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GOBP_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
  GOMF_UBIQUITIN_BINDING, GOMF_UBIQUITIN_BINDING
  GOBP_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS, GOBP_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS
  GOCC_INTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE, GOCC_INTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE
  GOBP_POSITIVE_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_PROCESS, GO
 GOBP_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY, GOBP_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY
 GOMF_ENDOPEPTIDASE_REGULATOR_ACTIVITY, GOMF_ENDOPEPTIDASE_REGULATOR_ACTIVITY
 GOBP_ERAD_PATHWAY, GOBP_ERAD_PATHWAY
  GOMF_CYSTEINE_TYPE_PEPTIDASE_ACTIVITY, GOMF_CYSTEINE_TYPE_PEPTIDASE_ACTIVITY
 GOBP_POSITIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS
 GOBP_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
 GOBP REGULATION OF HEMATOPOIETIC PROGENITOR CELL DIFFERENTIATION, GOBP REGULATION OF HEMATOPOIETIC PROGENITOR CELL DIFFERENTIATION
 GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I, GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I
 GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY, GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY
 GOBP_POSITIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULAR_PROCESS, GOBP_POSITIVE_REGULAR_PROCESS, GOBP_POSITIVE_REGULAR_PROCESS, GOBP_POSITIVE_REGULAR_PROCESS, GOBP_POSITIVE_REGULAR_PROCESS, GOBP_POSITIVE_REGULAR_PROCESS, GOBP_POSITI
 GOBP_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION, GOBP_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION
 GOBP_POSITIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
 GOCC_SAGA_TYPE_COMPLEX, GOCC_SAGA_TYPE_COMPLEX
 GOMF_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY, GOMF_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY
 GOBP_REGULATION_OF_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION
 GOCC_RNA_POLYMERASE_II_TRANSCRIPTION_REGULATOR_COMPLEX, GOCC_RNA_POLYMERASE_II_TRANSCRIPTION_REGULATOR_COMPLEX
- HP_GAIT_IMBALANCE, HP_GAIT_IMBALANCE
 GOMF_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GOMF_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING
 GOBP_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS, GOBP_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS
 GOCC_PROTEASOME_ACCESSORY_COMPLEX, GOCC_PROTEASOME_ACCESSORY_COMPLEX
 GOMF_OMEGA_PEPTIDASE_ACTIVITY, GOMF_OMEGA_PEPTIDASE_ACTIVITY
 GOBP_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION, GOBP_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION
 GOBP_TRANSCRIPTION_PREINITIATION_COMPLEX_ASSEMBLY, GOBP_TRANSCRIPTION_PREINITIATION_COMPLEX_ASSEMBLY
 HP_HYPERTRIGLYCERIDEMIA, HP_HYPERTRIGLYCERIDEMIA
 GOCC_TRANSCRIPTION_FACTOR_TFIID_COMPLEX, GOCC_TRANSCRIPTION_FACTOR_TFIID_COMPLEX
 GOBP_NEGATIVE_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_CATABOLICATABOLICATABOLICATABOLICATABOLICATABOLICATABOLICATABOLICATABOLICATABOLICATABOLICA
 HP_HIGHLY_ARCHED_EYEBROW, HP_HIGHLY_ARCHED_EYEBROW
 GOMF_RNA_POLYMERASE_II_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY, GOMF_RNA_POLYMERASE_II_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY
 GOMF_PROTEASOME_BINDING, GOMF_PROTEASOME_BINDING
 GOBP_PROTEASOMAL_UBIQUITIN_INDEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_PROTEASOMAL_UBIQUITIN_INDEPENDENT_PROTEIN_CATABOLIC_PROCESS
 GOBP_RNA_POLYMERASE_II_PREINITIATION_COMPLEX_ASSEMBLY, GOBP_RNA_POLYMERASE_II_PREINITIATION_COMPLEX_ASSEMBLY
 HP_FRONTAL_CORTICAL_ATROPHY, HP_FRONTAL_CORTICAL_ATROPHY
 HP_HYPERLIPIDEMIA, HP_HYPERLIPIDEMIA
 GOBP_PHOSPHATIDYLINOSITOL_BIOSYNTHETIC_PROCESS, GOBP_PHOSPHATIDYLINOSITOL_BIOSYNTHETIC_PROCESS
 GOBP_MEMBRANE_LIPID_BIOSYNTHETIC_PROCESS, GOBP_MEMBRANE_LIPID_BIOSYNTHETIC_PROCESS
 GOBP_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY, GOBP_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY
 GOBP_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
 GOBP_REGULATION_OF_STEM_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_STEM_CELL_DIFFERENTIATION
 GOMF LIPOPOLYSACCHARIDE BINDING, GOMF LIPOPOLYSACCHARIDE BINDING
 GOBP_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER, GOBP_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER
 GOBP_REGULATION_OF_CELLULAR_AMINE_METABOLIC_PROCESS, GOBP_REGULATION_OF_CELLULAR_AMINE_METABOLIC_PROCESS
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GOCC ENDOPEPTIDASE COMPLEX, GOCC ENDOPEPTIDASE COMPLEX

GOCC_RNA_POLYMERASE_COMPLEX, GOCC_RNA_POLYMERASE_COMPLEX

GOBP_PEPTIDYL_LYSINE_ACETYLATION, GOBP_PEPTIDYL_LYSINE_ACETYLATION

GOMF_UBIQUITIN_LIKE_PROTEIN_BINDING, GOMF_UBIQUITIN_LIKE_PROTEIN_BINDING

GOCC_RNA_POLYMERASE_II_HOLOENZYME, GOCC_RNA_POLYMERASE_II_HOLOENZYME

GOCC_PROTEIN_ACETYLTRANSFERASE_COMPLEX, GOCC_PROTEIN_ACETYLTRANSFERASE_COMPLEX

GOMF_NUCLEAR_RECEPTOR_COACTIVATOR_ACTIVITY, GOMF_NUCLEAR_RECEPTOR_COACTIVATOR_ACTIVITY

GOBP_ACTIVATION_OF_INNATE_IMMUNE_RESPONSE, GOBP_ACTIVATION_OF_INNATE_IMMUNE_RESPONSE

GOBP_MORPHOGENESIS_OF_A_POLARIZED_EPITHELIUM, GOBP_MORPHOGENESIS_OF_A_POLARIZED_EPITHELIUM

GOBP_POSITIVE_REGULATION_OF_WNT_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_WNT_SIGNALING_PATHWAY

GOCC_PROTEASOME_REGULATORY_PARTICLE_BASE_SUBCOMPLEX, GOCC_PROTEASOME_REGULATORY_PARTICLE_BASE_SUBCOMPLEX

GOBP_PROTEIN_K63_LINKED_DEUBIQUITINATION, GOBP_PROTEIN_K63_LINKED_DEUBIQUITINATION

HP_SLENDER_LONG_BONE, HP_SLENDER_LONG_BONE

HP_MUSCLE_FIBRILLATION, HP_MUSCLE_FIBRILLATION

GOBP_HISTONE_UBIQUITINATION, GOBP_HISTONE_UBIQUITINATION

GOCC_ENDOPLASMIC_RETICULUM_PROTEIN_CONTAINING_COMPLEX, GOCC_ENDOPLASMIC_RETICULUM_PROTEIN_CONTAINING_COMPLEX

GOBP_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_IN_RESPONSE_TO_STRESS, GOBP_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_IN_RESPONSE_TO_STRESS

GOBP_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS

GOBP_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA, GOBP_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESI

IDASE_COMPLEX, GOCC_PEPTIDASE_COMPLEX