

GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN
GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_DN, GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_DN
GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP, GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP
GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_DN, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_DN
GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_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
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_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP
GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN, GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP
GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP, GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP
GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN, GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN
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_BASOPHIL_VS_TH1_UP, GSE3982_BASOPHIL_VS_TH1_UP
GSE25087_TREG_VS_TCONV_FETUS_UP, GSE25087_TREG_VS_TCONV_FETUS_UP
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP
GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_UP
PECE_MAMMARY_STEM_CELL_DN, PECE_MAMMARY_STEM_CELL_DN
GSE41867_DAY15_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN, GSE41867_DAY15_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN
GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP, GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP
MIR548A1_MIR570_5P, MIR548A1_MIR570_5P
GAUSSMANN_MLL_AF4_FUSION_TARGETS_A_UP, GAUSSMANN_MLL_AF4_FUSION_TARGETS_A_UP
GSE37301_RAG2_KO_VS_RAG2_AND_ETSI1_KO_NK_CELL_UP, GSE37301_RAG2_KO_VS_RAG2_AND_ETSI1_KO_NK_CELL_UP
MIR558, MIR558
MIR6505_5P, MIR6505_5P
MIR548AO_5P, MIR548AX, MIR548AO_5P, MIR548AX
MIR194_5P, MIR194_5P
MIR585_5P, MIR585_5P
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
MIR8065, MIR8065
MIR3682_5P, MIR3682_5P
MIR4727_5P, MIR4727_5P
MIR12132, MIR12132
MIR584_5P, MIR584_5P
GOMF_INTEGRIN_BINDING, GOMF_INTEGRIN_BINDING
MIR4678, MIR4678
MIR642B_5P, MIR642B_5P
MIR889_5P, MIR889_5P
GOBP_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS, GOBP_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS
HP_THORACIC_HYPOPLASIA, HP_THORACIC_HYPOPLASIA
REACTOME_RHOG_GTPASE_CYCLE, REACTOME_RHOG_GTPASE_CYCLE
MIR4286, MIR4286
GSE39820_IL1B_IL6_VS_IL1B_IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_IL1B_IL6_VS_IL1B_IL6_IL23A_TREATED_CD4_TCELL_UP
MIR891A_3P, MIR891A_3P
GOBP_AMINOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_AMINOGLYCAN_BIOSYNTHETIC_PROCESS
MIR597_5P, MIR597_5P
INGRAM_SHH_TARGETS_UP, INGRAM_SHH_TARGETS_UP
MIR211_3P, MIR211_3P
MIR7549_3P, MIR7549_3P
MIR548Q, MIR548Q
MIR4797_3P, MIR4797_3P
MIR592, MIR592
REACTOME_CA2_PATHWAY, REACTOME_CA2_PATHWAY
MIR6866_5P, MIR6866_5P
TCTATGA_MIR376A, MIR376B, TCTATGA_MIR376A, MIR376B
MIR3120_5P, MIR3120_5P
HARALAMBIEVA_PBMC_M_M_R_IL_AGE_13_16YO_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_POST_TOP_DEG_UP, HARALAMBIEVA_PBMC_M_M_R_IL_AGE_13_16YO_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_POST_TOP_DEG_UP
GSE34205_RSV_VS_FLU_INF_INFANT_PBMC_UP, GSE34205_RSV_VS_FLU_INF_INFANT_PBMC_UP
MIR3682_3P, MIR3682_3P
MIR331_5P, MIR331_5P
RELA_DN.V1_UP, RELA_DN.V1_UP
MIR7112_3P, MIR7112_3P
HP_LYMPHEDEMA, HP_LYMPHEDEMA
GSE11924_TH1_VS_TH17_CD4_TCELL_UP, GSE11924_TH1_VS_TH17_CD4_TCELL_UP
WP_MECHANOREGULATION_AND_PATHOLOGY_OF_YAPTAZ_VIA_HIPPO_AND_NONHIPPO_MECHANISMS, WP_MECHANOREGULATION_AND_PATHOLOGY_OF_YAPTAZ_VIA_HIPPO_AND_NONHIPPO_MECHANISMS
MIR3142, MIR3142
SCHOEN_NFKB_SIGNALING, SCHOEN_NFKB_SIGNALING
GOBP_POLYSACCHARIDE_METABOLIC_PROCESS, GOBP_POLYSACCHARIDE_METABOLIC_PROCESS
GRAHAM_NORMAL_QUIESCENT_VS_NORMAL_DIVIDING_UP, GRAHAM_NORMAL_QUIESCENT_VS_NORMAL_DIVIDING_UP
KEGG_AXON_GUIDANCE, KEGG_AXON_GUIDANCE
TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_6HR_UP, TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_6HR_UP
MIR601, MIR601
MIR1224_5P, MIR1224_5P
GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY, GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY
RAGHAVACHARI_PLATELET_SPECIFIC_GENES, RAGHAVACHARI_PLATELET_SPECIFIC_GENES
HP_ABNORMALITY_OF_SKULL_OSSIFICATION, HP_ABNORMALITY_OF_SKULL_OSSIFICATION
GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS, GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS
BCAT_GD5748_DN, BCAT_GD5748_DN
GOBP_WNT_SIGNALING_PATHWAY_CALCIIUM_MODULATING_PATHWAY, GOBP_WNT_SIGNALING_PATHWAY_CALCIIUM_MODULATING_PATHWAY
WP_G_PROTEIN_SIGNALING_PATHWAYS, WP_G_PROTEIN_SIGNALING_PATHWAYS
MIR4539, MIR4539
HP_ABNORMAL_SPERM_MOTILITY, HP_ABNORMAL_SPERM_MOTILITY
HP_SHORT_RIBS, HP_SHORT_RIBS
GOBP_THYMUS_DEVELOPMENT, GOBP_THYMUS_DEVELOPMENT
REACTOME_PYROPTOSIS, REACTOME_PYROPTOSIS
HP_ENLARGED_THORAX, HP_ENLARGED_THORAX
GOMF_EXTRACELLULAR_MATRIX_BINDING, GOMF_EXTRACELLULAR_MATRIX_BINDING
GOBP_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
GOMF_COLLAGEN_BINDING, GOMF_COLLAGEN_BINDING
GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS, GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS
HP_BROAD_EYEBROW, HP_BROAD_EYEBROW
MIR769_5P, MIR769_5P
GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_DN, GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_DN
GOMF_UDP_GLYCOSYLTRANSFERASE_ACTIVITY, GOMF_UDP_GLYCOSYLTRANSFERASE_ACTIVITY
MIR145_3P, MIR145_3P
HP_ABNORMAL_DELIVERY, HP_ABNORMAL_DELIVERY
GOBP_REGULATION_OF_GOLGI_ORGANIZATION, GOBP_REGULATION_OF_GOLGI_ORGANIZATION
GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMC_UP, GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMC_UP
GOMF_PHOSPHATIDYLSELINE_BINDING, GOMF_PHOSPHATIDYLSELINE_BINDING
HP_MULTIPLE_PRENATAL_FRACTURES, HP_MULTIPLE_PRENATAL_FRACTURES
GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
HP_PLEURAL EFFUSION, HP_PLEURAL EFFUSION
HP_ATYPICAL_ABSENCE_SEIZURE, HP_ATYPICAL_ABSENCE_SEIZURE
GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
HP_DERMAL_TRANSLUCENCY, HP_DERMAL_TRANSLUCENCY
GOBP_ENERGY_RESERVE_METABOLIC_PROCESS, GOBP_ENERGY_RESERVE_METABOLIC_PROCESS
GOBP_REGULATION_OF_AMINO_ACID_TRANSPORT, GOBP_REGULATION_OF_AMINO_ACID_TRANSPORT
GOMF_ACETYLGALACTOSAMINYLTRANSFERASE_ACTIVITY, GOMF_ACETYLGALACTOSAMINYLTRANSFERASE_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_SYNAPTIC_TRANSMISSION, GOBP_POSITIVE_REGULATION_OF_SYNAPTIC_TRANSMISSION
GOBP_MESODERM_DEVELOPMENT, GOBP_MESODERM_DEVELOPMENT
HP_REDUCED_SPERM_MOTILITY, HP_REDUCED_SPERM_MOTILITY
GOBP_FIBRINOLYSIS, GOBP_FIBRINOLYSIS
PID_LYMPH_ANGIOGENESIS_PATHWAY, PID_LYMPH_ANGIOGENESIS_PATHWAY
GOBP_LABYRINTHINE_LAYER_MORPHOGENESIS, GOBP_LABYRINTHINE_LAYER_MORPHOGENESIS
BIOCARTA_UCALPAIN_PATHWAY, BIOCARTA_UCALPAIN_PATHWAY
LIEN_BREAST_CARCINOMA_METAPLASTIC, LIEN_BREAST_CARCINOMA_METAPLASTIC
WP_HEDGEHOG_SIGNALING_PATHWAY, WP_HEDGEHOG_SIGNALING_PATHWAY
GOCC_INSULIN_RESPONSIVE_COMPARTMENT, GOCC_INSULIN_RESPONSIVE_COMPARTMENT
REACTOME_MOLECULES_ASSOCIATED_WITH_ELASTIC_FIBRES, REACTOME_MOLECULES_ASSOCIATED_WITH_ELASTIC_FIBRES
GOBP_CHROMATIN_SILENCING_AT_TELOMERE, GOBP_CHROMATIN_SILENCING_AT_TELOMERE
FERRARI_RESPONSE_TO_FENRETINIDE_UP, FERRARI_RESPONSE_TO_FENRETINIDE_UP
HASLINGER_B_CLL_WITH_11Q23_DELETION, HASLINGER_B_CLL_WITH_11Q23_DELETION
REACTOME_SYNDECAN_INTERACTIONS, REACTOME_SYNDECAN_INTERACTIONS
MIR3677_5P, MIR3677_5P
GOMF_GLUTAMATE_RECEPTOR_BINDING, GOMF_GLUTAMATE_RECEPTOR_BINDING
TACCGGT_MIR99A, MIR100, MIR99B, TACCGGT_MIR99A, MIR100, MIR99B
REACTOME_SYNTHESIS_OF_PC, REACTOME_SYNTHESIS_OF_PC
GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION
HP_ABNORMALITY_OF_THE_UMBILICAL_CORD, HP_ABNORMALITY_OF_THE_UMBILICAL_CORD
GOBP_KIDNEY_MORPHOGENESIS, GOBP_KIDNEY_MORPHOGENESIS
GOBP_REGULATION_OF_PHOSPHATIDYLCHOLINE_METABOLIC_PROCESS, GOBP_REGULATION_OF_PHOSPHATIDYLCHOLINE_METABOLIC_PROCESS
GOBP_REGULATION_OF_EPIDERMIS_DEVELOPMENT, GOBP_REGULATION_OF_EPIDERMIS_DEVELOPMENT
REACTOME_ELASTIC_FIBRE_FORMATION, REACTOME_ELASTIC_FIBRE_FORMATION
GOBP_POSITIVE_REGULATION_OF_ORGANIC_ACID_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_ORGANIC_ACID_TRANSPORT
GOBP_NEGATIVE_REGULATION_OF_HORMONE_SECRETION, GOBP_NEGATIVE_REGULATION_OF_HORMONE_SECRETION
MIR9900, MIR9900
GOMF_MANNOSIDASE_ACTIVITY, GOMF_MANNOSIDASE_ACTIVITY
BIOCARTA_EDG1_PATHWAY, BIOCARTA_EDG1_PATHWAY
HP_DIAPHYSEAL_THICKENING, HP_DIAPHYSEAL_THICKENING
HP_ABNORMALITY_OF_THE_5TH_METACARPAL, HP_ABNORMALITY_OF_THE_5TH_METACARPAL
GOMF_MAP_KINASE_ACTIVITY, GOMF_MAP_KINASE_ACTIVITY
REACTOME_PROSTACYCLIN_SIGNALLING_THROUGH_PROSTACYCLIN_RECEPTOR, REACTOME_PROSTACYCLIN_SIGNALLING_THROUGH_PROSTACYCLIN_RECEPTOR
HP_ABNORMAL_PELVIS_BONE_OSSIFICATION, HP_ABNORMAL_PELVIS_BONE_OSSIFICATION
DORSEY_GAB2_TARGETS, DORSEY_GAB2_TARGETS
GOBP_STRIATED_MUSCLE_CELL_PROLIFERATION, GOBP_STRIATED_MUSCLE_CELL_PROLIFERATION
HINATA_NFKB_TARGETS, KERATINOCYTE_DN, HINATA_NFKB_TARGETS, KERATINOCYTE_DN
GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS
HP_HYPOCALCEMIC_TETANY, HP_HYPOCALCEMIC_TETANY
GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION
GOBP_ENDOCRINE_PROCESS, GOBP_ENDOCRINE_PROCESS
GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS, GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS
HP_GASTRITIS, HP_GASTRITIS
GOBP_NAD_BIOSYNTHETIC_PROCESS, GOBP_NAD_BIOSYNTHETIC_PROCESS
WP_SEROTONIN_TRANSPORTER_ACTIVITY, WP_SEROTONIN_TRANSPORTER_ACTIVITY
GOBP_RAB_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_RAB_PROTEIN_SIGNAL_TRANSDUCTION
GOBP_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEMBLY, GOBP_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEMBLY
GOMF_FIBRONECTIN_BINDING, GOMF_FIBRONECTIN_BINDING
GOBP_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
GOBP_MESONEPHRIC_TUBULE_MORPHOGENESIS, GOBP_MESONEPHRIC_TUBULE_MORPHOGENESIS
GOBP_POSITIVE_REGULATION_OF_AMINO_ACID_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_AMINO_ACID_TRANSPORT
WP_ANGIOGENESIS, WP_ANGIOGENESIS
BIOCARTA_IL5_PATHWAY, BIOCARTA_IL5_PATHWAY
HP_IMPACTED_TOOTH, HP_IMPACTED_TOOTH
GOBP_REGULATION_OF_DNA_LIGATION, GOBP_REGULATION_OF_DNA_LIGATION
LINDSTEDT_DENDRITIC_CELL_MATURATION_A, LINDSTEDT_DENDRITIC_CELL_MATURATION_A
REACTOME_CHONDROITIN_SULFATE_BIOSYNTHESIS, REACTOME_CHONDROITIN_SULFATE_BIOSYNTHESIS
REACTOME_GLUACAGON LIKE PEPTIDE_1_GLP1_REGULATES_INSULIN_SECRETION, REACTOME_GLUACAGON LIKE PEPTIDE_1_GLP1_REGULATES_INSULIN_SECRETION
GOBP_REGULATION_OF_PLASMINOGEN_ACTIVATION, GOBP_REGULATION_OF_PLASMINOGEN_ACTIVATION
HASLINGER_B_CLL_WITH_MUTATED_VH_GENES, HASLINGER_B_CLL_WITH_MUTATED_VH_GENES
KIM_ALL_DISORDERS_DURATION_CORR_UP, KIM_ALL_DISORDERS_DURATION_CORR_UP
WP_LTF_DANGER_SIGNAL_RESPONSE_PATHWAY, WP_LTF_DANGER_SIGNAL_RESPONSE_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_FIBRINOLYSIS, GOBP_NEGATIVE_REGULATION_OF_FIBRINOLYSIS
WP_TRYPTOPHAN_CATABOLISM_LEADING_TO_NAD_PRODUCTION, WP_TRYPTOPHAN_CATABOLISM_LEADING_TO_NAD_PRODUCTION
WP_NANOPARTICLEMEDIATED_ACTIVATION_OF_RECEPTOR_SIGNALING, WP_NANOPARTICLEMEDIATED_ACTIVATION_OF_RECEPTOR_SIGNALING
GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION
GOBP_PHOSPHATIDYLSELINE_METABOLIC_PROCESS, GOBP_PHOSPHATIDYLSELINE_METABOLIC_PROCESS
GOBP_NEPHRON_MORPHOGENESIS, GOBP_NEPHRON_MORPHOGENESIS
GOBP_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION
HP_APLASIA_HYPOPLASIA_OF_THE_5TH_METACARPAL, HP_APLASIA_HYPOPLASIA_OF_THE_5TH_METACARPAL
GOBP_NEGATIVE_REGULATION_OF_ENDOTHELIAL_CELL_CHEMOTAXIS, GOBP_NEGATIVE_REGULATION_OF_ENDOTHELIAL_CELL_CHEMOTAXIS
HP_ABSENT_OR_MINIMALLY_OSSIFIED_VERTEBRAL_BODIES, HP_ABSENT_OR_MINIMALLY_OSSIFIED_VERTEBRAL_BODIES
GOBP_ENDOCRINE_HORMONE_SECRETION, GOBP_ENDOCRINE_HORMONE_SECRETION
GOMF_C_X_C_CHEMOKINE_BINDING, GOMF_C_X_C_CHEMOKINE_BINDING
HP_HIGH_FREQUENCY_SENSORINEURAL_HEARING_IMPAIRMENT, HP_HIGH_FREQUENCY_SENSORINEURAL_HEARING_IMPAIRMENT
GOBP_POSITIVE_REGULATION_OF_DNA_LIGATION, GOBP_POSITIVE_REGULATION_OF_DNA_LIGATION
GOMF_C_X3_C_CHEMOKINE_BINDING, GOMF_C_X3_C_CHEMOKINE_BINDING
GOBP_CALMODULIN_DEPENDENT_KINASE_SIGNALING_PATHWAY, GOBP_CALMODULIN_DEPENDENT_KINASE_SIGNALING_PATHWAY
GOBP_KETONE_CATABOLIC_PROCESS, GOBP_KETONE_CATABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS, GOBP_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS
BIOCARTA_TFF_PATHWAY, BIOCARTA_TFF_PATHWAY
chr4p15, chr4p15
HP_REFLEX_SEIZURE, HP_REFLEX_SEIZURE
REACTOME_G_ALPHA_S_SIGNALLING_EVENTS, REACTOME_G_ALPHA_S_SIGNALLING_EVENTS
GOME_NEUREGULIN_BINDING, GOMF_NEUREGULIN_BINDING
PALOMERO_GSL_SENSITIVITY_DN, PALOMERO_GSL_SENSITIVITY_DN
HP_ANKYLOSIS, HP_ANKYLOSIS
chr15q23, chr15q23
HP_APLASIA_HYPOPLASIA_OF_THE_COLON, HP_APLASIA_HYPOPLASIA_OF_THE_COLON
HP_PSEUDOARTHROSIS, HP_PSEUDOARTHROSIS
GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION
HP_EEG_WITH_FOCAL_SHARP_SLOW WAVES, HP_EEG_WITH_FOCAL_SHARP_SLOW WAVES
GOBP_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
HP_POSTEXERTIONAL_MALAISE, HP_POSTEXERTIONAL_MALAISE
GOMF_OPIOID_RECEPTOR_BINDING, GOMF_OPIOID_RECEPTOR_BINDING
GOBP_NEGATIVE_REGULATION_OF_CGMP_MEDIATED_SIGNALING, GOBP_NEGATIVE_REGULATION_OF_CGMP_MEDIATED_SIGNALING
GOBP_NEGATIVE_REGULATION_OF_VASOCONSTRICTION, GOBP_NEGATIVE_REGULATION_OF_VASOCONSTRICTION
GOBP_REGULATION_OF_INWARD_RECTIFIER_POTASSIUM_CHANNEL_ACTIVITY, GOBP_REGULATION_OF_INWARD_RECTIFIER_POTASSIUM_CHANNEL_ACTIVITY
GOMF_CELL_MATRIX_ADHESION_MEDIATOR_ACTIVITY, GOMF_CELL_MATRIX_ADHESION_MEDIATOR_ACTIVITY
DESCARTES_MAIN_FETAL_INHIBITORY_INTERNEURONS, DESCARTES_MAIN_FETAL_INHIBITORY_INTERNEURONS
REACTOME_GLUACAGON_TYPE_LIGAND_RECEPTORS, REACTOME_GLUACAGON_TYPE_LIGAND_RECEPTORS
GOBP_PLASMINOGEN_ACTIVATION, GOBP_PLASMINOGEN_ACTIVATION
GOBP_ENDOTHELIAL_CELL_CHEMOTAXIS, GOBP_ENDOTHELIAL_CELL_CHEMOTAXIS
REACTOME_ADORA2B_MEDIATED_ANTINFLAMMATORY_CYTOKINES_PRODUCTION, REACTOME_ADORA2B_MEDIATED_ANTINFLAMMATORY_CYTOKINES_PRODUCTION