F_WATER_LOSS_VIA_SKIN, GO_REGULATION_OF_WATER_LOSS_VIA_SKIN

GO_WATER_HOMEOSTASIS, GO_WATER_HOMEOSTASIS
GO_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS, GO_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS
GO_TIGHT_JUNCTION, GO_TIGHT_JUNCTION
GO_SPHINGOLIPID_BIOSYNTHETIC_PROCESS, GO_SPHINGOLIPID_BIOSYNTHETIC_PROCESS
GO_LONG_CHAIN_FATTY_ACID_BIOSYNTHETIC_PROCESS, GO_LONG_CHAIN_FATTY_ACID_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEMBLY, GO_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEM
GO_MEMBRANE_LIPID_BIOSYNTHETIC_PROCESS, GO_MEMBRANE_LIPID_BIOSYNTHETIC_PROCESS
GO_SPHINGOID_METABOLIC_PROCESS, GO_SPHINGOID_METABOLIC_PROCESS
GO_REGULATION_OF_RETINOIC_ACID_RECEPTOR_SIGNALING_PATHWAY, GO_REGULATION_OF_RETINOIC_ACID_RECEPTOR_S
GO_PROXIMAL_DISTAL_PATTERN_FORMATION, GO_PROXIMAL_DISTAL_PATTERN_FORMATION
GO_ISOPRENOID_CATABOLIC_PROCESS, GO_ISOPRENOID_CATABOLIC_PROCESS
GO_LIPOSACCHARIDE_METABOLIC_PROCESS, GO_LIPOSACCHARIDE_METABOLIC_PROCESS
GO_ARACHIDONIC_ACID_METABOLIC_PROCESS, GO_ARACHIDONIC_ACID_METABOLIC_PROCESS
GO_TERPENOID_CATABOLIC_PROCESS, GO_TERPENOID_CATABOLIC_PROCESS
GO_RESPONSE_TO_INTERLEUKIN_18, GO_RESPONSE_TO_INTERLEUKIN_18
GO_ICOSANOID_METABOLIC_PROCESS, GO_ICOSANOID_METABOLIC_PROCESS
GO_LONG_CHAIN_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS, GO_LONG_CHAIN_FATTY_ACYL_COA_BIOSYNTHETIC_PROCES
GO_IRON_ION_BINDING, GO_IRON_ION_BINDING
GO_RETINOIC_ACID_METABOLIC_PROCESS, GO_RETINOIC_ACID_METABOLIC_PROCESS
GO_RETINOIC_ACID_4_HYDROXYLASE_ACTIVITY, GO_RETINOIC_ACID_4_HYDROXYLASE_ACTIVITY