

D_TRANSCRIPTION_IN_RESPONSE_TO_STRESS, GOBP_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_IN_RESPONSE_TO_STRESS

- GOBP_ENDOPLASMIC_RETICULUM_UNFOLDED_PROTEIN_RESPONSE, GOBP_ENDOPLASMIC_RETICULUM_UNFOLDED_PROTEIN_RESPONSE
- GOBP_MYELOID_CELL_HOMEOSTASIS, GOBP_MYELOID_CELL_HOMEOSTASIS
- 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_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN, GOBP_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN
- GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
- GOBP_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA, GOBP_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN
- GOBP_ERYTHROCYTE_HOMEOSTASIS, GOBP_ERYTHROCYTE_HOMEOSTASIS
- GOBP_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
- GOBP_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION, GOBP_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION
- GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I, GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN
- GOBP_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION, GOBP_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION
- GOBP_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION
- GOBP_REGULATION_OF_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION
- GOBP_PEROXISOME_ORGANIZATION, GOBP_PEROXISOME_ORGANIZATION
- GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION
- GOBP_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GOBP_TRANSLESION_SYNTHESIS, GOBP_TRANSLESION_SYNTHESIS
- GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_I
- GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_GAP_FILLING, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_GAP_FILLING
- GOBP_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS, GOBP_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS
- GOBP_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR, GOBP_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR
- GOBP_PEROXISOMAL_TRANSPORT, GOBP_PEROXISOMAL_TRANSPORT
- GOBP_REGULATION_OF_CELLULAR_AMINE_METABOLIC_PROCESS, GOBP_REGULATION_OF_CELLULAR_AMINE_METABOLIC_PROCESS
- GOBP_ERROR_PRONE_TRANSLESION_SYNTHESIS, GOBP_ERROR_PRONE_TRANSLESION_SYNTHESIS
- GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_DUPLEX_UNWINDING, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_DUPLEX_UNWINDING
- GOBP_ESTABLISHMENT_OF_MITOCHONDRION_LOCALIZATION, GOBP_ESTABLISHMENT_OF_MITOCHONDRION_LOCALIZATION
- GOBP_TEMPERATURE_HOMEOSTASIS, GOBP_TEMPERATURE_HOMEOSTASIS
- GOBP_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRIN
- GOBP_CELL_REDOX_HOMEOSTASIS, GOBP_CELL_REDOX_HOMEOSTASIS
- GOBP_POSITIVE_REGULATION_OF_WNT_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_WNT_SIGNALING_PATHWAY
- GOBP_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_DEVELOPMENT, GOBP_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_DEVELOPMENT
- GOBP_CELLULAR_RESPONSE_TO_GLUCOSE_STARVATION, GOBP_CELLULAR_RESPONSE_TO_GLUCOSE_STARVATION
- GOBP_PROTEIN_HYDROXYLATION, GOBP_PROTEIN_HYDROXYLATION
- GOBP_CELL_MATURATION, GOBP_CELL_MATURATION
- GOBP_MORPHOGENESIS_OF_A_POLARIZED_EPITHELIUM, GOBP_MORPHOGENESIS_OF_A_POLARIZED_EPITHELIUM
- GOBP_REGULATION_OF_TRANSLATIONAL_INITIATION_IN_RESPONSE_TO_STRESS, GOBP_REGULATION_OF_TRANSLATIONAL_INITIATION_IN_RESPONSE_TO_STRESS
- GOBP_ESTABLISHMENT_OF_TISSUE_POLARITY, GOBP_ESTABLISHMENT_OF_TISSUE_POLARITY
- GOBP_POSITIVE_REGULATION_OF_CANONICAL_WNT_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_CANONICAL_WNT_SIGNALING_PATHWAY
- GOBP_PROTEASOME_ASSEMBLY, GOBP_PROTEASOME_ASSEMBLY
- GOBP_CHAPERONE_MEDIATED_PROTEIN_FOLDING, GOBP_CHAPERONE_MEDIATED_PROTEIN_FOLDING
- GOBP_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER, GOBP_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER
- GOBP_RESPONSE_TO_AMINO_ACID_STARVATION, GOBP_RESPONSE_TO_AMINO_ACID_STARVATION
- GOBP_REGULATION_OF_RNA_POLYMERASE_II_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING, GOBP_REGULATION_OF_RNA_POLYMERASE_II_REGULATORY_REGION_SEQUENCE_SPECIFIC
- GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS