

3_KO_4H_LPS_STIM_MACROPHAGE_UP, GSE33162_HDAC3_KO_VS_HDAC3_KO_4H_LPS_STIM_MACROPHAGE_UP

GSE14000_TRANSLATED_RNA_VS_MRNA_4H_LPS_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_4H_LPS_DC_UP
GSE37416_0H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_0H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_DN, GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_DN
GSE1448_CTRL_VS_ANTL_VBETA5_DP_THYMOCYTE_UP, GSE1448_CTRL_VS_ANTL_VBETA5_DP_THYMOCYTE_UP
GSE3204_LY6C_HIGH_VS_LOW_MONOCYTE_DN, GSE3204_LY6C_HIGH_VS_LOW_MONOCYTE_DN
GOBP_CYTOSOLIC_TRANSPORT, GOBP_CYTOSOLIC_TRANSPORT
GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP, GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP
GSE3609_WT_VS_NLRP10_KO_DC_LPS_STIM_DN, GSE3609_WT_VS_NLRP10_KO_DC_LPS_STIM_DN
GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_DN, GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_DN
GSE27786_LSK_VS_BCELL_DN, GSE27786_LSK_VS_BCELL_DN
GOBP_VACUOLE_ORGANIZATION, GOBP_VACUOLE_ORGANIZATION
GSE17721_POLYIC_VS_PAMCSK4_0.5H_BMDC_DN, GSE17721_POLYIC_VS_PAMCSK4_0.5H_BMDC_DN
GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
GSE17721_0.5H_VS_4H_GARDQUIMOD_BMDC_DN, GSE17721_0.5H_VS_4H_GARDQUIMOD_BMDC_DN
GSE3920_IFNA_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP, GSE3920_IFNA_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP
GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_4H_UP, GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_4H_UP
GSE36888_STATS_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_UP, GSE36888_STATS_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_UP
GSE15330_WT_VS_IKAROS_KO_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN, GSE15330_WT_VS_IKAROS_KO_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN
GSE17721_POLYIC_VS_GARDQUIMOD_2H_BMDC_UP, GSE17721_POLYIC_VS_GARDQUIMOD_2H_BMDC_UP
GSE41087_WT_VS_FOXP3_MUT_ANTL_CD3_CD28_STEM_CD4_TCELL_UP, GSE41087_WT_VS_FOXP3_MUT_ANTL_CD3_CD28_STEM_CD4_TCELL_UP
GSE39192_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP, GSE39192_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP
GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN, GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN
GOBP_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI, GOBP_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI
GSE17721_LPS_VS_CPG_8H_BMDC_UP, GSE17721_LPS_VS_CPG_8H_BMDC_UP
GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP
GSE17721_LPS_VS_GARDQUIMOD_0.5H_BMDC_DN, GSE17721_LPS_VS_GARDQUIMOD_0.5H_BMDC_DN
GSE3939_CD4POS_VS_CD4POS_DC_IN_CULTURE_UP, GSE3939_CD4POS_VS_CD4POS_DC_IN_CULTURE_UP
GSE17721_CTRL_VS_POLYIC_24H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_24H_BMDC_DN
GOBP_AUTOPHAGOSOME_ORGANIZATION, GOBP_AUTOPHAGOSOME_ORGANIZATION
AACTGAC_MIR223, AACTGAC_MIR223
GSE3982_FOSINOPHIL_VS_TH1_UP, GSE3982_FOSINOPHIL_VS_TH1_UP
MIR6730_3P, MIR6730_3P
GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_DN, GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_DN
YAGL_AML_SURVIVAL, YAGL_AML_SURVIVAL
GOBP_ENDOSOME_ORGANIZATION, GOBP_ENDOSOME_ORGANIZATION
GOCC_EARLY_ENDOSOME_MEMBRANE, GOCC_EARLY_ENDOSOME_MEMBRANE
MIR642A_3P, MIR642B_3P, MIR642A_3P, MIR642B_3P
KEGG_NEUROTROPHIN_SIGNALING_PATHWAY, KEGG_NEUROTROPHIN_SIGNALING_PATHWAY
HP_TRIANGULAR_FACE, HP_TRIANGULAR_FACE
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_DN, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_DN
MIR5008_3P, MIR5008_3P
GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION, GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION
CCM4_ERBB2IP, CCM4_ERBB2IP
MIR6824_3P, MIR6824_3P
MIR6764_3P, MIR6764_3P
MIR125B_2_3P, MIR125B_2_3P
HP_PULMONIC_STENOSIS, HP_PULMONIC_STENOSIS
MIR223_3P, MIR223_3P
MIR338_3P, MIR338_3P
GSE3203_UNTREATED_VS_IFNB_TREATED_LN_BCELL_UP, GSE3203_UNTREATED_VS_IFNB_TREATED_LN_BCELL_UP
MIR7152_3P, MIR7152_3P
MORF_CDC2L5, MORF_CDC2L5
MIR1_5P, MIR1_5P
MIR18A_5P, MIR18B_5P, MIR18A_5P, MIR18B_5P
HP_ABNORMAL_PULMONARY_VALVE_PHYSIOLOGY, HP_ABNORMAL_PULMONARY_VALVE_PHYSIOLOGY
GSE41176_WT_VS_TAK1_KO_UNSTIM_BCELL_UP, GSE41176_WT_VS_TAK1_KO_UNSTIM_BCELL_UP
GOBP_NEGATIVE_REGULATION_OF_BINDING, GOBP_NEGATIVE_REGULATION_OF_BINDING
GOMF_TRANSCRIPTION_COREGULATOR_BINDING, GOMF_TRANSCRIPTION_COREGULATOR_BINDING
HP_WIDE_NOSE, HP_WIDE_NOSE
MODULE_503, MODULE_503
MOR1_MATURE_B_LYMPHOCYTE_UP, MOR1_MATURE_B_LYMPHOCYTE_UP
MIR1219, MIR1219
GOCC_EXTRINSIC_COMPONENT_OF_PLASMA_MEMBRANE, GOCC_EXTRINSIC_COMPONENT_OF_PLASMA_MEMBRANE
MODULE_576, MODULE_576
REACTOME_MTOR_SIGNALING, REACTOME_MTOR_SIGNALING
MIR4735_3P, MIR4735_3P
MIR2276_3P, MIR2276_3P
AAGGGAT_MIR188, AAGGGAT_MIR188
GSE15624_3H_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN, GSE15624_3H_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN
HP_ABNORMALITY_OF_EARLOBE, HP_ABNORMALITY_OF_EARLOBE
MIR5581_3P, MIR5581_3P
MIR383_3P, MIR383_3P
GSE13485_PRE_VS_POST_YF17D_VACCINATION_PBMC_UP, GSE13485_PRE_VS_POST_YF17D_VACCINATION_PBMC_UP
GOBP_ERBB_SIGNALING_PATHWAY, GOBP_ERBB_SIGNALING_PATHWAY
BURTON_ADIPOGENESIS_I1, BURTON_ADIPOGENESIS_I1
HP_PECTUS_CARINATUM, HP_PECTUS_CARINATUM
MIR6529_3P, MIR6529_3P
HP_ABNORMAL_FINGERTIP_MORPHOLOGY, HP_ABNORMAL_FINGERTIP_MORPHOLOGY
WP_ENDOMETRIAL_CANCER, WP_ENDOMETRIAL_CANCER
PID_TAPK3_PATHWAY, PID_TAPK3_PATHWAY
GOCC_TRANSCRIPTION_REPRESSOR_COMPLEX, GOCC_TRANSCRIPTION_REPRESSOR_COMPLEX
HP_ABNORMALITY_OF_THE_COLUMELLA, HP_ABNORMALITY_OF_THE_COLUMELLA
MIR203A_5P, MIR203A_5P
MIR581, MIR581
RIZ_ERYTHROID_DIFFERENTIATION_CCNE1, RIZ_ERYTHROID_DIFFERENTIATION_CCNE1
REACTOME_ENERGY_DEPENDENT_REGULATION_OF_MTOR_BY_LKB1_AMPK, REACTOME_ENERGY_DEPENDENT_REGULATION_OF_MTOR_BY_LKB1_AMPK
MIR155_3P, MIR155_3P
GOBP_NEGATIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION
WP_EGFR_TYROSINE_KINASE_INHIBITOR_RESISTANCE, WP_EGFR_TYROSINE_KINASE_INHIBITOR_RESISTANCE
PID_IL8_CXCR2_PATHWAY, PID_IL8_CXCR2_PATHWAY
PID_KIT_PATHWAY, PID_KIT_PATHWAY
GOBP_EPIDERMAL_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_EPIDERMAL_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
ATTACAT_MIR380BP, ATTACAT_MIR380BP
REACTOME_RHOH_GTPASE_CYCLE, REACTOME_RHOH_GTPASE_CYCLE
CROONQUIST_NRAS_VS_STROMAL_STIMULATION_UP, CROONQUIST_NRAS_VS_STROMAL_STIMULATION_UP
MIR1202, MIR1202
WP_NOTCH_SIGNALING_PATHWAY_NETPATH, WP_NOTCH_SIGNALING_PATHWAY_NETPATH
MIR6514_3P, MIR6514_3P
GOBP_PROTEIN_K63_LINKED_DEUBIQUITINATION, GOBP_PROTEIN_K63_LINKED_DEUBIQUITINATION
WP_CORTICOTROPINRELEASING_HORMONE_SIGNALING_PATHWAY, WP_CORTICOTROPINRELEASING_HORMONE_SIGNALING_PATHWAY
MIR434_3P, MIR434_3P
HP_APLASIA_HYPOPLASIA_OF_THE_EYEBROW, HP_APLASIA_HYPOPLASIA_OF_THE_EYEBROW
GOBP_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION, GOBP_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION
HP_SPARSE_EYEBROW, HP_SPARSE_EYEBROW
GOMF_BETA_CATENIN_BINDING, GOMF_BETA_CATENIN_BINDING
MIR3972, MIR3972
MIR708_5P, MIR708_5P
TATCTGG_MIR488, TATCTGG_MIR488
GOBP_LYSOSOME_LOCALIZATION, GOBP_LYSOSOME_LOCALIZATION
GOCC_PERIKARYON, GOCC_PERIKARYON
PID_NOTCH_PATHWAY, PID_NOTCH_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_CELL_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_CELL_DEVELOPMENT
GOCC_RETROMER_COMPLEX, GOCC_RETROMER_COMPLEX
GOBP_NEGATIVE_REGULATION_OF_DNA_BINDING, GOBP_NEGATIVE_REGULATION_OF_DNA_BINDING
MIR921, MIR921
GOBP_POSITIVE_REGULATION_OF_NEURON_DEATH, GOBP_POSITIVE_REGULATION_OF_NEURON_DEATH
MIR26_5P, MIR26_5P
MIR6511A_3P, MIR6511B_3P, MIR6511A_3P, MIR6511B_3P
MIR4445_5P, MIR4445_5P
MIR576_3P, MIR576_3P
YUAN_ZNF143_PARTNERS, YUAN_ZNF143_PARTNERS
GOBP_POSITIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION
GOBP_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY, GOBP_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY
MIR474_3P, MIR474_3P
GOMF_PHOSPHATIDYLINOSITOL_3_PHOSPHATE_BINDING, GOMF_PHOSPHATIDYLINOSITOL_3_PHOSPHATE_BINDING
MIR7843_5P, MIR7843_5P
MIR6861_3P, MIR6861_3P
MIR182_3P, MIR182_3P
GOBP_CENTRAL_NERVOUS_SYSTEM_NEURON_DEVELOPMENT, GOBP_CENTRAL_NERVOUS_SYSTEM_NEURON_DEVELOPMENT
HP_PANCREATIC_ADENOCARCINOMA, HP_PANCREATIC_ADENOCARCINOMA
MESIAHONXAI_01, MESIAHONXAI_01
GOBP_RESPONSE_TO_EPIDERMAL_GROWTH_FACTOR, GOBP_RESPONSE_TO_EPIDERMAL_GROWTH_FACTOR
PID_IL12_2PATHWAY, PID_IL12_2PATHWAY
GOBP_BONE_CELL_DEVELOPMENT, GOBP_BONE_CELL_DEVELOPMENT
GOBP_CENTRAL_NERVOUS_SYSTEM_NEURON_DIFFERENTIATION, GOBP_CENTRAL_NERVOUS_SYSTEM_NEURON_DIFFERENTIATION
MIR591, MIR591
MIR6658_3P, MIR6658_3P
MIR1306_5P, MIR1306_5P
BIOCARTA_CSK3_PATHWAY, BIOCARTA_CSK3_PATHWAY
GOBP_PROTEIN_TARGETING_TO_LYSOSOME, GOBP_PROTEIN_TARGETING_TO_LYSOSOME
MIR1289, MIR1289
GOBP_MATURATION_OF_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_S5U_RRNA_5_3S_RRNA_LSU_RRNA, GOBP_MATURATION_OF_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_S5U_RRNA_5_3S_RRNA_LSU_RRNA
MIR5009_5P, MIR5009_5P
GSE9946_LISTERIA_INF_MATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP, GSE9946_LISTERIA_INF_MATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP
MIR432_3P, MIR432_3P
MIR5694, MIR5694
WP_COPPER_HOMEOSTASIS, WP_COPPER_HOMEOSTASIS
MIR490_3P, MIR490_3P
SIG_IL4RECEPTOR_IN_B_LYPHOCYTES, SIG_IL4RECEPTOR_IN_B_LYPHOCYTES
GOBP_REGULATION_OF_VIRAL_INDUCED_CYTOPLASMIC_PATTERN_RECOGNITION, GOBP_REGULATION_OF_VIRAL_INDUCED_CYTOPLASMIC_PATTERN_RECOGNITION
HP_DEEP_PLANTAR_CREASES, HP_DEEP_PLANTAR_CREASES
GOBP INNER_CELL_MASS_CELL_PROLIFERATION, GOBP INNER_CELL_MASS_CELL_PROLIFERATION
GOMF_TRANSCRIPTION_COACTIVATOR_BINDING, GOMF_TRANSCRIPTION_COACTIVATOR_BINDING
GOBP_REGULATION_OF_LIPID_LOCALIZATION, GOBP_REGULATION_OF_LIPID_LOCALIZATION
GOBP_STEROL_TRANSPORT, GOBP_STEROL_TRANSPORT
MIR5589_5P, MIR5589_5P
SCHEIDERREIT_IKK_TARGETS, SCHEIDERREIT_IKK_TARGETS
MIR556_5P, MIR556_5P
REACTOME_REGULATION_OF_MECP2_EXPRESSION_AND_ACTIVITY, REACTOME_REGULATION_OF_MECP2_EXPRESSION_AND_ACTIVITY
BAUS_TP2_TARGETS_UP, BAUS_TP2_TARGETS_UP
GOBP_REGULATION_OF_RIG_1_SIGNALING_PATHWAY, GOBP_REGULATION_OF_RIG_1_SIGNALING_PATHWAY
KANG_DOXORUBICIN_RESISTANCE_DN, KANG_DOXORUBICIN_RESISTANCE_DN
GOBP_REGULATION_OF_POLYSACCHARIDE_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_POLYSACCHARIDE_BIOSYNTHETIC_PROCESS
HP_THIN_EYEBROW, HP_THIN_EYEBROW
MIR548AO_3P, MIR548AO_3P
MIR6834_3P, MIR6834_3P
HP_CUPPED_EAR, HP_CUPPED_EAR
KEGG_DORSO_VENTRAL_AXIS_FORMATION, KEGG_DORSO_VENTRAL_AXIS_FORMATION
REACTOME_EXTRA_NUCLEAR_ESTROGEN_SIGNALING, REACTOME_EXTRA_NUCLEAR_ESTROGEN_SIGNALING
GOBP_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE, GOBP_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE
REACTOME_CONSTITUTIVE_SIGNALING_BY_AKT1_E17K_IN_CANCER, REACTOME_CONSTITUTIVE_SIGNALING_BY_AKT1_E17K_IN_CANCER
BIOCARTA_HES_PATHWAY, BIOCARTA_HES_PATHWAY
REACTOME_FOXO_MEDIATED_TRANSCRIPTION_OF_OXIDATIVE_STRESS_METABOLIC_AND_NEURONAL_GENES, REACTOME_FOXO_MEDIATED_TRANSCRIPTION_OF_OXIDATIVE_STRESS_METABOLIC_AND_NEURONAL_GENES
MIR4761_3P, MIR4761_3P
MIR6503_3P, MIR6503_3P
REACTOME_SUMOYLATION_OF_DNA_METHYLATION_PROTEINS, REACTOME_SUMOYLATION_OF_DNA_METHYLATION_PROTEINS
HP_GNANULOCYTOPENIA, HP_GNANULOCYTOPENIA
MALTA_CURATED_STEMNESS_MARKERS, MALTA_CURATED_STEMNESS_MARKERS
OLSSON_E2F3_TARGETS_UP, OLSSON_E2F3_TARGETS_UP
PID_IL23_PATHWAY, PID_IL23_PATHWAY
GOBP_GLANDULAR_EPITHELIAL_CELL_DEVELOPMENT, GOBP_GLANDULAR_EPITHELIAL_CELL_DEVELOPMENT
VALK_AML_CLUSTER_1, VALK_AML_CLUSTER_1
GOBP_CENTRAL_NERVOUS_SYSTEM_NEURON_AXONOGENESIS, GOBP_CENTRAL_NERVOUS_SYSTEM_NEURON_AXONOGENESIS
HP_POSTERIOR_SUBCAPSULAR_CATARACT, HP_POSTERIOR_SUBCAPSULAR_CATARACT
WP_NANOPARTICLE_TRIGGERED_AUTOPHAGIC_CELL_DEATH, WP_NANOPARTICLE_TRIGGERED_AUTOPHAGIC_CELL_DEATH
MIR4712_3P, MIR4712_3P
REACTOME_CASPASE_ACTIVATION_VIA_DEPENDENCE_RECEPTORS_IN_THE_ABSENCE_OF_LIGAND, REACTOME_CASPASE_ACTIVATION_VIA_DEPENDENCE_RECEPTORS_IN_THE_ABSENCE_OF_LIGAND
GOBP_REGULATION_OF_LIPID_STORAGE, GOBP_REGULATION_OF_LIPID_STORAGE
GOCC_PRC1_COMPLEX, GOCC_PRC1_COMPLEX
GOBP_ENTEROENDOCRINE_CELL_DIFFERENTIATION, GOBP_ENTEROENDOCRINE_CELL_DIFFERENTIATION
HP_INCREASED_CIRCULATING_IGM_LEVEL, HP_INCREASED_CIRCULATING_IGM_LEVEL
MIR1468_5P, MIR1468_5P
GOBP_MARGINAL_ZONE_B_CELL_DIFFERENTIATION, GOBP_MARGINAL_ZONE_B_CELL_DIFFERENTIATION
HP_VERTEBRAL_COMPRESSION_FRACTURES, HP_VERTEBRAL_COMPRESSION_FRACTURES
GOBP_REGULATION_OF_NEURON_DIFFERENTIATION, GOBP_REGULATION_OF_NEURON_DIFFERENTIATION
HP_BICONCAVE_VERTEBRAL_BODIES, HP_BICONCAVE_VERTEBRAL_BODIES
WP_SEROTONIN_RECEPTOR_467_AND_NRC3_SIGNALING, WP_SEROTONIN_RECEPTOR_467_AND_NRC3_SIGNALING
SCHLINGEMANN_SKIN_CARCINOMA, SCHLINGEMANN_SKIN_CARCINOMA
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEUS, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEUS
GOBP_REGULATION_OF_POLYSACCHARIDE_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_POLYSACCHARIDE_BIOSYNTHETIC_PROCESS
GOBP_TYPE_B_PANCREATIC_CELL_DEVELOPMENT, GOBP_TYPE_B_PANCREATIC_CELL_DEVELOPMENT
GOBP_NUCLEOTIDE_BINDING_OLIGOMERIZATION_DOMAIN_CONTAINING_2_SIGNALING_PATHWAY, GOBP_NUCLEOTIDE_BINDING_OLIGOMERIZATION_DOMAIN_CONTAINING_2_SIGNALING_PATHWAY
WP_EXRNA_MECHANISM_OF_ACTION_AND_BIOGENESIS, WP_EXRNA_MECHANISM_OF_ACTION_AND_BIOGENESIS
BIOCARTA_PTEN_PATHWAY, BIOCARTA_PTEN_PATHWAY
HP_SHORT_CLAVICLES, HP_SHORT_CLAVICLES
HP_INTUSUSCEPTION, HP_INTUSUSCEPTION
HP_IVORY_EPHIPHYSES, HP_IVORY_EPHIPHYSES
GOBP_REGULATION_OF_TRIGLYCERIDE_METABOLIC_PROCESS, GOBP_REGULATION_OF_TRIGLYCERIDE_METABOLIC_PROCESS
REACTOME_G_PROTEIN_BETA_GAMMA_SIGNALING, REACTOME_G_PROTEIN_BETA_GAMMA_SIGNALING
MIR3147, MIR3147
GOBP_ENDOCRINE_PANCREAS_DEVELOPMENT, GOBP_ENDOCRINE_PANCREAS_DEVELOPMENT
REACTOME_PBP_REGULATES_TP53_ACETYLATION, REACTOME_PBP_REGULATES_TP53_ACETYLATION
GOBP_MIRNA_LOADING_ONTO_RISC_INVOLVED_IN_GENE_SILENCING_BY_MIRNA, GOBP_MIRNA_LOADING_ONTO_RISC_INVOLVED_IN_GENE_SILENCING_BY_MIRNA
GOBP_REGULATION_OF_NEUTROPHIL_MEDIATED_CYTOTOXICITY, GOBP_REGULATION_OF_NEUTROPHIL_MEDIATED_CYTOTOXICITY
GOBP_INTERLEUKIN_18_MEDIATED_SIGNALING_PATHWAY, GOBP_INTERLEUKIN_18_MEDIATED_SIGNALING_PATHWAY
GOBP_SNRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_III, GOBP_SNRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_III
GOBP_CELLULAR_RESPONSE_TO_PROSTAGLANDIN_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_PROSTAGLANDIN_STIMULUS
GOBP_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS, GOBP_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS
chr14q13, chr14q13
GOBP_POSITIVE_REGULATION_OF_RIG_1_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_RIG_1_SIGNALING_PATHWAY
HP_CERMINOMA, HP_CERMINOMA
MIR3768_5P, MIR376C_5P, MIR3768_5P, MIR376C_5P
GOBP_REGULATION_OF_AXON_GUIDANCE, GOBP_REGULATION_OF_AXON_GUIDANCE
GOMF_HMG_BOX_DOMAIN_BINDING, GOMF_HMG_BOX_DOMAIN_BINDING
GOBP_RESPONSE_TO_CATECHOLAMINE, GOBP_RESPONSE_TO_CATECHOLAMINE
HP_ABNORMALITY_OF_NAIL_COLOR, HP_ABNORMALITY_OF_NAIL_COLOR
HP_OVERWEIGHT, HP_OVERWEIGHT
WP_SMALL_LIGAND_GPCRS, WP_SMALL_LIGAND_GPCRS
HP_ABNORMAL_SWEAT_GLAND_MORPHOLOGY, HP_ABNORMAL_SWEAT_GLAND_MORPHOLOGY
MULLIGAN_NTF3_SIGNALING_VIA_INSR_AND_IGFIR_DN, MULLIGAN_NTF3_SIGNALING_VIA_INSR_AND_IGFIR_DN
PETRETTO_BLOOD_PRESSURE_DN, PETRETTO_BLOOD_PRESSURE_DN
HP_SHORT_NAIL, HP_SHORT_NAIL
CHESLER_BRAIN_D6MIT150_QTL_TRANS, CHESLER_BRAIN_D6MIT150_QTL_TRANS
HP_PROMINENT_UMBILICUS, HP_PROMINENT_UMBILICUS
BIOCARTA_CBR1_PATHWAY, BIOCARTA_CBR1_PATHWAY
GOBP_REGULATION_OF_FC_GAMMA_RECEPTOR_SIGNALING_PATHWAY_INVOLVED_IN_PHAGOCYTOSIS, GOBP_REGULATION_OF_FC_GAMMA_RECEPTOR_SIGNALING_PATHWAY_INVOLVED_IN_PHAGOCYTOSIS
GOCC_WNT_SIGNALOSOME, GOCC_WNT_SIGNALOSOME
GOBP_CENTRAL_NERVOUS_SYSTEM_PROJECTION_NEURON_AXONOGENESIS, GOBP_CENTRAL_NERVOUS_SYSTEM_PROJECTION_NEURON_AXONOGENESIS
HP_SPARSE_LATERAL_EYEBROW, HP_SPARSE_LATERAL_EYEBROW
GOBP_POSITIVE_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_SALMON_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_SALMON_DN
REACTOME_G_BETA_GAMMA_SIGNALING_THROUGH_PDKGAMMA, REACTOME_G_BETA_GAMMA_SIGNALING_THROUGH_PDKGAMMA
PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_5, PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_5
REACTOME_INTERLEUKIN_15_SIGNALING, REACTOME_INTERLEUKIN_15_SIGNALING
REACTOME_NOTCH2_INTRACELLULAR_DOMAIN_REGULATES_TRANSCRIPTION, REACTOME_NOTCH2_INTRACELLULAR_DOMAIN_REGULATES_TRANSCRIPTION
GOBP_MICROTUBULE_ANCHORING_AT_MICROTUBULE_ORGANIZING_CENTER, GOBP_MICROTUBULE_ANCHORING_AT_MICROTUBULE_ORGANIZING_CENTER
KAZMIN_PBMC_P_FALCIPARUM_RTSS_A01_AGE_UNKNOWN_CORRELATED_WITH_PROTECTION_56DY_POSITIVE, KAZMIN_PBMC_P_FALCIPARUM_RTSS_A01_AGE_UNKNOWN_CORRELATED_WITH_PROTECTION_56DY_POSITIVE
HP_HYPOPLASIA_OF_THE_PRIMARY_TEETH, HP_HYPOPLASIA_OF_THE_PRIMARY_TEETH
HP_ABSENT_FRONTAL_SINUSES, HP_ABSENT_FRONTAL_SINUSES
GOBP_PROTEIN_LOCALIZATION_TO_BASOLATERAL_PLASMA_MEMBRANE, GOBP_PROTEIN_LOCALIZATION_TO_BASOLATERAL_PLASMA_MEMBRANE
HP_IMPAIRED_GLUCOSE_TOLERANCE, HP_IMPAIRED_GLUCOSE_TOLERANCE
GOBP_REGULATION_OF_CHOLESTEROL_EFFLUX, GOBP_REGULATION_OF_CHOLESTEROL_EFFLUX
GOBP_MUSCLE_CELL_APOPTOTIC_PROCESS, GOBP_MUSCLE_CELL_APOPTOTIC_PROCESS
HP_ACNE_INVERSA, HP_ACNE_INVERSA
BIOCARTA_TERC_PATHWAY, BIOCARTA_TERC_PATHWAY
HP_SCLEROSIS_OF_FINGER_PHALANX, HP_SCLEROSIS_OF_FINGER_PHALANX
GOBP_GASTRULATION_WITH_MOUTH_FORMING_SECOND, GOBP_GASTRULATION_WITH_MOUTH_FORMING_SECOND
MIR1182, MIR1182
GOBP_ROUNDABOUT_SIGNALING_PATHWAY, GOBP_ROUNDABOUT_SIGNALING_PATHWAY
HP_EVERSION_OF_LATERAL_THIRD_OF_LOWER_EYELIDS, HP_EVERSION_OF_LATERAL_THIRD_OF_LOWER_EYELIDS
HP_PROMINENT_ANTITRAGUS, HP_PROMINENT_ANTITRAGUS
GOBP_GLOMERULUS_MORPHOGENESIS, GOBP_GLOMERULUS_MORPHOGENESIS
REACTOME_INTERLEUKIN_18_SIGNALING, REACTOME_INTERLEUKIN_18_SIGNALING
REACTOME_GLUATHIONE_SYNTHESIS_AND_RECYCLING, REACTOME_GLUATHIONE_SYNTHESIS_AND_RECYCLING
GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY