

ANTIOXIDANT_ACTIVITY, GOMF_ANTIOXIDANT_ACTIVITY

- GOBP_UNSATURATED_FATTY_ACID_METABOLIC_PROCESS, GOBP_UNSATURATED_FATTY_ACID_METABOLIC_PROCESS
- GOBP_DETOXIFICATION, GOBP_DETOXIFICATION
- GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PEROXIDE_AS_ACCEPTOR, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PEROXIDE_AS_ACCEPTOR
- GOBP_CELLULAR_OXIDANT_DETOXIFICATION, GOBP_CELLULAR_OXIDANT_DETOXIFICATION
- GOBP_CELLULAR_RESPONSE_TO_TOXIC_SUBSTANCE, GOBP_CELLULAR_RESPONSE_TO_TOXIC_SUBSTANCE
- GOBP_ICOSANOID_METABOLIC_PROCESS, GOBP_ICOSANOID_METABOLIC_PROCESS
- GOBP_OLEFINIC_COMPOUND_METABOLIC_PROCESS, GOBP_OLEFINIC_COMPOUND_METABOLIC_PROCESS
- GOBP_ARACHIDONIC_ACID_METABOLIC_PROCESS, GOBP_ARACHIDONIC_ACID_METABOLIC_PROCESS
- GOMF_IRON_ION_BINDING, GOMF_IRON_ION_BINDING
- GOBP_GLUTATHIONE_DERIVATIVE_METABOLIC_PROCESS, GOBP_GLUTATHIONE_DERIVATIVE_METABOLIC_PROCESS
- GOBP_RESPONSE_TO_XENOBIOTIC_STIMULUS, GOBP_RESPONSE_TO_XENOBIOTIC_STIMULUS
- GOMF_GLUTATHIONE_TRANSFERASE_ACTIVITY, GOMF_GLUTATHIONE_TRANSFERASE_ACTIVITY
- GOBP_UV_PROTECTION, GOBP_UV_PROTECTION
- GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_TO_CHEMICAL_STIMULUS, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_T
- HP_GOITER, HP_GOITER
- GOBP_REGULATION_OF_OXIDOREDUCTASE_ACTIVITY, GOBP_REGULATION_OF_OXIDOREDUCTASE_ACTIVITY
- GOMF_TRANSFERASE_ACTIVITY_TRANSFERRING_ALKYL_OR_ARYL_OTHER_THAN_METHYL_GROUPS, GOMF_TRANSFERASE_ACTIVITY_TRANSFERRING_ALKYL_OR_ARYL_OTHER_THAN_METHYL_GROUPS
- GOMF_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY, GOMF_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY
- GOCC_NUCLEAR_OUTER_MEMBRANE, GOCC_NUCLEAR_OUTER_MEMBRANE
- GOBP_LEYDIG_CELL_DIFFERENTIATION, GOBP_LEYDIG_CELL_DIFFERENTIATION