

GSE22527_ANTL_CD3_INVIVO_VS_UNTREATED_MOUSE_TREG_DN, GSE22527_ANTL_CD3_INVIVO_VS_UNTREATED_MOUSE_TREG_DN
GSE22033_UNTREATED_VS_ROSIGLITAZONE_TREATED_MEF_UP, GSE22033_UNTREATED_VS_ROSIGLITAZONE_TREATED_MEF_UP
GSE21033_CTRL_VS_POLYIC_STIM_DC_3H_UP, GSE21033_CTRL_VS_POLYIC_STIM_DC_3H_UP
GSE20500_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_DN, GSE20500_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_DN
GSE13484_UNSTIM_VS_3H_YF17D_VACCINE_STIM_PPMC_UP, GSE13484_UNSTIM_VS_3H_YF17D_VACCINE_STIM_PPMC_UP
GSE20152_SPHK1_KO_VS_WT_HTNFA_OVERXPRESS_ANKLE_DN, GSE20152_SPHK1_KO_VS_WT_HTNFA_OVERXPRESS_ANKLE_DN
GSE37301_LYMPHOID_PRIMED_MPP_VS_GRAN_MONO_PROGENITOR_DN, GSE37301_LYMPHOID_PRIMED_MPP_VS_GRAN_MONO_PROGENITOR_DN
GSE20152_SPHK1_KO_VS_HTNFA_OVEREXPRESS_ANKLE_DN, GSE20152_SPHK1_KO_VS_HTNFA_OVEREXPRESS_ANKLE_DN
GSE22033_UNTREATED_VS_MRL24_TREATED_MEF_UP, GSE22033_UNTREATED_VS_MRL24_TREATED_MEF_UP
GSE19401_RETINOIC_ACID_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN, GSE19401_RETINOIC_ACID_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN
GSE3982_MEMORY_CD4_TCELL_VS_TH1_UP, GSE3982_MEMORY_CD4_TCELL_VS_TH1_UP
GSE1925_3H_VS_24H_IFNG_STIM_MACROPHAGE_DN, GSE1925_3H_VS_24H_IFNG_STIM_MACROPHAGE_DN
MIR525_5P, MIR525_5P
GSE42021_CD24HI_VS_CD24INT_TCONV_THYMUS_UP, GSE42021_CD24HI_VS_CD24INT_TCONV_THYMUS_UP
MIR520A_5P, MIR520A_5P
GSE22886_NAIVE_CD8_TCELL_VS_MONOCYTE_UP, GSE22886_NAIVE_CD8_TCELL_VS_MONOCYTE_UP
MIR4257, MIR4257
GSE43955_TH0_VS_TGFB_IL6_THI7_ACT_CD4_TCELL_1H_UP, GSE43955_TH0_VS_TGFB_IL6_THI7_ACT_CD4_TCELL_1H_UP
GSE43955_10H_VS_30H_ACT_CD4_TCELL_WITH_TGFB_IL6_UP, GSE43955_10H_VS_30H_ACT_CD4_TCELL_WITH_TGFB_IL6_UP
CAGCCTC_MIR4855P, CAGCCTC_MIR4855P
GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_DN, GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_DN
MIR6131, MIR6131
AGGTGCA_MIR500, AGGTGCA_MIR500
GSE3982_BASOPHIL_VS_NKCELL_DN, GSE3982_BASOPHIL_VS_NKCELL_DN
ESC_V6.5_UP_EARLY.V1_UP, ESC_V6.5_UP_EARLY.V1_UP
ZHAN_MULTIPLE_MYELOMA_CD1_AND_CD2_DN, ZHAN_MULTIPLE_MYELOMA_CD1_AND_CD2_DN
GOBP_REGULATION_OF_PROTEIN_TARGETING, GOBP_REGULATION_OF_PROTEIN_TARGETING
REACTOME_RHOB_GTPASE_CYCLE, REACTOME_RHOB_GTPASE_CYCLE
WP_DNA_DAMAGE_RESPONSE_ONLY_ATM_DEPENDENT, WP_DNA_DAMAGE_RESPONSE_ONLY_ATM_DEPENDENT
REACTOME_CELL_DEATH_SIGNALLING_VIA_NRAGE_NRIF_AND_NADE, REACTOME_CELL_DEATH_SIGNALLING_VIA_NRAGE_NRIF_AND_NADE
COBLL1_TARGET_GENES, COBLL1_TARGET_GENES
MIR10401_5P, MIR10401_5P
REACTOME_METALLOPROTEASE_DUBS, REACTOME_METALLOPROTEASE_DUBS
GSE24972_WT_VS_IRF8_KO_SPLEEN_FOLLICULAR_BCELL_UP, GSE24972_WT_VS_IRF8_KO_SPLEEN_FOLLICULAR_BCELL_UP
GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY, GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY
GSE3982_MAC_VS_EFF_MEMORY_CD4_TCELL_DN, GSE3982_MAC_VS_EFF_MEMORY_CD4_TCELL_DN
MIR1286, MIR1286
GSE13306_TREG_RA_VS_TCONV_RA_DN, GSE13306_TREG_RA_VS_TCONV_RA_DN
PDGF_ERK_DN.V1_UP, PDGF_ERK_DN.V1_UP
MIR1233_5P, MIR1233_5P
PRC2_EED_UP.V1_UP, PRC2_EED_UP.V1_UP
GOMF_HISTONE_LYSINE_N_METHYLTRANSFERASE_ACTIVITY, GOMF_HISTONE_LYSINE_N_METHYLTRANSFERASE_ACTIVITY
WP_AMPACTIVATED_PROTEIN_KINASE_AMPK_SIGNALING, WP_AMPACTIVATED_PROTEIN_KINASE_AMPK_SIGNALING
RORA2_01, RORA2_01
GSE15659_RESTING_VS_ACTIVATED_TREG_UP, GSE15659_RESTING_VS_ACTIVATED_TREG_UP
GOCC_PERICENTRIC_HETEROCHROMATIN, GOCC_PERICENTRIC_HETEROCHROMATIN
GOMF_HSP90_PROTEIN_BINDING, GOMF_HSP90_PROTEIN_BINDING
MIR3907, MIR3907
WP_WNT_SIGNALING_PATHWAY_AND_PLURIPOTENCY, WP_WNT_SIGNALING_PATHWAY_AND_PLURIPOTENCY
GOMF_TAU_PROTEIN_BINDING, GOMF_TAU_PROTEIN_BINDING
MIR6749_3P, MIR6749_3P
GSE15659_CD45RA_NEG_CD4_TCELL_VS_ACTIVATED_TREG_UP, GSE15659_CD45RA_NEG_CD4_TCELL_VS_ACTIVATED_TREG_UP
BRACHAT_RESPONSE_TO_METHOTREXATE_UP, BRACHAT_RESPONSE_TO_METHOTREXATE_UP
GOBP_REGULATION_OF_INSULIN_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_INSULIN_RECEPTOR_SIGNALING_PATHWAY
LANDIS_ERBB2_BREAST_TUMORS_65_UP, LANDIS_ERBB2_BREAST_TUMORS_65_UP
MIR219A_2_3P, MIR219A_2_3P
REACTOME_REGULATION_OF_SIGNALING_BY_CBL, REACTOME_REGULATION_OF_SIGNALING_BY_CBL
GOBP_CARDIAC_SEPTUM_DEVELOPMENT, GOBP_CARDIAC_SEPTUM_DEVELOPMENT
GOCC_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX, GOCC_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX
GOBP_MATERNAL_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY, GOBP_MATERNAL_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY
GOBP_POSITIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MEMBRANE, GOBP_POSITIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MEMBRANE
GOMF_ESTROGEN_RECEPTOR_BINDING, GOMF_ESTROGEN_RECEPTOR_BINDING
REACTOME_CHAPERONE_MEDIATED_AUTOPHAGY, REACTOME_CHAPERONE_MEDIATED_AUTOPHAGY
GOBP_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS, GOBP_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS
GOMF_PHOSPHATIDYLINOSITOL_KINASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_KINASE_ACTIVITY
GOBP_REGULATION_OF_MITOCHONDRIAL_FISSION, GOBP_REGULATION_OF_MITOCHONDRIAL_FISSION
SHIN_B_CELL_LYMPHOMA_CLUSTER_3, SHIN_B_CELL_LYMPHOMA_CLUSTER_3
WP_SYNAPTIC_SIGNALING_PATHWAYS_ASSOCIATED_WITH_AUTISM_SPECTRUM_DISORDER, WP_SYNAPTIC_SIGNALING_PATHWAYS_ASSOCIATED_WITH_AUTISM_SPECTRUM_DISORDER
MODULE_327, MODULE_327
GOBP_PHOSPHATIDYLINOSITOL_3_PHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_PHOSPHATIDYLINOSITOL_3_PHOSPHATE_BIOSYNTHETIC_PROCESS
GOBP_CORONARY_VASCULATURE_DEVELOPMENT, GOBP_CORONARY_VASCULATURE_DEVELOPMENT
GOBP_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS
HP_URETEROPELVIC_JUNCTION_OBSTRUCTION, HP_URETEROPELVIC_JUNCTION_OBSTRUCTION
BRACHAT_RESPONSE_TO_CISPLATIN, BRACHAT_RESPONSE_TO_CISPLATIN
GOBP_PEPTIDYL_LYSINE_DIMETHYLATION, GOBP_PEPTIDYL_LYSINE_DIMETHYLATION
HP_GENERALIZED_HYPOPIGMENTATION, HP_GENERALIZED_HYPOPIGMENTATION
ZIRN_TRETINOIN_RESPONSE_WT1_UP, ZIRN_TRETINOIN_RESPONSE_WT1_UP
GOMF_PDZ_DOMAIN_BINDING, GOMF_PDZ_DOMAIN_BINDING
CCCACAT_MIR2993P, CCCACAT_MIR2993P
GOMF_RETINOIC_ACID_RECEPTOR_BINDING, GOMF_RETINOIC_ACID_RECEPTOR_BINDING
MCCLUNG_DELTA_FOSB_TARGETS_8WK, MCCLUNG_DELTA_FOSB_TARGETS_8WK
GOBP_REGULATION_OF_PROTEIN_TARGETING_TO_MEMBRANE, GOBP_REGULATION_OF_PROTEIN_TARGETING_TO_MEMBRANE
HP_ABNORMAL_CIRCULATING_NON_PROTEINOGENIC_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_NON_PROTEINOGENIC_AMINO_ACID_CONCENTRATION
REACTOME_HS_GAG_BIOSYNTHESIS, REACTOME_HS_GAG_BIOSYNTHESIS
REACTOME_DEACTIVATION_OF_THE_BETA_CATENIN_TRANSACTIVATING_COMPLEX, REACTOME_DEACTIVATION_OF_THE_BETA_CATENIN_TRANSACTIVATING_COMPLEX
HP_ELEVATED_MEAN_ARTERIAL_PRESSURE, HP_ELEVATED_MEAN_ARTERIAL_PRESSURE
REACTOME_NEGATIVE_REGULATION_OF_MET_ACTIVITY, REACTOME_NEGATIVE_REGULATION_OF_MET_ACTIVITY
HP_HYPOSMIA, HP_HYPOSMIA
GOBP_CELL_VOLUME_HOMEOSTASIS, GOBP_CELL_VOLUME_HOMEOSTASIS
GOBP_MESODERMAL_CELL_DIFFERENTIATION, GOBP_MESODERMAL_CELL_DIFFERENTIATION
CHEOK_RESPONSE_TO_MERCAPTOPURINE_UP, CHEOK_RESPONSE_TO_MERCAPTOPURINE_UP
HP_MULTIPLE_GASTRIC_POLYPS, HP_MULTIPLE_GASTRIC_POLYPS
GOCC_MLL3_4_COMPLEX, GOCC_MLL3_4_COMPLEX
GOBP_REGULATION_OF_MUSCLE_ORGAN_DEVELOPMENT, GOBP_REGULATION_OF_MUSCLE_ORGAN_DEVELOPMENT
TOMLINS_METASTASIS_UP, TOMLINS_METASTASIS_UP
MODULE_156, MODULE_156
GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_UP, GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_UP
GOBP_CAMP_METABOLIC_PROCESS, GOBP_CAMP_METABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY
MODULE_547, MODULE_547
GOMF_LAMIN_BINDING, GOMF_LAMIN_BINDING
GEISS_RESPONSE_TO_DSRNA_DN, GEISS_RESPONSE_TO_DSRNA_DN
GOMF_ANION_CATION_SYMPORTER_ACTIVITY, GOMF_ANION_CATION_SYMPORTER_ACTIVITY
HP_ABNORMALITY_OF_SMALL_INTESTINAL_VILLUS_MORPHOLOGY, HP_ABNORMALITY_OF_SMALL_INTESTINAL_VILLUS_MORPHOLOGY
chr16q12, chr16q12
GOBP_POSITIVE_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING, GOBP_POSITIVE_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING
LOPEZ_MESOTHELIOMA_SURVIVAL_WORST_VS_BEST_DN, LOPEZ_MESOTHELIOMA_SURVIVAL_WORST_VS_BEST_DN
GOBP_SMAD_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_SMAD_PROTEIN_SIGNAL_TRANSDUCTION
GOCC_CD40_RECEPTOR_COMPLEX, GOCC_CD40_RECEPTOR_COMPLEX
KEGG_MELANOGENESIS, KEGG_MELANOGENESIS
GOBP_AORTA_DEVELOPMENT, GOBP_AORTA_DEVELOPMENT
MODULE_240, MODULE_240
REACTOME_INTERLEUKIN_RECEPTOR_SHC_SIGNALING, REACTOME_INTERLEUKIN_RECEPTOR_SHC_SIGNALING
CGCTGCT_MIR503, CGCTGCT_MIR503
MODULE_313, MODULE_313
GOBP_AMMONIUM_TRANSPORT, GOBP_AMMONIUM_TRANSPORT
GOMF_AMMONIUM_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_AMMONIUM_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
BIOCARTA_GABA_PATHWAY, BIOCARTA_GABA_PATHWAY
WP_MAJOR_RECEPTORS_TARGETED_BY_EPINEPHRINE_AND_NOREPINEPHRINE, WP_MAJOR_RECEPTORS_TARGETED_BY_EPINEPHRINE_AND_NOREPINEPHRINE
REACTOME_LISTERIA_MONOCYTOGENES_ENTRY_INTO_HOST_CELLS, REACTOME_LISTERIA_MONOCYTOGENES_ENTRY_INTO_HOST_CELLS
GOBP_CELLULAR_RESPONSE_TO_MURAMYL_DIPEPTIDE, GOBP_CELLULAR_RESPONSE_TO_MURAMYL_DIPEPTIDE
JIANG_CORE_DUPLICATION_GENES, JIANG_CORE_DUPLICATION_GENES
HP_GONADAL_DYSGENESIS, HP_GONADAL_DYSGENESIS
GOBP_TACHYKININ_RECEPTOR_SIGNALING_PATHWAY, GOBP_TACHYKININ_RECEPTOR_SIGNALING_PATHWAY
GOBP_RESPONSE_TO_HISTAMINE, GOBP_RESPONSE_TO_HISTAMINE
GOBP_GLIAL_CELL_FATE_COMMITMENT, GOBP_GLIAL_CELL_FATE_COMMITMENT
GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H3_K4_SPECIFIC, GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H3_K4_SPECIFIC
MIR616_3P, MIR616_3P
GOBP_MESODERM_MORPHOGENESIS, GOBP_MESODERM_MORPHOGENESIS
AGARWAL_AKT_PATHWAY_TARGETS, AGARWAL_AKT_PATHWAY_TARGETS
GOBP_OXIDATIVE_DNA_DEMETHYLATION, GOBP_OXIDATIVE_DNA_DEMETHYLATION
REACTOME_GABA_B_RECEPTOR_ACTIVATION, REACTOME_GABA_B_RECEPTOR_ACTIVATION
REACTOME_SIGNALING_BY_NOTCH2, REACTOME_SIGNALING_BY_NOTCH2
GOBP_HEPARAN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_HEPARAN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS
GOBP_MITRAL_VALVE_DEVELOPMENT, GOBP_MITRAL_VALVE_DEVELOPMENT
BIOCARTA_IL4_PATHWAY, BIOCARTA_IL4_PATHWAY
REACTOME_EGFR_DOWNREGULATION, REACTOME_EGFR_DOWNREGULATION
HP_PROTEIN_LOSING_ENTEROPATHY, HP_PROTEIN_LOSING_ENTEROPATHY
MIKKELSEN_IPS_1CP_WITH_H3K27ME3, MIKKELSEN_IPS_1CP_WITH_H3K27ME3
GOBP_HYPOTHALAMUS_CELL_DIFFERENTIATION, GOBP_HYPOTHALAMUS_CELL_DIFFERENTIATION
GOBP_CELLULAR_RESPONSE_TO_HISTAMINE, GOBP_CELLULAR_RESPONSE_TO_HISTAMINE
REACTOME_CATION_COUPLED_CHLORIDE_COTRANSPORTERS, REACTOME_CATION_COUPLED_CHLORIDE_COTRANSPORTERS
HP_CHRONIC_AXONAL_NEUROPATHY, HP_CHRONIC_AXONAL_NEUROPATHY
GOBP_HEPARIN_METABOLIC_PROCESS, GOBP_HEPARIN_METABOLIC_PROCESS
GOBP_VESICLE_TETHERING_TO_GOLGI, GOBP_VESICLE_TETHERING_TO_GOLGI
WP_TYPE_I_COLLAGEN_SYNTHESIS_IN_THE_CONTEXT_OF_OSTEOGENESIS_IMPERFECTA, WP_TYPE_I_COLLAGEN_SYNTHESIS_IN_THE_CONTEXT_OF_OSTEOGENESIS_IMPERFECTA
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_SALMON_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_SALMON_DN
PID_LPA4_PATHWAY, PID_LPA4_PATHWAY
GOBP_DORSAL_AORTA_DEVELOPMENT, GOBP_DORSAL_AORTA_DEVELOPMENT
SHLSPARC_TARGETS_DN, SHLSPARC_TARGETS_DN
ZNF112_TARGET_GENES, ZNF112_TARGET_GENES
GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION
GOBP_AUTOCRINE_SIGNALING, GOBP_AUTOCRINE_SIGNALING
GOBP_DORSAL_AORTA_MORPHOGENESIS, GOBP_DORSAL_AORTA_MORPHOGENESIS
GOBP_VENTRICULAR_COMPACT_MYOCARDIUM_MORPHOGENESIS, GOBP_VENTRICULAR_COMPACT_MYOCARDIUM_MORPHOGENESIS
HP_SUBMUCOUS_CLEFT_SOFT_PALATE, HP_SUBMUCOUS_CLEFT_SOFT_PALATE
HP_FIBROSARCOMA, HP_FIBROSARCOMA
GOCC_HFE_TRANSFERRIN_RECEPTOR_COMPLEX, GOCC_HFE_TRANSFERRIN_RECEPTOR_COMPLEX
chr4q12, chr4q12
GOBP_VENTRICULAR TRABECULA_MYOCARDIUM_MORPHOGENESIS, GOBP_VENTRICULAR TRABECULA_MYOCARDIUM_MORPHOGENESIS
REACTOME DISSOLUTION_OF_FIBRIN_CLOT, REACTOME DISSOLUTION_OF_FIBRIN_CLOT
GOBP_POSITIVE_REGULATION_OF_MUSCLE_TISSUE_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_MUSCLE_TISSUE_DEVELOPMENT
REACTOME_DISEASES_ASSOCIATED_WITH_SURFACTANT_METABOLISM, REACTOME_DISEASES_ASSOCIATED_WITH_SURFACTANT_METABOLISM
GOBP_REGULATION_OF_SMAD_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_SMAD_PROTEIN_SIGNAL_TRANSDUCTION
HP_MENORRHAGIA, HP_MENORRHAGIA
GOBP_ACTIVATION_OF_ADENYLATE_CYCLASE_ACTIVITY, GOBP_ACTIVATION_OF_ADENYLATE_CYCLASE_ACTIVITY
HP_CORONARY_ARTERY_STENOSIS, HP_CORONARY_ARTERY_STENOSIS
GOMF_INSULIN_BINDING, GOMF_INSULIN_BINDING
GOMF_CYTOSKELETON_NUCLEAR_MEMBRANE_ANCHOR_ACTIVITY, GOMF_CYTOSKELETON_NUCLEAR_MEMBRANE_ANCHOR_ACTIVITY
HP_ABNORMALITY_OF_THE_MASTOID, HP_ABNORMALITY_OF_THE_MASTOID
REACTOME_TGF_BETA_RECEPTOR_SIGNALING_IN_EMT_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, REACTOME_TGF_BETA_RECEPTOR_SIGNALING_IN_EMT_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
BIOCARTA_PLATELETAPP_PATHWAY, BIOCARTA_PLATELETAPP_PATHWAY
GOMF_POTASSIUM_CHLORIDE_SYMPORTER_ACTIVITY, GOMF_POTASSIUM_CHLORIDE_SYMPORTER_ACTIVITY
REACTOME_MYOCLONIC_EPILEPSY_OF_LAFORE, REACTOME_MYOCLONIC_EPILEPSY_OF_LAFORE
GOMF_TRANSFERASE_ACTIVITY_TRANSFERRING_SULFUR_CONTAINING_GROUPS, GOMF_TRANSFERASE_ACTIVITY_TRANSFERRING_SULFUR_CONTAINING_GROUPS
GOMF_G_PROTEIN_COUPLED_RECEPTOR_KINASE_ACTIVITY, GOMF_G_PROTEIN_COUPLED_RECEPTOR_KINASE_ACTIVITY
HP_DECREASED_LEVEL_OF_GABA_IN_SERUM, HP_DECREASED_LEVEL_OF_GABA_IN_SERUM
WP_BONE_MORPHOGENIC_PROTEIN_BMP_SIGNALLING_AND_REGULATION, WP_BONE_MORPHOGENIC_PROTEIN_BMP_SIGNALLING_AND_REGULATION
HP_CRANIOFACIAL_OSTEOSCLEROSIS, HP_CRANIOFACIAL_OSTEOSCLEROSIS
GOBP_ANTEROGRADE_NEURONAL_DENSE_CORE_VESICLE_TRANSPORT, GOBP_ANTEROGRADE_NEURONAL_DENSE_CORE_VESICLE_TRANSPORT
BIOCARTA_WNT_LRP6_PATHWAY, BIOCARTA_WNT_LRP6_PATHWAY
HP_INCREASED_URINE_PROTEINOGENIC_AMINO_ACID_DERIVATIVE_LEVEL, HP_INCREASED_URINE_PROTEINOGENIC_AMINO_ACID_DERIVATIVE_LEVEL
GOBP_BMP_SIGNALING_PATHWAY_INVOLVED_IN_HEART_DEVELOPMENT, GOBP_BMP_SIGNALING_PATHWAY_INVOLVED_IN_HEART_DEVELOPMENT
GOCC_CORTICAL_ENDOPLASMIC_RETICULUM, GOCC_CORTICAL_ENDOPLASMIC_RETICULUM
MIR3678_5P, MIR3678_5P
DESCARTES_MAIN_FETAL_AFP_ALB_POSITIVE_CELLS, DESCARTES_MAIN_FETAL_AFP_ALB_POSITIVE_CELLS
YIH_RESPONSE_TO_ARSENITE_C5, YIH_RESPONSE_TO_ARSENITE_C5
REACTOME_INLA_MEDIATED_ENTRY_OF_LISTERIA_MONOCYTOGENES_INTO_HOST_CELLS, REACTOME_INLA_MEDIATED_ENTRY_OF_LISTERIA_MONOCYTOGENES_INTO_HOST_CELLS
GOMF_GUANOSINE_DIPHOSPHATASE_ACTIVITY, GOMF_GUANOSINE_DIPHOSPHATASE_ACTIVITY
GOBP_MESENCHYMAL STEM_CELL_DIFFERENTIATION, GOBP_MESENCHYMAL STEM_CELL_DIFFERENTIATION
WP_MOLYBDENUM_COFACTOR_MOCO_BIOSYNTHESIS, WP_MOLYBDENUM_COFACTOR_MOCO_BIOSYNTHESIS
GOBP_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_GROWTH, GOBP_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_GROWTH
HP_PRECOCIOUS_PUBERTY_IN_FEMALES, HP_PRECOCIOUS_PUBERTY_IN_FEMALES
GOBP_NEURAL_PLATE_PATTERN_SPECIFICATION, GOBP_NEURAL_PLATE_PATTERN_SPECIFICATION
GOMF_1_PHOSPHATIDYLINOSITOL_4_PHOSPHATE_3_KINASE_ACTIVITY, GOMF_1_PHOSPHATIDYLINOSITOL_4_PHOSPHATE_3_KINASE_ACTIVITY
KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS, HEPARAN_SULFATE, KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS, HEPARAN_SULFATE