379	THE COURT OF THE C	2.31	2.2e-04	2.7e-03
MRNA SPLICE SITE SELECTION%GOBP%GO:0006376	HET HITCHIEF CONTRACTOR CONTRACTO	2.13	2.2e-04	2.7e-03
REGULATION OF CHROMATIN SILENCING%GOBP%GO:0031935		2.21	2.2e-04	2.7e-03
CELL ACTIVATION%GOBP%GO:0001775		-1.64	1.4e-04	2.6e-03
OXOACID METABOLIC PROCESS%GOBP%GO:0043436		-1.59	1.4e-04	2.6e-03
SECRETION BY CELL%GOBP%GO:0032940		-1.64	1.4e-04	2.6e-03
ORGANIC ACID METABOLIC PROCESS%GOBP%GO:0006082		-1.62	1.4e-04	2.6e-03
SECRETION%GOBP%GO:0046903		-1.64	1.4e-04	2.6e-03
MOVEMENT OF CELL OR SUBCELLULAR COMPONENT%GOBP%GO:0006928		-1.50	1.4e-04	2.6e-03
REGULATION OF CELLULAR COMPONENT BIOGENESIS%GOBP%GO:0044087		-1.62	1.4e-04	2.6e-03
ORGANOPHOSPHATE METABOLIC PROCESS%GOBP%GO:0019637		-1.78	1.4e-04	2.6e-03
IMMUNE RESPONSE%GOBP%GO:0006955		-1.56	1.4e-04	2.6e-03
PHOSPHORYLATION%GOBP%GO:0016310		-1.43	1.3e-04	2.6e-03
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