N_1_RICH_GRANULE, GO_FICOLIN_1_RICH_GRANULE

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GO PEPTIDASE COMPLEX, GO PEPTIDASE COMPLEX
GO ANAPHASE PROMOTING COMPLEX DEPENDENT CATABOLIC PROCESS, GO ANAPHASE PROMOTING COMPLEX DEPENDENT CATABOLIC PROCESS
GO_ENDOPEPTIDASE_COMPLEX, GO_ENDOPEPTIDASE_COMPLEX
GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN, GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN
GO_AZUROPHIL_GRANULE, GO_AZUROPHIL_GRANULE
GO_POSITIVE_REGULATION_OF_DNA_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_DNA_METABOLIC_PROCESS
GO_FC_EPSILON_RECEPTOR_SIGNALING_PATHWAY, GO_FC_EPSILON_RECEPTOR_SIGNALING_PATHWAY
GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I, GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I
GO_DNA_BIOSYNTHETIC_PROCESS, GO_DNA_BIOSYNTHETIC_PROCESS
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- GO_LAMELLIPODIUM, GO_LAMELLIPODIUM GO_PROTEASOME_ACCESSORY_COMPLEX, GO_PROTEASOME_ACCESSORY_COMPLEX
- GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I, GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I
- GO_FC_RECEPTOR_SIGNALING_PATHWAY, GO_FC_RECEPTOR_SIGNALING_PATHWAY
- GO_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
- GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION, GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION
- GO_POSITIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS, GO_POSITIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS
- GO_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY, GO_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY
- GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS, GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS
- GO_BLOOD_MICROPARTICLE, GO_BLOOD_MICROPARTICLE
- GO_PIGMENT_GRANULE, GO_PIGMENT_GRANULE
- GO_T_CELL_RECEPTOR_SIGNALING_PATHWAY, GO_T_CELL_RECEPTOR_SIGNALING_PATHWAY
- GO_SITE_OF_DOUBLE_STRAND_BREAK, GO_SITE_OF_DOUBLE_STRAND_BREAK
- GO_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEUS, GO_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEUS
- GO_SITE_OF_POLARIZED_GROWTH, GO_SITE_OF_POLARIZED_GROWTH
- GO_HEAT_SHOCK_PROTEIN_BINDING, GO_HEAT_SHOCK_PROTEIN_BINDING
- GO_RIBONUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS, GO_RIBONUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS
- GO_CYSTEINE_TYPE_PEPTIDASE_ACTIVITY, GO_CYSTEINE_TYPE_PEPTIDASE_ACTIVITY
- GO_PROTEIN_IMPORT, GO_PROTEIN_IMPORT
- GO_RIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS, GO_RIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS
- GO_ORGANOPHOSPHATE_CATABOLIC_PROCESS, GO_ORGANOPHOSPHATE_CATABOLIC_PROCESS
- GO_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY, GO_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY
- GO_PYRIMIDINE_NUCLEOBASE_METABOLIC_PROCESS, GO_PYRIMIDINE_NUCLEOBASE_METABOLIC_PROCESS
- GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_OR_POLYSACCHARIDE_ANTIGEN_VIA_MHC_CLASS_II, GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_OR_POLYSACCHARIDE_ANTIGEN_

GO_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA, GO_REGULATION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA, GO_REGULATION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA, GO_REGULATION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA, GO_REGULATION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RNA_POLYMERASE_II_PROMOTER_IN_RNA_POLYMERASE_II_PROMOTER_IN_RNA_POLYMERASE_II_PROMOTER_IN_RNA_POLYMERASE_II_PROMOTER_IN_RNA_POLYMERASE_II_PROMOTER_IN_RNA_POLYMERASE_II_PROMOTER_IN_RNA_POLYMERASE_II_PROMOTER_IN_RNA_POLYMERASE_II_PROMOTER_IN_RNA_POLYMERASE_II_PROMOTER_IN_RNA_POLYMERASE_II_PROMOTER_IN_RNA_POLYMERASE_II_PROMOTER_IN_IN_RNA_POLYMERASE_II_PROMOTER_IN_IN_RNA_POLYMERASE_II_PROMOTER_IN

- GO_NADP_METABOLIC_PROCESS, GO_NADP_METABOLIC_PROCESS
- GO_REGULATION_OF_MACROPHAGE_CHEMOTAXIS, GO_REGULATION_OF_MACROPHAGE_CHEMOTAXIS
- GO_PODOSOME, GO_PODOSOME
- GO_CARBOHYDRATE_CATABOLIC_PROCESS, GO_CARBOHYDRATE_CATABOLIC_PROCESS
- GO_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS, GO_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS
- GO_TERTIARY_GRANULE, GO_TERTIARY_GRANULE
- GO_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GO_NUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS, GO_NUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS
- GO_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT, GO_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT
- GO_MHC_CLASS_II_PROTEIN_COMPLEX_BINDING, GO_MHC_CLASS_II_PROTEIN_COMPLEX_BINDING
- GO_ENDOCYTIC_VESICLE_LUMEN, GO_ENDOCYTIC_VESICLE_LUMEN
- GO_MAP_KINASE_ACTIVITY, GO_MAP_KINASE_ACTIVITY
- GO NUCLEOBASE CONTAINING SMALL MOLECULE BIOSYNTHETIC PROCESS, GO NUCLEOBASE CONTAINING SMALL MOLECULE BIOSYNTHETIC PROCESS
- GO_RAC_PROTEIN_SIGNAL_TRANSDUCTION, GO_RAC_PROTEIN_SIGNAL_TRANSDUCTION
- GO_ESTABLISHMENT_OF_MITOTIC_SPINDLE_LOCALIZATION, GO_ESTABLISHMENT_OF_MITOTIC_SPINDLE_LOCALIZATION
- GO_RNA_POLYMERASE_II_CTD_HEPTAPEPTIDE_REPEAT_KINASE_ACTIVITY, GO_RNA_POLYMERASE_II_CTD_HEPTAPEPTIDE_REPEAT_KINASE_ACTIVITY
- GO_POSITIVE_REGULATION_OF_TELOMERASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_TELOMERASE_ACTIVITY
- GO PROTEASOME REGULATORY PARTICLE BASE SUBCOMPLEX, GO PROTEASOME REGULATORY PARTICLE BASE SUBCOMPLEX
- GO UNFOLDED PROTEIN BINDING, GO UNFOLDED PROTEIN BINDING
- GO_NUCLEOSIDE_METABOLIC_PROCESS, GO_NUCLEOSIDE_METABOLIC_PROCESS
- GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_IN_RESPONSE_TO_STRESS, GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_IN_RESPONSE_TO_STRESS
- GO_CELLULAR_METABOLIC_COMPOUND_SALVAGE, GO_CELLULAR_METABOLIC_COMPOUND_SALVAGE
- GO NEGATIVE REGULATION OF FIBROBLAST PROLIFERATION, GO NEGATIVE REGULATION OF FIBROBLAST PROLIFERATION
- GO_EXECUTION_PHASE_OF_APOPTOSIS, GO_EXECUTION_PHASE_OF_APOPTOSIS
- GO_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION, GO_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION
- GO_PURINE_NUCLEOSIDE_METABOLIC_PROCESS, GO_PURINE_NUCLEOSIDE_METABOLIC_PROCESS
- GO_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY, GO_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY
- GO_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER
- GO_ESTABLISHMENT_OF_SPINDLE_ORIENTATION, GO_ESTABLISHMENT_OF_SPINDLE_ORIENTATION
- GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER
- GO_PURINE_NUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS, GO_PURINE_NUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS
- GO_PROTEASOME_ASSEMBLY, GO_PROTEASOME_ASSEMBLY
- GO_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION, GO_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION