

_TO_GLUCOSE_STARVATION, GO_CELLULAR_RESPONSE_TO_GLUCOSE_STARVATION

- GO_INTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE, GO_INTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE
- GO_ENDOPLASMIC_RETICULUM_UNFOLDED_PROTEIN_RESPONSE, GO_ENDOPLASMIC_RETICULUM_UNFOLDED_PROTEIN_RESPONSE
- GO_PHOSPHATASE_BINDING, GO_PHOSPHATASE_BINDING
- GO_IRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE, GO_IRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE
- GO_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GO_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
- GO_POSITIVE_REGULATION_OF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_PRODUCTION, GO_POSITIVE_REGULATION_OF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_PRODUCTION
- GO_PHAGOCYTIC_VESICLE_MEMBRANE, GO_PHAGOCYTIC_VESICLE_MEMBRANE
- GO_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN, GO_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN
- GO_AUTOPHAGOSOME, GO_AUTOPHAGOSOME
- GO_CELL_MATURATION, GO_CELL_MATURATION
- GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
- GO_REGULATION_OF_TOR_SIGNALING, GO_REGULATION_OF_TOR_SIGNALING
- GO_REGULATION_OF_PROTEIN_COMPLEX_STABILITY, GO_REGULATION_OF_PROTEIN_COMPLEX_STABILITY
- GO_PROTEIN_TARGETING_TO_VACUOLE, GO_PROTEIN_TARGETING_TO_VACUOLE
- GO_CIRCADIAN_REGULATION_OF_GENE_EXPRESSION, GO_CIRCADIAN_REGULATION_OF_GENE_EXPRESSION
- GO_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ENDOPLASMIC_RETICULUM, GO_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ENDOPLASMIC_RETICULUM
- GO_NEGATIVE_REGULATION_OF_MYOTUBE_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_MYOTUBE_DIFFERENTIATION
- GO_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_STRESS, GO_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_STRESS
- GO_EXTRINSIC_COMPONENT_OF_ORGANELLE_MEMBRANE, GO_EXTRINSIC_COMPONENT_OF_ORGANELLE_MEMBRANE
- GO_ORGANELLE_MEMBRANE_CONTACT_SITE, GO_ORGANELLE_MEMBRANE_CONTACT_SITE
- GO_NEGATIVE_REGULATION_OF_TOR_SIGNALING, GO_NEGATIVE_REGULATION_OF_TOR_SIGNALING
- GO_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX, GO_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX
- GO_PHAGOPHORE_ASSEMBLY_SITE_MEMBRANE, GO_PHAGOPHORE_ASSEMBLY_SITE_MEMBRANE
- GO_LYMPHOCYTE_HOMEOSTASIS, GO_LYMPHOCYTE_HOMEOSTASIS
- GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
- GO_MIRNA_BINDING, GO_MIRNA_BINDING
- GO_PHOSPHATIDYLINOSITOL_METABOLIC_PROCESS, GO_PHOSPHATIDYLINOSITOL_METABOLIC_PROCESS
- GO_PYRIMIDINE_RIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS, GO_PYRIMIDINE_RIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS
- GO_ATF6_MEDIATED_UNFOLDED_PROTEIN_RESPONSE, GO_ATF6_MEDIATED_UNFOLDED_PROTEIN_RESPONSE