ROMEDINU_IN_T_CELL_LINE_6H_DN, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_DN
GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN
GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP
GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_DN
GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN, GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN
GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP
GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP, GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP
GSE3982_NEUTROPHIL_VS_EFF_MEMORY_CD4_TCELL_UP, GSE3982_NEUTROPHIL_VS_EFF_MEMORY_CD4_TCELL_UP
GSE38696_LIGHT_ZONE_VS_DARK_ZONE_BCELL_UP, GSE38696_LIGHT_ZONE_VS_DARK_ZONE_BCELL_UP
GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN
GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP
GSE42021_TCONV_PLN_VS_TREG_PRECURSORS_THYMUS_UP, GSE42021_TCONV_PLN_VS_TREG_PRECURSORS_THYMUS_UP
GSE25087_TREG_VS_TCONV_FETUS_UP, GSE25087_TREG_VS_TCONV_FETUS_UP
GSE40666_UNTREATED_VS_IFNA_STIM_STAT4_KO_EFFECTOR_CD8_TCELL_90MIN_DN, GSE40666_UNTREATED_VS_IFNA_STIM_STAT4_KO_EFFECTOR_CD8_TCELL_90MIN_DN
GSE3982_CTRL_VS_IGE_STIM_MAST_CELL_UP, GSE3982_CTRL_VS_IGE_STIM_MAST_CELL_UP
GSE3982_DC_VS_EFF_MEMORY_CD4_TCELL_UP, GSE3982_DC_VS_EFF_MEMORY_CD4_TCELL_UP
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_UP
GSE3982_BASOPHIL_VS_TH1_UP, GSE3982_BASOPHIL_VS_TH1_UP
GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP, GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP
GSE15767_MED_VS_SCS_MAC_LN_UP, GSE15767_MED_VS_SCS_MAC_LN_UP
GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN, GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN
GOBP_REGULATION_OF_GENE_SILENCING_BY_RNA, GOBP_REGULATION_OF_GENE_SILENCING_BY_RNA
GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_UP, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_UP
GSE37301_RAG2_KO_VS_RAG2_AND_ETS1_KO_NK_CELL_UP, GSE37301_RAG2_KO_VS_RAG2_AND_ETS1_KO_NK_CELL_UP
STEARMAN_LUNG_CANCER_EARLY_VS_LATE_UP, STEARMAN_LUNG_CANCER_EARLY_VS_LATE_UP
PECE_MAMMARY_STEM_CELL_DN, PECE_MAMMARY_STEM_CELL_DN
GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_UP, GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_UP
MIR194_5P, MIR194_5P
GSE42021_TREG_PLN_VS_TREG_PRECURSORS_THYMUS_UP, GSE42021_TREG_PLN_VS_TREG_PRECURSORS_THYMUS_UP
MIR500A_5P, MIR500A_5P
GSE37532_VISCERAL_ADIPOSE_TISSUE_VS_LN_DERIVED_TREG_CD4_TCELL_UP, GSE37532_VISCERAL_ADIPOSE_TISSUE_VS_LN_DERIVED_TREG_CD4_TCELL_UP
MIR548AI_MIR570_5P, MIR548AI_MIR570_5P
GOMF_INTEGRIN_BINDING, GOMF_INTEGRIN_BINDING
GSE7548_DAY7_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP, GSE7548_DAY7_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP
MIR6505_5P, MIR6505_5P
GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN
GSE17721_CTRL_VS_LPS_2H_BMDC_DN, GSE17721_CTRL_VS_LPS_2H_BMDC_DN
GOBP_INTEGRIN_MEDIATED_SIGNALING_PATHWAY, GOBP_INTEGRIN_MEDIATED_SIGNALING_PATHWAY
HP_THORACIC_HYPOPLASIA, HP_THORACIC_HYPOPLASIA
MIR4727_5P, MIR4727_5P
MIR585_5P, MIR585_5P
GOBP_MEGAKARYOCYTE_DIFFERENTIATION, GOBP_MEGAKARYOCYTE_DIFFERENTIATION
MIR488_3P, MIR488_3P
MIR8065, MIR8065
GOMF_HORMONE_RECEPTOR_BINDING, GOMF_HORMONE_RECEPTOR_BINDING
SENESE_HDAC2_TARGETS_UP, SENESE_HDAC2_TARGETS_UP
MIR891A_3P, MIR891A_3P
MIR12132, MIR12132
GOBP_POSITIVE_REGULATION_OF_HEMOPOIESIS, GOBP_POSITIVE_REGULATION_OF_HEMOPOIESIS
REACTOME_SUMOYLATION_OF_TRANSCRIPTION_COFACTORS, REACTOME_SUMOYLATION_OF_TRANSCRIPTION_COFACTORS
MIR642B_5P, MIR642B_5P
MIR4293, MIR4293
GOBP_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION
MIR558, MIR558
MIR7849_3P, MIR7849_3P
GSE32423_CTRL_VS_IL7_MEMORY_CD8_TCELL_DN, GSE32423_CTRL_VS_IL7_MEMORY_CD8_TCELL_DN
GOBP_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE, GOBP_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE
MIR892A, MIR892A
GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_DN
MIR4286, MIR4286
GRAHAM_CML_DIVIDING_VS_NORMAL_QUIESCENT_DN, GRAHAM_CML_DIVIDING_VS_NORMAL_QUIESCENT_DN
GOCC_CUL3_RING_UBIQUITIN_LIGASE_COMPLEX, GOCC_CUL3_RING_UBIQUITIN_LIGASE_COMPLEX
HALLMARK_COAGULATION, HALLMARK_COAGULATION
MIR4456, MIR4456
MIR597_5P, MIR597_5P
MIR495_5P, MIR495_5P
MIR3682_3P, MIR3682_3P
KEGG_AXON_GUIDANCE, KEGG_AXON_GUIDANCE
HP_STAGE_5_CHRONIC_KIDNEY_DISEASE, HP_STAGE_5_CHRONIC_KIDNEY_DISEASE
GOBP_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS, GOBP_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS
MIR5004_5P, MIR5004_5P
TCTATGA_MIR376A_MIR376B, TCTATGA_MIR376A_MIR376B
MIR584_5P, MIR584_5P
GOBP_AMINOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_AMINOGLYCAN_BIOSYNTHETIC_PROCESS
GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN
MIR339_5P, MIR339_5P
MOREAUX_B_LYMPHOCYTE_MATURATION_BY_TACI_UP, MOREAUX_B_LYMPHOCYTE_MATURATION_BY_TACI_UP
GRAHAM_NORMAL_QUIESCENT_VS_NORMAL_DIVIDING_UP, GRAHAM_NORMAL_QUIESCENT_VS_NORMAL_DIVIDING_UP
MIR148B_5P, MIR148B_5P
PID_RETINOIC_ACID_PATHWAY, PID_RETINOIC_ACID_PATHWAY
CASORELLI_APL_SECONDARY_VS_DE_NOVO_UP, CASORELLI_APL_SECONDARY_VS_DE_NOVO_UP
GUO_HEX_TARGETS_UP, GUO_HEX_TARGETS_UP
GOBP_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
MIR629_5P, MIR629_5P
BCAT_GDS748_DN, BCAT_GDS748_DN
GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY, GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY
COATES_MACROPHAGE_M1_VS_M2_DN, COATES_MACROPHAGE_M1_VS_M2_DN
COMP1_01, COMP1_01
GSE34205_RSV_VS_FLU_INF_INFANT_PBMC_UP, GSE34205_RSV_VS_FLU_INF_INFANT_PBMC_UP
MIR1178_3P, MIR1178_3P
GOBP_POSITIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT
MIR1912_3P, MIR1912_3P
MIR4797_3P, MIR4797_3P
GOBP_LIPID_STORAGE, GOBP_LIPID_STORAGE
GOBP_REGULATION_OF_WOUND_HEALING, GOBP_REGULATION_OF_WOUND_HEALING
GOMF_CORECEPTOR_ACTIVITY, GOMF_CORECEPTOR_ACTIVITY
MIR548Q, MIR548Q
WP_HEDGEHOG_SIGNALING_PATHWAY, WP_HEDGEHOG_SIGNALING_PATHWAY
WP_NUCLEOTIDEBINDING_OLIGOMERIZATION_DOMAIN_NOD_PATHWAY, WP_NUCLEOTIDEBINDING_OLIGOMERIZATION_DOMAIN_NOD_PATHWAY
GOBP_PROTEOGLYCAN_METABOLIC_PROCESS, GOBP_PROTEOGLYCAN_METABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_CELL_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_CELL_DEVELOPMENT
SANA_TNF_SIGNALING_DN, SANA_TNF_SIGNALING_DN
GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS
HP_ABNORMAL_PLATELET_FUNCTION, HP_ABNORMAL_PLATELET_FUNCTION
HARALAMBIEVA_PBMC_M_M_R_II_AGE_13_16YO_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_POST_TOP_DEG_UP, HARALAMBIEVA_PBMC_M_M_R_II_AGE_13_16YO_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_POST_TOP_DEG_UP
MIR6866_5P, MIR6866_5P
MIR4274, MIR4274
MIR1224_5P, MIR1224_5P
GOBP_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY, GOBP_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY
KYNG_DNA_DAMAGE_BY_4NQO_OR_UV, KYNG_DNA_DAMAGE_BY_4NQO_OR_UV
GOMF_EXTRACELLULAR_MATRIX_BINDING, GOMF_EXTRACELLULAR_MATRIX_BINDING
REACTOME_SLC_TRANSPORTER_DISORDERS, REACTOME_SLC_TRANSPORTER_DISORDERS
HP_INAPPROPRIATE_BEHAVIOR, HP_INAPPROPRIATE_BEHAVIOR
MIR6851_3P, MIR6851_3P
MIR12113, MIR12113
GOBP_POLYSACCHARIDE_METABOLIC_PROCESS, GOBP_POLYSACCHARIDE_METABOLIC_PROCESS
MIR5588_3P, MIR5588_3P
HP_PLEURAL_EFFUSION, HP_PLEURAL_EFFUSION
BROWNE_HCMV_INFECTION_16HR_DN, BROWNE_HCMV_INFECTION_16HR_DN
GOBP_NEGATIVE_REGULATION_OF_NERVOUS_SYSTEM_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_NERVOUS_SYSTEM_DEVELOPMENT
BECKER_TAMOXIFEN_RESISTANCE_DN, BECKER_TAMOXIFEN_RESISTANCE_DN
GOBP_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS
MIR595, MIR595
BIOCARTA_PPARA_PATHWAY, BIOCARTA_PPARA_PATHWAY
GOMF_COLLAGEN_BINDING, GOMF_COLLAGEN_BINDING
HP_ABNORMALITY_OF_SKULL_OSSIFICATION, HP_ABNORMALITY_OF_SKULL_OSSIFICATION
GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION
GOBP_NEGATIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
GTAAGAT_MIR200A, GTAAGAT_MIR200A
MCCLUNG_COCAIN_REWARD_4WK, MCCLUNG_COCAIN_REWARD_4WK
GOBP_MACROPHAGE_MIGRATION, GOBP_MACROPHAGE_MIGRATION
MIR3160_3P, MIR3160_3P
GOBP_CHONDROITIN_SULFATE_BIOSYNTHETIC_PROCESS, GOBP_CHONDROITIN_SULFATE_BIOSYNTHETIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_LIPID_LOCALIZATION, GOBP_POSITIVE_REGULATION_OF_LIPID_LOCALIZATION
GOBP_POLYSACCHARIDE_CATABOLIC_PROCESS, GOBP_POLYSACCHARIDE_CATABOLIC_PROCESS
GOBP_MACROPHAGE_CHEMOTAXIS, GOBP_MACROPHAGE_CHEMOTAXIS
MIR3972, MIR3972
GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY
GOBP_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
GOBP_WNT_SIGNALING_PATHWAY_CALCIUM_MODULATING_PATHWAY, GOBP_WNT_SIGNALING_PATHWAY_CALCIUM_MODULATING_PATHWAY
GOBP_INTERLEUKIN_1_BETA_PRODUCTION, GOBP_INTERLEUKIN_1_BETA_PRODUCTION
GOCC_DENDRITE_CYTOPLASM, GOCC_DENDRITE_CYTOPLASM
GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION
GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN, GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN
REACTOME_CA2_PATHWAY, REACTOME_CA2_PATHWAY
GOBP_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
GOBP_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING, GOBP_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING
GOBP_RESPONSE_TO_PROGESTERONE, GOBP_RESPONSE_TO_PROGESTERONE
GOBP_REGULATION_OF_ORGANIC_ACID_TRANSPORT, GOBP_REGULATION_OF_ORGANIC_ACID_TRANSPORT
GOBP_INTERLEUKIN_1_PRODUCTION, GOBP_INTERLEUKIN_1_PRODUCTION
MIR4712_3P, MIR4712_3P
HP_SHORT_RIBS, HP_SHORT_RIBS
GOBP_APOPTOTIC_CELL_CLEARANCE, GOBP_APOPTOTIC_CELL_CLEARANCE
KANG_DOXORUBICIN_RESISTANCE_DN, KANG_DOXORUBICIN_RESISTANCE_DN
GOBP_REGULATION_OF_LIPID_LOCALIZATION, GOBP_REGULATION_OF_LIPID_LOCALIZATION
REACTOME_CHONDROITIN_SULFATE_BIOSYNTHESIS, REACTOME_CHONDROITIN_SULFATE_BIOSYNTHESIS
GOMF_PHOSPHATIDYLSERINE_BINDING, GOMF_PHOSPHATIDYLSERINE_BINDING
HP_CORPUS_CALLOSUM_ATROPHY, HP_CORPUS_CALLOSUM_ATROPHY
EVI1_02, EVI1_02
GOBP_NEGATIVE_REGULATION_OF_TRANSLATIONAL_INITIATION, GOBP_NEGATIVE_REGULATION_OF_TRANSLATIONAL_INITIATION
HP_BLUE_IRIDES, HP_BLUE_IRIDES
TATCTGG_MIR488, TATCTGG_MIR488
KANG_IMMORTALIZED_BY_TERT_UP, KANG_IMMORTALIZED_BY_TERT_UP
BIOCARTA_UCALPAIN_PATHWAY, BIOCARTA_UCALPAIN_PATHWAY
MIR7112_3P, MIR7112_3P
PID_IL12_STAT4_PATHWAY, PID_IL12_STAT4_PATHWAY
GOMF_STEROL_TRANSPORTER_ACTIVITY, GOMF_STEROL_TRANSPORTER_ACTIVITY
PID_S1P_S1P3_PATHWAY, PID_S1P_S1P3_PATHWAY
GOBP_POSITIVE_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GOBP_POSITIVE_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS
GOCC_MITOTIC_SPINDLE_MICROTUBULE, GOCC_MITOTIC_SPINDLE_MICROTUBULE
GOBP_REGULATION_OF_HORMONE_METABOLIC_PROCESS, GOBP_REGULATION_OF_HORMONE_METABOLIC_PROCESS
GOCC_AXONAL_GROWTH_CONE, GOCC_AXONAL_GROWTH_CONE
GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
MIR601, MIR601
KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE, KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE
HP_ABNORMALITY_OF_THE_UMBILICAL_CORD, HP_ABNORMALITY_OF_THE_UMBILICAL_CORD
HP_IMPAIRED_RISTOCETIN_INDUCED_PLATELET_AGGREGATION, HP_IMPAIRED_RISTOCETIN_INDUCED_PLATELET_AGGREGATION
HP_ABNORMAL_PLATELET_MEMBRANE_PROTEIN_EXPRESSION, HP_ABNORMAL_PLATELET_MEMBRANE_PROTEIN_EXPRESSION
GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_1_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_1_PRODUCTION
HP_RECURRENT_INFECTION_OF_THE_GASTROINTESTINAL_TRACT, HP_RECURRENT_INFECTION_OF_THE_GASTROINTESTINAL_TRACT
GOMF_1_ACYLGLYCEROPHOSPHOCHOLINE_O_ACYLTRANSFERASE_ACTIVITY, GOMF_1_ACYLGLYCEROPHOSPHOCHOLINE_O_ACYLTRANSFERASE_ACTIVITY
HP_ABNORMALITY_OF_THE_5TH_METACARPAL, HP_ABNORMALITY_OF_THE_5TH_METACARPAL
HP_ABNORMAL_PLATELET_MORPHOLOGY, HP_ABNORMAL_PLATELET_MORPHOLOGY
GOBP_NEGATIVE_REGULATION_OF_LIPID_STORAGE, GOBP_NEGATIVE_REGULATION_OF_LIPID_STORAGE
GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_MIGRATION, GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_MIGRATION
GOBP_NEGATIVE_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING, GOBP_NEGATIVE_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING
GOBP_THYMUS_DEVELOPMENT, GOBP_THYMUS_DEVELOPMENT
MIR512_5P, MIR512_5P
MIR3677_5P, MIR3677_5P
VERRECCHIA_RESPONSE_TO_TGFB1_C1, VERRECCHIA_RESPONSE_TO_TGFB1_C1
GOBP_TISSUE_REGENERATION, GOBP_TISSUE_REGENERATION
GOBP_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GOBP_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS
GOBP_POSITIVE_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS
REACTOME_RUNX2_REGULATES_BONE_DEVELOPMENT, REACTOME_RUNX2_REGULATES_BONE_DEVELOPMENT
REACTOME_REGULATION_OF_RUNX1_EXPRESSION_AND_ACTIVITY, REACTOME_REGULATION_OF_RUNX1_EXPRESSION_AND_ACTIVITY
HP_ABNORMAL_PLACENTA_MORPHOLOGY, HP_ABNORMAL_PLACENTA_MORPHOLOGY
MARTIN_NFKB_TARGETS_UP, MARTIN_NFKB_TARGETS_UP
WP_SEROTONIN_TRANSPORTER_ACTIVITY, WP_SEROTONIN_TRANSPORTER_ACTIVITY
GUENTHER_GROWTH_SPHERICAL_VS_ADHERENT_DN, GUENTHER_GROWTH_SPHERICAL_VS_ADHERENT_DN
GLI1_UP.V1_DN, GLI1_UP.V1_DN
GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_DEPENDENT_DECAY, GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_DEPEND
HASLINGER_B_CLL_WITH_MUTATED_VH_GENES, HASLINGER_B_CLL_WITH_MUTATED_VH_GENES
GOBP_MONOCYTE_CHEMOTACTIC_PROTEIN_1_PRODUCTION, GOBP_MONOCYTE_CHEMOTACTIC_PROTEIN_1_PRODUCTION
GOBP_POSITIVE_REGULATION_OF_FIBROBLAST_MIGRATION, GOBP_POSITIVE_REGULATION_OF_FIBROBLAST_MIGRATION
HP_APLASIA_HYPOPLASIA_OF_THE_4TH_METACARPAL, HP_APLASIA_HYPOPLASIA_OF_THE_4TH_METACARPAL
MIR4487, MIR4487
WP_ANGIOGENESIS, WP_ANGIOGENESIS
FERRARI_RESPONSE_TO_FENRETINIDE_UP, FERRARI_RESPONSE_TO_FENRETINIDE_UP
GOBP_LABYRINTHINE_LAYER_DEVELOPMENT, GOBP_LABYRINTHINE_LAYER_DEVELOPMENT
GOBP_REGULATION_OF_MACROPHAGE_CHEMOTAXIS, GOBP_REGULATION_OF_MACROPHAGE_CHEMOTAXIS
HP_REDUCED_SPERM_MOTILITY, HP_REDUCED_SPERM_MOTILITY
BIOCARTA_ERK_PATHWAY, BIOCARTA_ERK_PATHWAY
WP_MFAP5MEDIATED_OVARIAN_CANCER_CELL_MOTILITY_AND_INVASIVENESS, WP_MFAP5MEDIATED_OVARIAN_CANCER_CELL_MOTILITY_AND_INVASIVENESS
GOBP_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS, GOBP_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT
GOBP_ENERGY_RESERVE_METABOLIC_PROCESS, GOBP_ENERGY_RESERVE_METABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION
GOBP_CHROMATIN_SILENCING_AT_TELOMERE, GOBP_CHROMATIN_SILENCING_AT_TELOMERE
REACTOME_FORMATION_OF_APOPTOSOME, REACTOME_FORMATION_OF_APOPTOSOME
HP_ABNORMALITY_OF_THE_4TH_METACARPAL, HP_ABNORMALITY_OF_THE_4TH_METACARPAL
BIOCARTA_PTEN_PATHWAY, BIOCARTA_PTEN_PATHWAY
HALLMARK_ANGIOGENESIS, HALLMARK_ANGIOGENESIS
GOMF_IONOTROPIC_GLUTAMATE_RECEPTOR_BINDING, GOMF_IONOTROPIC_GLUTAMATE_RECEPTOR_BINDING
ABE_VEGFA_TARGETS_2HR, ABE_VEGFA_TARGETS_2HR
HP_PROMINENT_UMBILICUS, HP_PROMINENT_UMBILICUS
GOBP_PHOSPHATIDYLSERINE_METABOLIC_PROCESS, GOBP_PHOSPHATIDYLSERINE_METABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_CELL_JUNCTION_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_CELL_JUNCTION_ASSEMBLY
REACTOME_GLUCAGON_LIKE_PEPTIDE_1_GLP1_REGULATES_INSULIN_SECRETION, REACTOME_GLUCAGON_LIKE_PEPTIDE_1_GLP1_REGULATES_INSULIN_SECRETION
HP_CONE_CONE_ROD_DYSTROPHY, HP_CONE_CONE_ROD_DYSTROPHY
GOBP_KYNURENINE_METABOLIC_PROCESS, GOBP_KYNURENINE_METABOLIC_PROCESS
WP_PREGNANE_X_RECEPTOR_PATHWAY, WP_PREGNANE_X_RECEPTOR_PATHWAY
GOBP_RESPONSE_TO_CATECHOLAMINE, GOBP_RESPONSE_TO_CATECHOLAMINE
GOBP_NEUROINFLAMMATORY_RESPONSE, GOBP_NEUROINFLAMMATORY_RESPONSE
GOBP_PLASMINOGEN_ACTIVATION, GOBP_PLASMINOGEN_ACTIVATION
GOBP_NAD_BIOSYNTHETIC_PROCESS, GOBP_NAD_BIOSYNTHETIC_PROCESS
GOBP_REGULATION_OF_PRODUCTION_OF_SMALL_RNA_INVOLVED_IN_GENE_SILENCING_BY_RNA, GOBP_REGULATION_OF_PRODUCTION_OF_SMALL_RNA_INVOLVED_IN_GENE_SILENCING_BY_RNA
GOCC_HAUS_COMPLEX, GOCC_HAUS_COMPLEX
GOBP_NEGATIVE_REGULATION_OF_PRODUCTION_OF_MIRNAS_INVOLVED_IN_GENE_SILENCING_BY_MIRNA, GOBP_NEGATIVE_REGULATION_OF_PRODUCTION_OF_MIRNAS_INVOLVED_IN_GENE_SILENCING_BY_MIRNA
GOBP_REGULATION_OF_FATTY_ACID_TRANSPORT, GOBP_REGULATION_OF_FATTY_ACID_TRANSPORT
PID_ANTHRAX_PATHWAY, PID_ANTHRAX_PATHWAY
GOBP_COLLAGEN_BIOSYNTHETIC_PROCESS, GOBP_COLLAGEN_BIOSYNTHETIC_PROCESS
MAHAJAN_RESPONSE_TO_IL1A_UP, MAHAJAN_RESPONSE_TO_IL1A_UP
BIOCARTA_BARR_MAPK_PATHWAY, BIOCARTA_BARR_MAPK_PATHWAY
HP_LARGE_PLACENTA, HP_LARGE_PLACENTA
GOMF_CELL_MATRIX_ADHESION_MEDIATOR_ACTIVITY, GOMF_CELL_MATRIX_ADHESION_MEDIATOR_ACTIVITY
BIOCARTA_MSP_PATHWAY, BIOCARTA_MSP_PATHWAY
HP_ELEVATED_DIASTOLIC_BLOOD_PRESSURE, HP_ELEVATED_DIASTOLIC_BLOOD_PRESSURE
GOBP_POSITIVE_REGULATION_OF_MONOCYTE_CHEMOTACTIC_PROTEIN_1_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_MONOCYTE_CHEMOTACTIC_PROTEIN_1_PRODUCTION
MCBRYAN_TERMINAL_END_BUD_DN, MCBRYAN_TERMINAL_END_BUD_DN
PHONG_TNF_TARGETS_DN, PHONG_TNF_TARGETS_DN
HP_REFLEX_SEIZURE, HP_REFLEX_SEIZURE
HP_ABNORMALITY_OF_THE_CHOROID_PLEXUS, HP_ABNORMALITY_OF_THE_CHOROID_PLEXUS
GOBP_REGULATION_OF_GRANULOCYTE_CHEMOTAXIS, GOBP_REGULATION_OF_GRANULOCYTE_CHEMOTAXIS
GOMF_GLUTAMATE_RECEPTOR_BINDING, GOMF_GLUTAMATE_RECEPTOR_BINDING
WP_STEROID_BIOSYNTHESIS, WP_STEROID_BIOSYNTHESIS
GOBP_REGULATION_OF_AMINO_ACID_TRANSPORT, GOBP_REGULATION_OF_AMINO_ACID_TRANSPORT
TSENG_ADIPOGENIC_POTENTIAL_DN, TSENG_ADIPOGENIC_POTENTIAL_DN
GOCC_MICROVILLUS_MEMBRANE, GOCC_MICROVILLUS_MEMBRANE
PID_INTEGRIN3_PATHWAY, PID_INTEGRIN3_PATHWAY
GOBP_DENDRITE_EXTENSION, GOBP_DENDRITE_EXTENSION
HP_ANKYLOSIS, HP_ANKYLOSIS
GOBP_POSITIVE_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION, GOBP_POSITIVE_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION
GOMF_GAMMA_CATENIN_BINDING, GOMF_GAMMA_CATENIN_BINDING
WP_OSTEOPONTIN_SIGNALING, WP_OSTEOPONTIN_SIGNALING
GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_CHEMOTAXIS, GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_CHEMOTAXIS
BIOCARTA_BARRESTIN_SRC_PATHWAY, BIOCARTA_BARRESTIN_SRC_PATHWAY
PID_S1P_S1P1_PATHWAY, PID_S1P_S1P1_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION
GOBP_CELLULAR_RESPONSE_TO_CADMIUM_ION, GOBP_CELLULAR_RESPONSE_TO_CADMIUM_ION
PID_SYNDECAN_4_PATHWAY, PID_SYNDECAN_4_PATHWAY
REACTOME_RUNX2_REGULATES_OSTEOBLAST_DIFFERENTIATION, REACTOME_RUNX2_REGULATES_OSTEOBLAST_DIFFERENTIATION
WP_PRADERWILLI_AND_ANGELMAN_SYNDROME, WP_PRADERWILLI_AND_ANGELMAN_SYNDROME
KIM_ALL_DISORDERS_DURATION_CORR_UP, KIM_ALL_DISORDERS_DURATION_CORR_UP
GOBP_CARDIAC_MUSCLE_FIBER_DEVELOPMENT, GOBP_CARDIAC_MUSCLE_FIBER_DEVELOPMENT
GOBP_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEMBLY, GOBP_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEMBLY
GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_MIGRATION, GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_MIGRATION
GOBP_CELL_CELL_ADHESION_MEDIATED_BY_INTEGRIN, GOBP_CELL_CELL_ADHESION_MEDIATED_BY_INTEGRIN
GOBP_FIBRINOLYSIS, GOBP_FIBRINOLYSIS
GOBP_REPRODUCTIVE_BEHAVIOR, GOBP_REPRODUCTIVE_BEHAVIOR
GOBP_MAINTENANCE_OF_DNA_REPEAT_ELEMENTS, GOBP_MAINTENANCE_OF_DNA_REPEAT_ELEMENTS
GOBP_HYPOXIA_INDUCIBLE_FACTOR_1ALPHA_SIGNALING_PATHWAY, GOBP_HYPOXIA_INDUCIBLE_FACTOR_1ALPHA_SIGNALING_PATHWAY
GOBP_MUSCLE_FIBER_DEVELOPMENT, GOBP_MUSCLE_FIBER_DEVELOPMENT
HP_ATYPICAL_ABSENCE_SEIZURE, HP_ATYPICAL_ABSENCE_SEIZURE
GOBP_ACIDIC_AMINO_ACID_TRANSPORT, GOBP_ACIDIC_AMINO_ACID_TRANSPORT
GOBP_REGULATION_OF_CONNECTIVE_TISSUE_REPLACEMENT, GOBP_REGULATION_OF_CONNECTIVE_TISSUE_REPLACEMENT
GOBP_POSITIVE_REGULATION_OF_AMINE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_AMINE_TRANSPORT
GOBP_REGULATION_OF_NEUROTRANSMITTER_UPTAKE, GOBP_REGULATION_OF_NEUROTRANSMITTER_UPTAKE
GOBP_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS, GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS
HP_CEPHALOHEMATOMA, HP_CEPHALOHEMATOMA
REACTOME_VEGF_LIGAND_RECEPTOR_INTERACTIONS, REACTOME_VEGF_LIGAND_RECEPTOR_INTERACTIONS
FAN_EMBRYONIC_CTX_MICROGLIA_2, FAN_EMBRYONIC_CTX_MICROGLIA_2
HP_DERMAL_TRANSLUCENCY, HP_DERMAL_TRANSLUCENCY
RIZ_ERYTHROID_DIFFERENTIATION_HEMGN, RIZ_ERYTHROID_DIFFERENTIATION_HEMGN
GOBP_POSITIVE_REGULATION_OF_MONOCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MONOCYTE_DIFFERENTIATION
GOBP_AROMATIC_AMINO_ACID_FAMILY_CATABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_CATABOLIC_PROCESS
PID_S1P_S1P4_PATHWAY, PID_S1P_S1P4_PATHWAY
PID_VEGF_VEGFR_PATHWAY, PID_VEGF_VEGFR_PATHWAY
GOBP_REGULATION_OF_NATURAL_KILLER_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_NATURAL_KILLER_CELL_DIFFERENTIATION
GOBP_POSITIVE_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOP
BIOCARTA_AGR_PATHWAY, BIOCARTA_AGR_PATHWAY
GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY, GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY
HP_STIFF_SKIN, HP_STIFF_SKIN
GOBP_REGULATION_OF_MACROPHAGE_PROLIFERATION, GOBP_REGULATION_OF_MACROPHAGE_PROLIFERATION
HP_EEG_WITH_FOCAL_SHARP_SLOW_WAVES, HP_EEG_WITH_FOCAL_SHARP_SLOW_WAVES
GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS
GOBP_FOAM_CELL_DIFFERENTIATION, GOBP_FOAM_CELL_DIFFERENTIATION
GOBP_NEUROTRANSMITTER_REUPTAKE, GOBP_NEUROTRANSMITTER_REUPTAKE
HP_GENERALIZED_CEREBRAL_ATROPHY_HYPOPLASIA, HP_GENERALIZED_CEREBRAL_ATROPHY_HYPOPLASIA
GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT
GOBP_LEUKOCYTE_ADHESION_TO_VASCULAR_ENDOTHELIAL_CELL, GOBP_LEUKOCYTE_ADHESION_TO_VASCULAR_ENDOTHELIAL_CELL
KEGG_ETHER_LIPID_METABOLISM, KEGG_ETHER_LIPID_METABOLISM
GOMF_LAMININ_BINDING, GOMF_LAMININ_BINDING
GOBP_DICARBOXYLIC_ACID_TRANSPORT, GOBP_DICARBOXYLIC_ACID_TRANSPORT
GOBP_REGULATION_OF_DENDRITIC_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_DENDRITIC_CELL_DIFFERENTIATION
IRITANI_MAD1_TARGETS_UP, IRITANI_MAD1_TARGETS_UP
HP_GASTRITIS, HP_GASTRITIS
GOMF_GABA_RECEPTOR_BINDING, GOMF_GABA_RECEPTOR_BINDING
GOBP_IMMATURE_T_CELL_PROLIFERATION, GOBP_IMMATURE_T_CELL_PROLIFERATION
REACTOME_GLUCAGON_SIGNALING_IN_METABOLIC_REGULATION, REACTOME_GLUCAGON_SIGNALING_IN_METABOLIC_REGULATION
GOBP_MATING_BEHAVIOR, GOBP_MATING_BEHAVIOR
WP_LTF_DANGER_SIGNAL_RESPONSE_PATHWAY, WP_LTF_DANGER_SIGNAL_RESPONSE_PATHWAY
GOBP_REGULATORY_T_CELL_DIFFERENTIATION, GOBP_REGULATORY_T_CELL_DIFFERENTIATION
HP_THROMBOPHLEBITIS, HP_THROMBOPHLEBITIS
GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_RECEPTOR_2_BINDING, GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_RECEPTOR_2_BINDING
GOMF_OPIOID_RECEPTOR_BINDING, GOMF_OPIOID_RECEPTOR_BINDING
HP_OVERTUBULATED_LONG_BONES, HP_OVERTUBULATED_LONG_BONES
GOBP_SMOOTH_MUSCLE_ADAPTATION, GOBP_SMOOTH_MUSCLE_ADAPTATION
GOBP_REGULATION_OF_MACROPHAGE_FUSION, GOBP_REGULATION_OF_MACROPHAGE_FUSION
GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_BINDING, GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_BINDING
GOMF_FIBRINOGEN_BINDING, GOMF_FIBRINOGEN_BINDING
REACTOME_MET_INTERACTS_WITH_TNS_PROTEINS, REACTOME_MET_INTERACTS_WITH_TNS_PROTEINS
PALOMERO_GSI_SENSITIVITY_DN, PALOMERO_GSI_SENSITIVITY_DN
GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_UPTAKE, GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_UPTAKE
HP_LOSS_OF_SUBCUTANEOUS_ADIPOSE_TISSUE_IN_LIMBS, HP_LOSS_OF_SUBCUTANEOUS_ADIPOSE_TISSUE_IN_LIMBS
HP_ENTEROCOLITIS, HP_ENTEROCOLITIS
GOBP_TRANSITION_BETWEEN_FAST_AND_SLOW_FIBER, GOBP_TRANSITION_BETWEEN_FAST_AND_SLOW_FIBER
GOCC_FIBRINOGEN_COMPLEX, GOCC_FIBRINOGEN_COMPLEX
BIOCARTA_EPHA4_PATHWAY, BIOCARTA_EPHA4_PATHWAY
HP_ENTRAPMENT_NEUROPATHY, HP_ENTRAPMENT_NEUROPATHY
WP_NAD_BIOSYNTHESIS_II_FROM_TRYPTOPHAN, WP_NAD_BIOSYNTHESIS_II_FROM_TRYPTOPHAN
REACTOME_RUNX2_REGULATES_GENES_INVOLVED_IN_CELL_MIGRATION, REACTOME_RUNX2_REGULATES_GENES_INVOLVED_IN_CELL_MIGRATION
WP_TRYPTOPHAN_CATABOLISM_LEADING_TO_NAD_PRODUCTION, WP_TRYPTOPHAN_CATABOLISM_LEADING_TO_NAD_PRODUCTION
GRAHAM_CML_QUIESCENT_VS_CML_DIVIDING_UP, GRAHAM_CML_QUIESCENT_VS_CML_DIVIDING_UP
REACTOME_REMOVAL_OF_AMINOTERMINAL_PROPEPTIDES_FROM_GAMMA_CARBOXYLATED_PROTEINS, REACTOME_REMOVAL_OF_AMINOTERMINAL_PROPEPTIDES_FROM_GAMMA_CARBOXYLATED_PROTEINS
FREDERICK_PRKCI_TARGETS, FREDERICK_PRKCI_TARGETS
REACTOME_SIGNAL_ATTENUATION, REACTOME_SIGNAL_ATTENUATION
GOBP_TRYPTOPHAN_CATABOLIC_PROCESS, GOBP_TRYPTOPHAN_CATABOLIC_PROCESS
GOBP_ACTIVATION_OF_PROTEIN_KINASE_B_ACTIVITY, GOBP_ACTIVATION_OF_PROTEIN_KINASE_B_ACTIVITY
GOBP_MACROPHAGE_FUSION, GOBP_MACROPHAGE_FUSION
GOBP_RESPONSE_TO_HYDROSTATIC_PRESSURE, GOBP_RESPONSE_TO_HYDROSTATIC_PRESSURE
GOBP_LOOP_OF_HENLE_DEVELOPMENT, GOBP_LOOP_OF_HENLE_DEVELOPMENT
GOBP_METANEPHRIC_PART_OF_URETERIC_BUD_DEVELOPMENT, GOBP_METANEPHRIC_PART_OF_URETERIC_BUD_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_ORGANIC_ACID_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_ORGANIC_ACID_TRANSPORT
HINATA_NFKB_TARGETS_KERATINOCYTE_DN, HINATA_NFKB_TARGETS_KERATINOCYTE_DN
GOBP_NEGATIVE_REGULATION_OF_SMOOTH_MUSCLE_CONTRACTION, GOBP_NEGATIVE_REGULATION_OF_SMOOTH_MUSCLE_CONTRACTION
GOBP_RESPONSE_TO_NICOTINE, GOBP_RESPONSE_TO_NICOTINE
GOMF_C_X_C_CHEMOKINE_BINDING, GOMF_C_X_C_CHEMOKINE_BINDING
GOMF_COLLAGEN_BINDING_INVOLVED_IN_CELL_MATRIX_ADHESION, GOMF_COLLAGEN_BINDING_INVOLVED_IN_CELL_MATRIX_ADHESION
WP_TCA_CYCLE_NUTRIENT_UTILIZATION_AND_INVASIVENESS_OF_OVARIAN_CANCER, WP_TCA_CYCLE_NUTRIENT_UTILIZATION_AND_INVASIVENESS_OF_OVARIAN_CANCER
HP_INAPPROPRIATE_SEXUAL_BEHAVIOR, HP_INAPPROPRIATE_SEXUAL_BEHAVIOR
GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS, GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS
GOBP_CARBON_CATABOLITE_REGULATION_OF_TRANSCRIPTION, GOBP_CARBON_CATABOLITE_REGULATION_OF_TRANSCRIPTION
GOCC_CHECKPOINT_CLAMP_COMPLEX, GOCC_CHECKPOINT_CLAMP_COMPLEX
GOCC_IPAF_INFLAMMASOME_COMPLEX, GOCC_IPAF_INFLAMMASOME_COMPLEX
GOMF_CALMODULIN_DEPENDENT_PROTEIN_PHOSPHATASE_ACTIVITY, GOMF_CALMODULIN_DEPENDENT_PROTEIN_PHOSPHATASE_ACTIVITY
GOBP_PROTEIN_SULFATION, GOBP_PROTEIN_SULFATION
GOBP_NEPHRIC_DUCT_DEVELOPMENT, GOBP_NEPHRIC_DUCT_DEVELOPMENT
GOMF_INSULIN_LIKE_GROWTH_FACTOR_RECEPTOR_BINDING, GOMF_INSULIN_LIKE_GROWTH_FACTOR_RECEPTOR_BINDING
WP_KYNURENINE_PATHWAY_AND_LINKS_TO_CELLULAR_SENESCENCE, WP_KYNURENINE_PATHWAY_AND_LINKS_TO_CELLULAR_SENESCENCE
GOBP_MESODERMAL_CELL_DIFFERENTIATION, GOBP_MESODERMAL_CELL_DIFFERENTIATION
GOBP_REGULATION_OF_CARDIAC_MUSCLE_CELL_PROLIFERATION, GOBP_REGULATION_OF_CARDIAC_MUSCLE_CELL_PROLIFERATION
MATZUK_CUMULUS_EXPANSION, MATZUK_CUMULUS_EXPANSION
HP_APLASIA_HYPOPLASIA_OF_THE_COLON, HP_APLASIA_HYPOPLASIA_OF_THE_COLON
GOBP_IMMATURE_T_CELL_PROLIFERATION_IN_THYMUS, GOBP_IMMATURE_T_CELL_PROLIFERATION_IN_THYMUS
GOBP_POSITIVE_REGULATION_OF_UNSATURATED_FATTY_ACID_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_UNSATURATED_FATTY_ACID_BIOSYNTHETIC_PROCESS
GOBP_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT, GOBP_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT
GOBP_NEGATIVE_REGULATION_OF_VASOCONSTRICTION, GOBP_NEGATIVE_REGULATION_OF_VASOCONSTRICTION
TURASHVILI_BREAST_CARCINOMA_DUCTAL_VS_LOBULAR_DN, TURASHVILI_BREAST_CARCINOMA_DUCTAL_VS_LOBULAR_DN
GOBP_REGULATION_OF_SPONTANEOUS_SYNAPTIC_TRANSMISSION, GOBP_REGULATION_OF_SPONTANEOUS_SYNAPTIC_TRANSMISSION
SASAKI_TARGETS_OF_TP73_AND_TP63, SASAKI_TARGETS_OF_TP73_AND_TP63
GOBP_OVULATION_FROM_OVARIAN_FOLLICLE, GOBP_OVULATION_FROM_OVARIAN_FOLLICLE
GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_DIFFERENTIATION
SMID_BREAST_CANCER_RELAPSE_IN_LIVER_UP, SMID_BREAST_CANCER_RELAPSE_IN_LIVER_UP
GOBP_TRYPTOPHAN_METABOLIC_PROCESS, GOBP_TRYPTOPHAN_METABOLIC_PROCESS
LIU_CMYB_TARGETS_DN, LIU_CMYB_TARGETS_DN
GOBP_CALCIUM_ION_IMPORT_INTO_SARCOPLASMIC_RETICULUM, GOBP_CALCIUM_ION_IMPORT_INTO_SARCOPLASMIC_RETICULUM
GOBP_TAURINE_TRANSPORT, GOBP_TAURINE_TRANSPORT
GOBP_REGULATION_OF_DNA_LIGATION, GOBP_REGULATION_OF_DNA_LIGATION
GOBP_CHORIO_ALLANTOIC_FUSION, GOBP_CHORIO_ALLANTOIC_FUSION
GOBP_REGULATION_OF_ADIPONECTIN_SECRETION, GOBP_REGULATION_OF_ADIPONECTIN_SECRETION
GOBP_KERATINOCYTE_APOPTOTIC_PROCESS, GOBP_KERATINOCYTE_APOPTOTIC_PROCESS
REACTOME_AQUAPORIN_MEDIATED_TRANSPORT, REACTOME_AQUAPORIN_MEDIATED_TRANSPORT
GOBP_RESPONSE_TO_DOPAMINE, GOBP_RESPONSE_TO_DOPAMINE
GOBP_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION
GOBP_TROPHECTODERMAL_CELL_DIFFERENTIATION, GOBP_TROPHECTODERMAL_CELL_DIFFERENTIATION
GOBP_SPONTANEOUS_SYNAPTIC_TRANSMISSION, GOBP_SPONTANEOUS_SYNAPTIC_TRANSMISSION
GOBP_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION, GOBP_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION
SA_MMP_CYTOKINE_CONNECTION, SA_MMP_CYTOKINE_CONNECTION
GOBP_REGULATION_OF_DENDRITIC_CELL_ANTIGEN_PROCESSING_AND_PRESENTATION, GOBP_REGULATION_OF_DENDRITIC_CELL_ANTIGEN_PROCESSING_AND_PRESENTATION
BIOCARTA_IL5_PATHWAY, BIOCARTA_IL5_PATHWAY
GOBP_T_HELPER_1_CELL_DIFFERENTIATION, GOBP_T_HELPER_1_CELL_DIFFERENTIATION
GOBP_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS, GOBP_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS
GOBP_POSITIVE_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE, GOBP_POSITIVE_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE
GOBP_REGULATION_OF_PHOTORECEPTOR_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_PHOTORECEPTOR_CELL_DIFFERENTIATION
GOCC_FILTRATION_DIAPHRAGM, GOCC_FILTRATION_DIAPHRAGM
REACTOME_INTERLEUKIN_1_PROCESSING, REACTOME_INTERLEUKIN_1_PROCESSING
GOBP_RESPONSE_TO_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR, GOBP_RESPONSE_TO_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR
GOBP_REGULATION_OF_LYMPHOID_PROGENITOR_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_LYMPHOID_PROGENITOR_CELL_DIFFERENTIATION
GOBP_NEGATIVE_REGULATION_OF_CGMP_MEDIATED_SIGNALING, GOBP_NEGATIVE_REGULATION_OF_CGMP_MEDIATED_SIGNALING
GOBP_HYALOID_VASCULAR_PLEXUS_REGRESSION, GOBP_HYALOID_VASCULAR_PLEXUS_REGRESSION
GOBP_NEGATIVE_REGULATION_OF_ENDOTHELIAL_CELL_CHEMOTAXIS, GOBP_NEGATIVE_REGULATION_OF_ENDOTHELIAL_CELL_CHEMOTAXIS
GOMF_GAMMA_AMINOBUTYRIC_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_GAMMA_AMINOBUTYRIC_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_PROSTAGLANDIN_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_PROSTAGLANDIN_BIOSYNTHETIC_PROCESS
GOBP_REGULATION_OF_CDC42_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_CDC42_PROTEIN_SIGNAL_TRANSDUCTION
REACTOME_SIGNALING_BY_MAPK_MUTANTS, REACTOME_SIGNALING_BY_MAPK_MUTANTS
GOBP_NEGATIVE_REGULATION_OF_MAST_CELL_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_MAST_CELL_ACTIVATION
GOCC_CHROMAFFIN_GRANULE, GOCC_CHROMAFFIN_GRANULE
GOMF_OUTWARD_RECTIFIER_POTASSIUM_CHANNEL_ACTIVITY, GOMF_OUTWARD_RECTIFIER_POTASSIUM_CHANNEL_ACTIVITY
GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION
GOBP_REGULATION_OF_SYNAPTIC_VESICLE_CYCLE, GOBP_REGULATION_OF_SYNAPTIC_VESICLE_CYCLE
GOBP_L_SERINE_BIOSYNTHETIC_PROCESS, GOBP_L_SERINE_BIOSYNTHETIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTIO
WACKER_HYPOXIA_TARGETS_OF_VHL, WACKER_HYPOXIA_TARGETS_OF_VHL
HP_INTESTINAL_HYPOPLASIA, HP_INTESTINAL_HYPOPLASIA
GOBP_POSITIVE_REGULATION_OF_CHEMOKINE_C_X_C_MOTIF_LIGAND_2_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_CHEMOKINE_C_X_C_MOTIF_LIGAND_2_PRODUCTION
GOBP_ADENYLATE_CYCLASE_ACTIVATING_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY, GOBP_ADENYLATE_CYCLASE_ACTIVATING_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY
GOBP_L_SERINE_METABOLIC_PROCESS, GOBP_L_SERINE_METABOLIC_PROCESS
HP_ABNORMAL_BLOOD_CHLORIDE_CONCENTRATION, HP_ABNORMAL_BLOOD_CHLORIDE_CONCENTRATION
GOBP_NATURAL_KILLER_CELL_DIFFERENTIATION, GOBP_NATURAL_KILLER_CELL_DIFFERENTIATION
GOBP_DENDRITIC_CELL_ANTIGEN_PROCESSING_AND_PRESENTATION, GOBP_DENDRITIC_CELL_ANTIGEN_PROCESSING_AND_PRESENTATION
GOBP_REGULATION_OF_KERATINOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_KERATINOCYTE_DIFFERENTIATION
chr4p15, chr4p15
GOBP_KIDNEY_MESENCHYME_DEVELOPMENT, GOBP_KIDNEY_MESENCHYME_DEVELOPMENT
GOCC_GTPASE_COMPLEX, GOCC_GTPASE_COMPLEX
GOBP_AMINO_ACID_NEUROTRANSMITTER_REUPTAKE, GOBP_AMINO_ACID_NEUROTRANSMITTER_REUPTAKE
REN_MIF_TARGETS_DN, REN_MIF_TARGETS_DN
HP_DUMBBELL_SHAPED_LONG_BONE, HP_DUMBBELL_SHAPED_LONG_BONE
GOBP_T_HELPER_1_CELL_CYTOKINE_PRODUCTION, GOBP_T_HELPER_1_CELL_CYTOKINE_PRODUCTION
GOBP_NEGATIVE_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS
REACTOME_RUNX2_REGULATES_CHONDROCYTE_MATURATION, REACTOME_RUNX2_REGULATES_CHONDROCYTE_MATURATION
HP_HELICOBACTER_PYLORI_INFECTION, HP_HELICOBACTER_PYLORI_INFECTION
HP_INCREASED_FIBULAR_DIAMETER, HP_INCREASED_FIBULAR_DIAMETER
GOBP_TRYPTOPHAN_CATABOLIC_PROCESS_TO_KYNURENINE, GOBP_TRYPTOPHAN_CATABOLIC_PROCESS_TO_KYNURENINE
GOBP_POSITIVE_REGULATION_OF_IMMATURE_T_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_IMMATURE_T_CELL_PROLIFERATION
GOBP_QUINOLINATE_METABOLIC_PROCESS, GOBP_QUINOLINATE_METABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_DNA_LIGATION, GOBP_POSITIVE_REGULATION_OF_DNA_LIGATION
GOBP_CELLULAR_RESPONSE_TO_WATER_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_WATER_STIMULUS
GOBP_REGULATION_OF_RETINAL_GANGLION_CELL_AXON_GUIDANCE, GOBP_REGULATION_OF_RETINAL_GANGLION_CELL_AXON_GUIDANCE
GOBP_MOTOR_NEURON_MIGRATION, GOBP_MOTOR_NEURON_MIGRATION
HP_FOOT_ACROOSTEOLYSIS, HP_FOOT_ACROOSTEOLYSIS
GOBP_CALMODULIN_DEPENDENT_KINASE_SIGNALING_PATHWAY, GOBP_CALMODULIN_DEPENDENT_KINASE_SIGNALING_PATHWAY
HP_PSEUDOHYPOALDOSTERONISM, HP_PSEUDOHYPOALDOSTERONISM
GOBP_POSITIVE_REGULATION_OF_GLUTAMATE_SECRETION, GOBP_POSITIVE_REGULATION_OF_GLUTAMATE_SECRETION
GOBP_MESENDODERM_DEVELOPMENT, GOBP_MESENDODERM_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_CELL_ADHESION_MOLECULE_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_CELL_ADHESION_MOLECULE_PRODUCTION
GOCC_CHROMAFFIN_GRANULE_MEMBRANE, GOCC_CHROMAFFIN_GRANULE_MEMBRANE
GOBP_DE_NOVO_NAD_BIOSYNTHETIC_PROCESS, GOBP_DE_NOVO_NAD_BIOSYNTHETIC_PROCESS
GOBP_REGULATION_OF_STORE_OPERATED_CALCIUM_ENTRY, GOBP_REGULATION_OF_STORE_OPERATED_CALCIUM_ENTRY
GOBP_CHEMOKINE_C_X_C_MOTIF_LIGAND_2_PRODUCTION, GOBP_CHEMOKINE_C_X_C_MOTIF_LIGAND_2_PRODUCTION
REACTOME_GLUCAGON_TYPE_LIGAND_RECEPTORS, REACTOME_GLUCAGON_TYPE_LIGAND_RECEPTORS
GOBP_B_CELL_RECEPTOR_SIGNALING_PATHWAY, GOBP_B_CELL_RECEPTOR_SIGNALING_PATHWAY
GOBP_VASCULAR_ENDOTHELIAL_CELL_PROLIFERATION, GOBP_VASCULAR_ENDOTHELIAL_CELL_PROLIFERATION
DESCARTES_FETAL_INTESTINE_LYMPHATIC_ENDOTHELIAL_CELLS, DESCARTES_FETAL_INTESTINE_LYMPHATIC_ENDOTHELIAL_CELLS
GOBP_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION, GOBP_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION
CLAUS_PGR_POSITIVE_MENINGIOMA_DN, CLAUS_PGR_POSITIVE_MENINGIOMA_DN
GOBP_RENAL_SYSTEM_VASCULATURE_MORPHOGENESIS, GOBP_RENAL_SYSTEM_VASCULATURE_MORPHOGENESIS
REACTOME_FIBRONECTIN_MATRIX_FORMATION, REACTOME_FIBRONECTIN_MATRIX_FORMATION
HP_BAND_KERATOPATHY, HP_BAND_KERATOPATHY
GOBP_METANEPHRIC_S_SHAPED_BODY_MORPHOGENESIS, GOBP_METANEPHRIC_S_SHAPED_BODY_MORPHOGENESIS
GOBP_NEURAL_CREST_CELL_MIGRATION_INVOLVED_IN_AUTONOMIC_NERVOUS_SYSTEM_DEVELOPMENT, GOBP_NEURAL_CREST_CELL_MIGRATION_INVOLVED_IN_AUTONOMIC_NERVOUS_SYSTEM_DEVELOPMENT
GOMF_CORTICOTROPIN_RELEASING_HORMONE_RECEPTOR_BINDING, GOMF_CORTICOTROPIN_RELEASING_HORMONE_RECEPTOR_BINDING
GALI_TP53_TARGETS_APOPTOTIC_DN, GALI_TP53_TARGETS_APOPTOTIC_DN
GOBP_NEGATIVE_REGULATION_OF_TYPE_B_PANCREATIC_CELL_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_TYPE_B_PANCREATIC_CELL_APOPTOTIC_PROCESS
GOCC_INTEGRAL_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_ENDOPLASMIC_RETICULUM_MEMBRANE, GOCC_INTEGRAL_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_ENDOPLASMIC_RETICULUM_MEMBRANE
GOBP_REGULATION_OF_EPIDERMIS_DEVELOPMENT, GOBP_REGULATION_OF_EPIDERMIS_DEVELOPMENT
GOBP_MATING, GOBP_MATING
GOBP_NEGATIVE_REGULATION_OF_OSTEOBLAST_PROLIFERATION, GOBP_NEGATIVE_REGULATION_OF_OSTEOBLAST_PROLIFERATION
HP_CENTRAL_OPACIFICATION_OF_THE_CORNEA, HP_CENTRAL_OPACIFICATION_OF_THE_CORNEA
GOBP_CARDIAC_NEURAL_CREST_CELL_DIFFERENTIATION_INVOLVED_IN_HEART_DEVELOPMENT, GOBP_CARDIAC_NEURAL_CREST_CELL_DIFFERENTIATION_INVOLVED_IN_HEART_DEVELOPMENT
HP_SMALL_PLACENTA, HP_SMALL_PLACENTA
GOBP_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CONTRACTION, GOBP_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CONTRACTION
GOBP_SYMPATHETIC_GANGLION_DEVELOPMENT, GOBP_SYMPATHETIC_GANGLION_DEVELOPMENT
GOBP_REGULATION_OF_MACROPHAGE_DERIVED_FOAM_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_MACROPHAGE_DERIVED_FOAM_CELL_DIFFERENTIATION
HP_MENOMETRORRHAGIA, HP_MENOMETRORRHAGIA
GOBP_POSITIVE_REGULATION_OF_AXON_EXTENSION_INVOLVED_IN_AXON_GUIDANCE, GOBP_POSITIVE_REGULATION_OF_AXON_EXTENSION_INVOLVED_IN_AXON_GUIDANCE
GOBP_REGULATION_OF_CALCIDIOL_1_MONOOXYGENASE_ACTIVITY, GOBP_REGULATION_OF_CALCIDIOL_1_MONOOXYGENASE_ACTIVITY
GOBP_REGULATION_OF_INWARD_RECTIFIER_POTASSIUM_CHANNEL_ACTIVITY, GOBP_REGULATION_OF_INWARD_RECTIFIER_POTASSIUM_CHANNEL_ACTIVITY
REACTOME_DEFECTIVE_FACTOR_VIII_CAUSES_HEMOPHILIA_A, REACTOME_DEFECTIVE_FACTOR_VIII_CAUSES_HEMOPHILIA_A
GOBP_REGULATION_OF_SKELETAL_MUSCLE_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_SKELETAL_MUSCLE_CELL_DIFFERENTIATION
chr1q43, chr1q43
GOBP_MONOCYTE_CHEMOTAXIS, GOBP_MONOCYTE_CHEMOTAXIS
BIOCARTA_TFF_PATHWAY, BIOCARTA_TFF_PATHWAY
GOBP_NEUROTRANSMITTER_UPTAKE, GOBP_NEUROTRANSMITTER_UPTAKE
MIR3186_5P, MIR3186_5P
HP_SHORT_UMBILICAL_CORD, HP_SHORT_UMBILICAL_CORD
HP_CONGENITAL_PSEUDOARTHROSIS_OF_THE_CLAVICLE, HP_CONGENITAL_PSEUDOARTHROSIS_OF_THE_CLAVICLE
GOBP_REGULATION_OF_INOSITOL_1_4_5_TRISPHOSPHATE_SENSITIVE_CALCIUM_RELEASE_CHANNEL_ACTIVITY, GOBP_REGULATION_OF_INOSITOL_1_4_5_TRISPHOSPHATE_SENSITIVE_CALCIUM_RELEASE_CHANNEL_ACTIVITY
GOBP_REGULATION_OF_GAP_JUNCTION_ASSEMBLY, GOBP_REGULATION_OF_GAP_JUNCTION_ASSEMBLY
REACTOME_REELIN_SIGNALLING_PATHWAY, REACTOME_REELIN_SIGNALLING_PATHWAY
GOBP_S_SHAPED_BODY_MORPHOGENESIS, GOBP_S_SHAPED_BODY_MORPHOGENESIS
WP_ASPIRIN_AND_MIRNAS, WP_ASPIRIN_AND_MIRNAS
GOBP_MYOBLAST_DIFFERENTIATION, GOBP_MYOBLAST_DIFFERENTIATION
WP_RAS_AND_BRADYKININ_PATHWAYS_IN_COVID19, WP_RAS_AND_BRADYKININ_PATHWAYS_IN_COVID19
HP_IMPACTED_TOOTH, HP_IMPACTED_TOOTH
GOBP_POSITIVE_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_MIGRATION, GOBP_POSITIVE_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_MIGRATION
GOBP_GROWTH_PLATE_CARTILAGE_DEVELOPMENT, GOBP_GROWTH_PLATE_CARTILAGE_DEVELOPMENT
GOBP_NEGATIVE_REGULATION_OF_DENDRITE_MORPHOGENESIS, GOBP_NEGATIVE_REGULATION_OF_DENDRITE_MORPHOGENESIS
GOBP_REGULATION_OF_NITRIC_OXIDE_MEDIATED_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_NITRIC_OXIDE_MEDIATED_SIGNAL_TRANSDUCTION
GOBP_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_ACTIVITY
LE_SKI_TARGETS_UP, LE_SKI_TARGETS_UP
BIOCARTA_RACC_PATHWAY, BIOCARTA_RACC_PATHWAY
GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_DERIVED_FOAM_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_DERIVED_FOAM_CELL_DIFFERENTIATION
GOCC_INTRINSIC_COMPONENT_OF_EXTERNAL_SIDE_OF_PLASMA_MEMBRANE, GOCC_INTRINSIC_COMPONENT_OF_EXTERNAL_SIDE_OF_PLASMA_MEMBRANE