

GO_STRUCTURAL_CONSTITUENT, GO_EXTRACELLULAR_MATRIX_STRUCTURAL_CONSTITUENT

- GO_HEPARIN_BINDING, GO_HEPARIN_BINDING
- GO_INTEGRIN_BINDING, GO_INTEGRIN_BINDING
- GO_VACUOLAR_LUMEN, GO_VACUOLAR_LUMEN
- GO_BASEMENT_MEMBRANE, GO_BASEMENT_MEMBRANE
- GO_COLLAGEN_METABOLIC_PROCESS, GO_COLLAGEN_METABOLIC_PROCESS
- GO_AMINOGLYCAN_METABOLIC_PROCESS, GO_AMINOGLYCAN_METABOLIC_PROCESS
- GO_CARBOHYDRATE_DERIVATIVE_CATABOLIC_PROCESS, GO_CARBOHYDRATE_DERIVATIVE_CATABOLIC_PROCESS
- GO_COLLAGEN_TRIMER, GO_COLLAGEN_TRIMER
- GO_GROWTH_FACTOR_BINDING, GO_GROWTH_FACTOR_BINDING
- GO_LYSOSOMAL_LUMEN, GO_LYSOSOMAL_LUMEN
- GO_PEPTIDE_CROSS_LINKING, GO_PEPTIDE_CROSS_LINKING
- GO_COLLAGEN_BINDING, GO_COLLAGEN_BINDING
- GO_EXTRACELLULAR_MATRIX_BINDING, GO_EXTRACELLULAR_MATRIX_BINDING
- GO_SCAVENGER_RECEPTOR_ACTIVITY, GO_SCAVENGER_RECEPTOR_ACTIVITY
- GO_EYE_MORPHOGENESIS, GO_EYE_MORPHOGENESIS
- GO_NEGATIVE_REGULATION_OF_WOUND_HEALING, GO_NEGATIVE_REGULATION_OF_WOUND_HEALING
- GO_BLOOD_MICROPARTICLE, GO_BLOOD_MICROPARTICLE
- GO_EXTRACELLULAR_MATRIX_ASSEMBLY, GO_EXTRACELLULAR_MATRIX_ASSEMBLY
- GO_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS, GO_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS
- GO_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE, GO_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE
- GO_NEPHRON_DEVELOPMENT, GO_NEPHRON_DEVELOPMENT
- GO_AMINOGLYCAN_BIOSYNTHETIC_PROCESS, GO_AMINOGLYCAN_BIOSYNTHETIC_PROCESS
- GO TRABECULA FORMATION, GO TRABECULA FORMATION
- GO_AMINOGLYCAN_CATABOLIC_PROCESS, GO_AMINOGLYCAN_CATABOLIC_PROCESS
- GO_ENDODERM_FORMATION, GO_ENDODERM_FORMATION
- GO_PLATELET_DEGRANULATION, GO_PLATELET_DEGRANULATION
- GO_BONE TRABECULA MORPHOGENESIS, GO_BONE TRABECULA MORPHOGENESIS
- GO_FIBRINOLYSIS, GO_FIBRINOLYSIS
- GO_SULFUR_COMPOUND_BIOSYNTHETIC_PROCESS, GO_SULFUR_COMPOUND_BIOSYNTHETIC_PROCESS
- GO_PLASMINOGEN_ACTIVATION, GO_PLASMINOGEN_ACTIVATION
- GO_EMBRYONIC_EYE_MORPHOGENESIS, GO_EMBRYONIC_EYE_MORPHOGENESIS
- GO_CELLULAR_RESPONSE_TO_AMINO_ACID_STIMULUS, GO_CELLULAR_RESPONSE_TO_AMINO_ACID_STIMULUS
- GO_PLATELET_AGGREGATION, GO_PLATELET_AGGREGATION
- GO_POSITIVE_REGULATION_OF_VASOCONSTRICTION, GO_POSITIVE_REGULATION_OF_VASOCONSTRICTION
- GO_NEGATIVE_REGULATION_OF_COAGULATION, GO_NEGATIVE_REGULATION_OF_COAGULATION
- GO_ENAMEL_MINERALIZATION, GO_ENAMEL_MINERALIZATION
- GO_ENDODERM_DEVELOPMENT, GO_ENDODERM_DEVELOPMENT
- GO_ZYMOGEN_ACTIVATION, GO_ZYMOGEN_ACTIVATION
- GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_WOUNDING, GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_WOUNDING
- GO_CHONDROITIN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS, GO_CHONDROITIN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS
- GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY, GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THR
- GO_REGULATION_OF_COAGULATION, GO_REGULATION_OF_COAGULATION