

E_RECEPTOR_BINDING, GO_NUCLEAR_HORMONE_RECEPTOR_BINDING

- GO_POSITIVE_REGULATION_OF_HEMOPOIESIS, GO_POSITIVE_REGULATION_OF_HEMOPOIESIS
- GO_BRANCHING_MORPHOGENESIS_OF_AN_EPITHELIAL_TUBE, GO_BRANCHING_MORPHOGENESIS_OF_AN_EPITHELIAL_TUBE
- GO_RNA_POLYMERASE_II_REPRESSING_TRANSCRIPTION_FACTOR_BINDING, GO_RNA_POLYMERASE_II_REPRESSING_TRANSCRIPTION_FACTOR_BINDING
- GO_POSITIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION
- GO_NUCLEOSOME_BINDING, GO_NUCLEOSOME_BINDING
- GO_GLAND_MORPHOGENESIS, GO_GLAND_MORPHOGENESIS
- GO_CIRCADIAN_REGULATION_OF_GENE_EXPRESSION, GO_CIRCADIAN_REGULATION_OF_GENE_EXPRESSION
- GO_POSITIVE_REGULATION_OF_PROTEIN_BINDING, GO_POSITIVE_REGULATION_OF_PROTEIN_BINDING
- GO_CELL_FATE_DETERMINATION, GO_CELL_FATE_DETERMINATION
- GO_ADIPOSE_TISSUE_DEVELOPMENT, GO_ADIPOSE_TISSUE_DEVELOPMENT
- GO_REGULATION_OF_NOTCH_SIGNALING_PATHWAY, GO_REGULATION_OF_NOTCH_SIGNALING_PATHWAY
- GO_MEDIATOR_COMPLEX, GO_MEDIATOR_COMPLEX
- GO_EMBRYONIC_PLACENTA_DEVELOPMENT, GO_EMBRYONIC_PLACENTA_DEVELOPMENT
- GO_REGULATION_OF_CHONDROCYTE_DIFFERENTIATION, GO_REGULATION_OF_CHONDROCYTE_DIFFERENTIATION
- GO_REGULATION_OF_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE, GO_REGULATION_OF_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE
- GO_NEGATIVE_REGULATION_OF_GENE_SILENCING_BY_MIRNA, GO_NEGATIVE_REGULATION_OF_GENE_SILENCING_BY_MIRNA
- GO_OUTFLOW_TRACT_MORPHOGENESIS, GO_OUTFLOW_TRACT_MORPHOGENESIS
- GO_REPRESSING_TRANSCRIPTION_FACTOR_BINDING, GO_REPRESSING_TRANSCRIPTION_FACTOR_BINDING
- GO_FACE_DEVELOPMENT, GO_FACE_DEVELOPMENT
- GO_NEGATIVE_REGULATION_OF_PROTEIN_BINDING, GO_NEGATIVE_REGULATION_OF_PROTEIN_BINDING
- GO_POSITIVE_REGULATION_OF_ORGANELLE_ASSEMBLY, GO_POSITIVE_REGULATION_OF_ORGANELLE_ASSEMBLY
- GO_POSITIVE_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION
- GO_METANEPHRIC_NEPHRON_DEVELOPMENT, GO_METANEPHRIC_NEPHRON_DEVELOPMENT
- GO_CELLULAR_RESPONSE_TO_KETONE, GO_CELLULAR_RESPONSE_TO_KETONE
- GO_SIGNAL_TRANSDUCTION_INVOLVED_IN_REGULATION_OF_GENE_EXPRESSION, GO_SIGNAL_TRANSDUCTION_INVOLVED_IN_REGULATION_OF_GENE_EXPRESSION
- GO_CARDIAC_CHAMBER_MORPHOGENESIS, GO_CARDIAC_CHAMBER_MORPHOGENESIS
- GO_NOTCH_SIGNALING_PATHWAY, GO_NOTCH_SIGNALING_PATHWAY
- GO_AXIS_ELONGATION, GO_AXIS_ELONGATION
- GO_NEGATIVE_REGULATION_OF_PRODUCTION_OF_MIRNAS_INVOLVED_IN_GENE_SILENCING_BY_MIRNA, GO_NEGATIVE_REGULATION_OF_PRODUCTION_OF_MIRNAS_INVOLVED_IN_GENE_SILENCING_BY_MIRNA
- GO_LENS_FIBER_CELL_DIFFERENTIATION, GO_LENS_FIBER_CELL_DIFFERENTIATION