

NES

SET

6.824		GOBP_MITOCHONDRIAL_GENE_EXPRESSION
6.498		GOBP_TRANSLATIONAL_TERMINATION
6.491		GOBP_MITOCHONDRIAL_TRANSLATION
6.484		GOBP_TRANSLATIONAL_ELONGATION
6.180		GOBP_MITOCHONDRIAL_TRANSLATIONAL_TERMINATION
5.933		GOCC_MITOCHONDRIAL_PROTEIN_CONTAINING_COMPLEX
5.765		GOCC_MITOCHONDRIAL_LARGE_RIBOSOMAL_SUBUNIT
5.716		GOCC_MITOCHONDRIAL_MATRIX
5.686		GOCC_ORGANELLAR_RIBOSOME
4.773		GOBP_CELLULAR_PROTEIN_COMPLEX_DISASSEMBLY
4.468		GOBP_TRNA_PROCESSING
4.451		GOCC_ORGANELLE_INNER_MEMBRANE
4.447		GOBP_PROTEIN_CONTAINING_COMPLEX_DISASSEMBLY
-4.420		GOBP_REGULATION_OF_RAS_PROTEIN_SIGNAL_TRANSDUCTION
4.389		GOBP_TRNA_METABOLIC_PROCESS
4.387		GOBP_TRNA_WOBBLE_BASE_MODIFICATION
4.343		GOCC_ELONGATOR_HOLOENZYME_COMPLEX
-4.264		GOBP_PROTEIN_K11_LINKED_UBIQUITINATION
4.262		GOBP_HISTONE_H2A_ACETYLATION
4.201		GOCC_H4_H2A_HISTONE_ACETYLTRANSFERASE_COMPLEX
4.185		GOBP_RNA_SPLICING_VIA_TRANSESTERIFICATION_REACTIONS
4.166		GOBP_NUCLEOSIDE_BISPHOSPHATE_BIOSYNTHETIC_PROCESS
-4.149		GOBP_RNA_CATABOLIC_PROCESS
-3.976		GOBP_P_BODY_ASSEMBLY
3.973		GOMF_TRANSCRIPTION_REGULATOR_ACTIVITY
3.947		GOBP_MRNA_PROCESSING
3.918		GOBP_THIOESTER_BIOSYNTHETIC_PROCESS
3.848		GOBP_THIOESTER_METABOLIC_PROCESS
3.846		GOMF_DNA_BINDING_TRANSCRIPTION_FACTOR_ACTIVITY
-3.843		GOBP_CELL_PROLIFERATION_INVOLVED_IN_HEART_MORPHOGENESIS
3.831		GOBP_RNA_PROCESSING
-3.825		GOBP_RESPONSE_TO_NERVE_GROWTH_FACTOR
3.820		GOBP_NUCLEOSIDE_BISPHOSPHATE_METABOLIC_PROCESS
-3.803		GOBP_DOUBLE_STRAND_BREAK_REPAIR
3.799		GOBP_ACETYL_COA_METABOLIC_PROCESS
3.781		GOMF_SEQUENCE_SPECIFIC_DNA_BINDING
3.771		GOMF_RNA_BINDING
3.740		GOBP_RNA_SPLICING
3.706		GOBP_ACETYL_COA_BIOSYNTHETIC_PROCESS
3.701		GOBP_TRNA_THREONYLCARBAMOYLADENOSINE_METABOLIC_PROCESS
3.658		GOBP_DOSAGE_COMPENSATION
-3.647		GOBP_POSITIVE_REGULATION_OF_NUCLEAR_CELL_CYCLE_DNA_REPLICATION
3.637		GOCC_MITOCHONDRIAL_ENVELOPE
3.614		GOBP_REGULATION_OF_PROTEIN_FOLDING
3.613		GOBP_ACETYL_COA_BIOSYNTHETIC_PROCESS_FROM_PYRUVATE