

PROTEIN_CONTAINING_COMPLEX, GOCC_ENDOPLASMIC_RETICULUM_PROTEIN_CONTAINING_COMPLEX

- GOCC_PHAGOCYTIC_VESICLE, GOCC_PHAGOCYTIC_VESICLE
- GOCC_ENDOPEPTIDASE_COMPLEX, GOCC_ENDOPEPTIDASE_COMPLEX
- GOCC_PEPTIDASE_COMPLEX, GOCC_PEPTIDASE_COMPLEX
- GOMF_UBIQUITIN_LIKE_PROTEIN_BINDING, GOMF_UBIQUITIN_LIKE_PROTEIN_BINDING
- GOBP_ERAD_PATHWAY, GOBP_ERAD_PATHWAY
- GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I, GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I
- GOCC_PHAGOCYTIC_VESICLE_MEMBRANE, GOCC_PHAGOCYTIC_VESICLE_MEMBRANE
- GOMF_UBIQUITIN_BINDING, GOMF_UBIQUITIN_BINDING
- GOCC_ENDOCYTIC_VESICLE_MEMBRANE, GOCC_ENDOCYTIC_VESICLE_MEMBRANE
- GOCC_INTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE, GOCC_INTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE
- GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY, GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY
- GOBP_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GOBP_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
- GOBP_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT, GOBP_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT
- GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I, GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I
- GOBP_IRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE, GOBP_IRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE
- GOCC_PIGMENT_GRANULE, GOCC_PIGMENT_GRANULE
- GOBP_PHOSPHATIDYLINOSITOL_BIOSYNTHETIC_PROCESS, GOBP_PHOSPHATIDYLINOSITOL_BIOSYNTHETIC_PROCESS
- GOBP_LIPOSACCHARIDE_METABOLIC_PROCESS, GOBP_LIPOSACCHARIDE_METABOLIC_PROCESS
- GOBP_REGULATION_OF_JUN_KINASE_ACTIVITY, GOBP_REGULATION_OF_JUN_KINASE_ACTIVITY
- GOBP_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT, GOBP_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT
- GOMF_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_BINDING, GOMF_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_BINDING
- GOBP_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
- GOCC_MHC_PROTEIN_COMPLEX, GOCC_MHC_PROTEIN_COMPLEX
- GOBP_POSITIVE_REGULATION_OF_JUN_KINASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_JUN_KINASE_ACTIVITY
- GOMF_PALMITOYLTRANSFERASE_ACTIVITY, GOMF_PALMITOYLTRANSFERASE_ACTIVITY
- GOBP_SPHINGOLIPID_METABOLIC_PROCESS, GOBP_SPHINGOLIPID_METABOLIC_PROCESS
- GOMF_UBIQUITIN_CONJUGATING_ENZYME_BINDING, GOMF_UBIQUITIN_CONJUGATING_ENZYME_BINDING
- GOBP_MITOCHONDRIAL_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT, GOBP_MITOCHONDRIAL_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
- GOBP_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS
- GOBP_CRISTAE_FORMATION, GOBP_CRISTAE_FORMATION
- GOMF_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GOMF_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING
- GOBP_REGULATION_OF_PRODUCTION_OF_MOLECULAR_MEDIATOR_OF_IMMUNE_RESPONSE, GOBP_REGULATION_OF_PRODUCTION_OF_MOLECULAR_MEDIATOR_OF_IMMUNE_RESPONSE
- GOBP_POSTTRANSLATIONAL_PROTEIN_TARGETING_TO_ENDOPLASMIC_RETICULUM_MEMBRANE, GOBP_POSTTRANSLATIONAL_PROTEIN_TARGETING_TO_ENDOPLASMIC_RETICULUM_MEMBRANE
- GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I_TAP_INDEPENDENT, GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I_TAP_INDEPENDENT
- GOBP_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS, GOBP_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS
- GOBP_MEMBRANE_LIPID_BIOSYNTHETIC_PROCESS, GOBP_MEMBRANE_LIPID_BIOSYNTHETIC_PROCESS
- GOBP_POSITIVE_REGULATION_OF_CYTOKINE_PRODUCTION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_CYTOKINE_PRODUCTION_INVOLVED_IN_IMMUNE_RESPONSE
- GOMF_ISOMERASE_ACTIVITY, GOMF_ISOMERASE_ACTIVITY
- GOCC_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX, GOCC_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX
- GOBP_MANNOSYLATION, GOBP_MANNOSYLATION
- GOBP_LIPOPROTEIN_BIOSYNTHETIC_PROCESS, GOBP_LIPOPROTEIN_BIOSYNTHETIC_PROCESS
- GOCC_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT, GOCC_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT
- GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_BIOTIC_STIMULUS, GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_BIOTIC_STIMULUS
- GOMF_PROTON_TRANSPORTING_ATP_SYNTHASE_ACTIVITY_ROTATIONAL_MECHANISM, GOMF_PROTON_TRANSPORTING_ATP_SYNTHASE_ACTIVITY_ROTATIONAL_MECHANISM
- GOBP_PROTEIN_MANNOSYLATION, GOBP_PROTEIN_MANNOSYLATION
- GOMF_MOLECULAR_CARRIER_ACTIVITY, GOMF_MOLECULAR_CARRIER_ACTIVITY
- GOBP_PROTEIN_INSERTION_INTO_MEMBRANE, GOBP_PROTEIN_INSERTION_INTO_MEMBRANE
- GOBP_PROTEIN_N_LINKED_GLYCOSYLATION, GOBP_PROTEIN_N_LINKED_GLYCOSYLATION
- GOBP_PHOSPHATIDYLINOSITOL_METABOLIC_PROCESS, GOBP_PHOSPHATIDYLINOSITOL_METABOLIC_PROCESS
- GOBP_NEGATIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS, GOBP_NEGATIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS
- GOBP_PROTEIN_O_LINKED_MANNOSYLATION, GOBP_PROTEIN_O_LINKED_MANNOSYLATION
- GOBP_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT, GOBP_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
- GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_A_SULFUR_GROUP_OF_DONORS, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_A_SULFUR_GROUP_OF_DONORS
- GOCC_MHC_CLASS_I_PROTEIN_COMPLEX, GOCC_MHC_CLASS_I_PROTEIN_COMPLEX
- GOBP_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM, GOBP_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM
- GOBP_SELECTIVE_AUTOPHAGY, GOBP_SELECTIVE_AUTOPHAGY
- GOBP_REGULATION_OF_ER_TO_GOLGI_VESICLE_MEDIATED_TRANSPORT, GOBP_REGULATION_OF_ER_TO_GOLGI_VESICLE_MEDIATED_TRANSPORT
- GOBP_PROTEIN_TRANSMEMBRANE_TRANSPORT, GOBP_PROTEIN_TRANSMEMBRANE_TRANSPORT
- GOBP_PROTEIN_DEGLYCOSYLATION, GOBP_PROTEIN_DEGLYCOSYLATION
- GOCC_OLIGOSACCHARYLTRANSFERASE_COMPLEX, GOCC_OLIGOSACCHARYLTRANSFERASE_COMPLEX
- GOMF_DISULFIDE_OXIDOREDUCTASE_ACTIVITY, GOMF_DISULFIDE_OXIDOREDUCTASE_ACTIVITY
- GOBP_RESPONSE_TO_MISFOLDED_PROTEIN, GOBP_RESPONSE_TO_MISFOLDED_PROTEIN
- GOCC_TRANSLOCON_COMPLEX, GOCC_TRANSLOCON_COMPLEX
- GOMF_VITAMIN_B6_BINDING, GOMF_VITAMIN_B6_BINDING
- GOBP_GPI_ANCHOR_METABOLIC_PROCESS, GOBP_GPI_ANCHOR_METABOLIC_PROCESS
- GOBP_POSITIVE_REGULATION_OF_PRODUCTION_OF_MOLECULAR_MEDIATOR_OF_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_PRODUCTION_OF_MOLECULAR_MEDIATOR_OF_IMMUNE_RESPONSE
- GOBP_PROTON_TRANSMEMBRANE_TRANSPORT, GOBP_PROTON_TRANSMEMBRANE_TRANSPORT
- GOMF_MEMBRANE_INSERTASE_ACTIVITY, GOMF_MEMBRANE_INSERTASE_ACTIVITY
- GOBP_POSITIVE_REGULATION_OF_PROTEIN_BINDING, GOBP_POSITIVE_REGULATION_OF_PROTEIN_BINDING
- GOMF_PROTON_CHANNEL_ACTIVITY, GOMF_PROTON_CHANNEL_ACTIVITY
- GOCC_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_CATALYTIC_DOMAIN, GOCC_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_CATALYTIC_DOMAIN
- GOBP_CHAPERONE_MEDIATED_PROTEIN_FOLDING, GOBP_CHAPERONE_MEDIATED_PROTEIN_FOLDING
- GOCC_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX, GOCC_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX