



- GOBP_TRANSLATIONAL_ELONGATION,
- GOBP_MITOCHONDRIAL_TRANSLATIONAL_TERMINATION,
- GOCC_ORGANELLAR_RIBOSOME,
- GOBP_TRANSLATIONAL_TERMINATION,
- GOBP_MITOCHONDRIAL_TRANSLATION,
- GOBP_MITOCHONDRIAL_GENE_EXPRESSION,
- GOCC_MITOCHONDRIAL_LARGE_RIBOSOMAL_SUBUNIT,
- GOBP_REGULATION_OF_PROTEIN_FOLDING,
- GOBP_TRNA_METABOLIC_PROCESS,
- GOBP_TRNA_PROCESSING,
- GOBP_RNA_MODIFICATION,
- GOBP_TRNA_WOBBLE_BASE_MODIFICATION,
- GOCC_H4_H2A_HISTONE_ACETYLTRANSFERASE_COMPLEX,
- GOBP_HISTONE_H2A_ACETYLATION,
- GOCC_ELONGATOR_HOLOENZYME_COMPLEX,
- GOCC_MITOCHONDRIAL_SMALL_RIBOSOMAL_SUBUNIT,
- GOBP_THIOESTER_BIOSYNTHETIC_PROCESS,
- GOBP_NUCLEOSIDE_BISPHOSPHATE_BIOSYNTHETIC_PROCESS,
- GOCC_RIBOSOMAL_SUBUNIT,
- GOMF_STRUCTURAL_CONSTITUENT_OF_RIBOSOME,
- GOCC_LARGE_RIBOSOMAL_SUBUNIT,
- GOCC_SPLICEOSOMAL_COMPLEX,
- GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSLATION,
- GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATION,
- GOMF_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS,
- GOBP_POSITIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS,
- GOBP_POSITIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_COMPLEXES,
- GOBP_CYTOPLASMIC_TRANSLATION,
- GOBP_TELOMERE_ORGANIZATION,
- GOCC_PROTEIN_ACETYLTRANSFERASE_COMPLEX,
- GOCC_INO80_TYPE_COMPLEX,
- GOCC_KINETOCHORE,
- GOCC_CHROMOSOME_CENTROMERIC_REGION,
- GOBP_REGULATION_OF_PEROXISOME_PROLIFERATOR_ACTIVATION,
- GOBP_THIOESTER_METABOLIC_PROCESS,
- GOBP_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE,