

GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_CELL_MAST_CELL_CELL_UP, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_CELL_MAST_CELL_CELL_UP

GSE37416_0H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_0H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_UP

GSE37416_0H_VS_12H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_0H_VS_12H_F_TULARENSIS_LVS_NEUTROPHIL_UP

GSE25988_IL4_VS_IL4_AND_ROSGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_UP, GSE25988_IL4_VS_IL4_AND_ROSGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_UP

GSE369_S0CS3_KO_VS_WT_LIVER_DN, GSE369_S0CS3_KO_VS_WT_LIVER_DN

GSE37416_12H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_12H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_UP

GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_DN, GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_DN

GSE4984_LPS_VS_VEHICLE_CTRL_TREATED_DC_DN, GSE4984_LPS_VS_VEHICLE_CTRL_TREATED_DC_DN

GSE32255_WT_UNSTIM_VS_JM1DZ_KNOCKDOWN_4H_LPS_STIM_DC_UP, GSE32255_WT_UNSTIM_VS_JM1DZ_KNOCKDOWN_4H_LPS_STIM_DC_UP

GSE21360_TERTIARY_VS_QUATERNARY_MEMORY_CD8_TCELL_UP, GSE21360_TERTIARY_VS_QUATERNARY_MEMORY_CD8_TCELL_UP

GSE22611_NOD2_VS_CTRL_TRANSDUCED_HEK293T_CELL_DN, GSE22611_NOD2_VS_CTRL_TRANSDUCED_HEK293T_CELL_DN

GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_24H_DN, GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_24H_DN

GSE30971_WBP7_HET_VS_KO_MACROPHAGE_2H_LPS_STIM_DN, GSE30971_WBP7_HET_VS_KO_MACROPHAGE_2H_LPS_STIM_DN

GSE4748_CTRL_VS_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP, GSE4748_CTRL_VS_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP

GSE30971_WBP7_HET_VS_KO_MACROPHAGE_DN, GSE30971_WBP7_HET_VS_KO_MACROPHAGE_DN

GSE39556_CD8A_DC_VS_NK_CELL_DN, GSE39556_CD8A_DC_VS_NK_CELL_DN

GSE6674_ANTI_IJGM_VS_ANTI_IJGM_AND_CPG_STIM_BCELL_UP, GSE6674_ANTI_IJGM_VS_ANTI_IJGM_AND_CPG_STIM_BCELL_UP

GSE27786_LIN_NEG_VS_NEUTROPHIL_DN, GSE27786_LIN_NEG_VS_NEUTROPHIL_DN

GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_UP, GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_UP

GSE1635_JFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP, GSE1635_JFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP

GSE7852_TREG_VS_TCONV_DN, GSE7852_TREG_VS_TCONV_DN

GSE11961_FOLLICULAR_BCELL_VS_GERMINAL_CENTER_BCELL_DAY40_UP, GSE11961_FOLLICULAR_BCELL_VS_GERMINAL_CENTER_BCELL_DAY40_UP

GSE21670_UNTREATED_VS_TGFB_TREATED_CD4_TCELL_DN, GSE21670_UNTREATED_VS_TGFB_TREATED_CD4_TCELL_DN

GCAA4AAA_MIR129, GCAA4AAA_MIR129

GSE2180_NON_TFH_VS_TFH_CD4_TCELL_UP, GSE2180_NON_TFH_VS_TFH_CD4_TCELL_UP

GSE1831_CORTX_VS_MEDULLA_THYMUS_DN, GSE1831_CORTX_VS_MEDULLA_THYMUS_DN

GSE14699_NAIVE_VS_DELETIONAL_TOLERANCE_CD8_TCELL_DN, GSE14699_NAIVE_VS_DELETIONAL_TOLERANCE_CD8_TCELL_DN

MIR348_3P, MIR348_3P

GSE22611_UNSTIM_VS_6H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_DN, GSE22611_UNSTIM_VS_6H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_DN

MIR4782_3P, MIR4782_3P

MIR6756_3P, MIR6756_3P

GSE3982_DC_VS_NEUTROPHIL_DN, GSE3982_DC_VS_NEUTROPHIL_DN

IKEDA_MIR30_TARGETS_UP, IKEDA_MIR30_TARGETS_UP

MIR219A_5P, MIR219A_5P

GSE46606_UNSTIM_VS_CD40_IL2_IL5_DAY3_STIMULATED_BCELL_UP, GSE46606_UNSTIM_VS_CD40_IL2_IL5_DAY3_STIMULATED_BCELL_UP

GSE29164_DAYS_VS_DAY7_UNTREATED_MELANOMA_UP, GSE29164_DAYS_VS_DAY7_UNTREATED_MELANOMA_UP

MIR6882_5P, MIR6882_5P

MEON_FETAL_KIDNEY_9_ENDOTHELIAL_CELLS, MEON_FETAL_KIDNEY_9_ENDOTHELIAL_CELLS

GSE3982_EOSINOPHIL_VS_DC_UP, GSE3982_EOSINOPHIL_VS_DC_UP

GOMB_PHOSPHATASE_BINDING, GOMB_PHOSPHATASE_BINDING

GSE10240_CTRL_VS_IL7_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN, GSE10240_CTRL_VS_IL7_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN

GSE40277_GATA1_AND_SATB1_TRANSDUCED_CTRL_CD4_TCELL_DN, GSE40277_GATA1_AND_SATB1_TRANSDUCED_CTRL_CD4_TCELL_DN

GOMB_AUTOPHAGOSOME_ORGANIZATION, GOMB_AUTOPHAGOSOME_ORGANIZATION

GSE40274_CTRL_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_CTRL_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP

GSE20152_HTNFA_OVERPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_DN, GSE20152_HTNFA_OVERPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_DN

GSE15271_CXCR4_POS_VS_NEG_GC_BCELL_DN, GSE15271_CXCR4_POS_VS_NEG_GC_BCELL_DN

MIR3679_3P, MIR3679_3P

GSE25988_CTRL_VS_IL4_STIM_MACROPHAGE_DN, GSE25988_CTRL_VS_IL4_STIM_MACROPHAGE_DN

REACTOME_RHO_GTPASES_ACTIVATE_PKNS, REACTOME_RHO_GTPASES_ACTIVATE_PKNS

KEGG_NEUROTROPHIN_SIGNALING_PATHWAY, KEGG_NEUROTROPHIN_SIGNALING_PATHWAY

MIR4744, MIR4744

MIR190A_5P, MIR190A_5P

BIOCARTA_MAPK_PATHWAY, BIOCARTA_MAPK_PATHWAY

HOLLERN_EMT_TUMOR_UP, HOLLERN_EMT_TUMOR_UP

KEGG_FOCAL_ADHESION, KEGG_FOCAL_ADHESION

MIR190B_5P, MIR190B_5P

GSE17721_PAMC5K4_VS_GADQUIMOD_0.5H_BMDC_DN, GSE17721_PAMC5K4_VS_GADQUIMOD_0.5H_BMDC_DN

GSE6259_BCELL_VS_CD8_TCELL_UP, GSE6259_BCELL_VS_CD8_TCELL_UP

GSE16450_IMMATURE_VS_MATURE_NEURON_CTRL_LINE_6H_IFNA_STIM_UP, GSE16450_IMMATURE_VS_MATURE_NEURON_CTRL_LINE_6H_IFNA_STIM_UP

GOMB_MEGAKARYOCYTE_DIFFERENTIATION, GOMB_MEGAKARYOCYTE_DIFFERENTIATION

GSE3920_IFNA_VS_IFNB_TREATED_ENDOTHELIAL_CELL_UP, GSE3920_IFNA_VS_IFNB_TREATED_ENDOTHELIAL_CELL_UP

KEGG_SYSTEMIC_LUPUS_ERYTHEMATOSUS, KEGG_SYSTEMIC_LUPUS_ERYTHEMATOSUS

MIR580_3P, MIR580_3P

GOMB_PROTEIN_PHOSPHATASE_BINDING, GOMB_PROTEIN_PHOSPHATASE_BINDING

WP_REGULATION_OF_ACTIN_CYTOSKELETON, WP_REGULATION_OF_ACTIN_CYTOSKELETON

GOCC_CYTOPLASMIC_SIDE_OF_MEMBRANE, GOCC_CYTOPLASMIC_SIDE_OF_MEMBRANE

MIR1343_3P, MIR1343_3P

MULLIGHAN_NPM1_MUTATED_SIGNATURE_2_UP, MULLIGHAN_NPM1_MUTATED_SIGNATURE_2_UP

GOMB_JNK_CASCADE, GOMB_JNK_CASCADE

GSE22886_NEUTROPHIL_VS_MONOCYTE_UP, GSE22886_NEUTROPHIL_VS_MONOCYTE_UP

KEGG_COLORECTAL_CANCER, KEGG_COLORECTAL_CANCER

MIR4680_5P, MIR4680_5P

MIR6858_5P, MIR6858_5P

HP_ABNORMAL_DISTAL_PHALANX_MORPHOLOGY_OF_FINGER, HP_ABNORMAL_DISTAL_PHALANX_MORPHOLOGY_OF_FINGER

AMIT_EGF_RESPONSE_120_HELA, AMIT_EGF_RESPONSE_120_HELA

MIR378A_5P, MIR378A_5P

GGTGTGT_MIR329, GGTGTGT_MIR329

MIR6132, MIR6132

MIR4308, MIR4308

KEGG_ACUTE_MYELOID_LEUKEMIA, KEGG_ACUTE_MYELOID_LEUKEMIA

MIR329_3P, MIR329_3P

MIR532_5P, MIR532_5P

MIR362_3P, MIR362_3P

MIR6730_3P, MIR6730_3P

GSE13306_RA_VS_UNTREATED_TREG_DN, GSE13306_RA_VS_UNTREATED_TREG_DN

AIZARANI_LIVER_C28_NK_NKT_CELLS_6, AIZARANI_LIVER_C28_NK_NKT_CELLS_6

REACTOME_REGULATION_OF_PTEIN_GENE_TRANSCRIPTION, REACTOME_REGULATION_OF_PTEIN_GENE_TRANSCRIPTION

GOMB_PHOSPHATIDYLYNOSITOL_3_KINASE_SIGNALING, GOMB_PHOSPHATIDYLYNOSITOL_3_KINASE_SIGNALING

KEGG_PANCREATIC_CANCER, KEGG_PANCREATIC_CANCER

HP_CARDIAC_ARREST, HP_CARDIAC_ARREST

TNCATNTCCYR_UNKNOWN, TNCATNTCCYR_UNKNOWN

WP_DNA_DAMAGE_RESPONSE_ONLY_ATM_DEPENDENT, WP_DNA_DAMAGE_RESPONSE_ONLY_ATM_DEPENDENT

MIR6509_5P, MIR6509_5P

GOMB_REGULATION_OF_PHOSPHATIDYLYNOSITOL_3_KINASE_SIGNALING, GOMB_REGULATION_OF_PHOSPHATIDYLYNOSITOL_3_KINASE_SIGNALING

LEIN_NEURON_MARKERS, LEIN_NEURON_MARKERS

MIR3117_5P, MIR3117_5P

GOCC_I_BAND, GOCC_I_BAND

HP_INTESTINAL_POLYP, HP_INTESTINAL_POLYP

KEGG_GLIOMA, KEGG_GLIOMA

ATGTAGC_MIR221, ATGTAGC_MIR221

REACTOME_RESPONSE_TO_ELEVATED_PLATELET_CYTOSOLIC_CA2, REACTOME_RESPONSE_TO_ELEVATED_PLATELET_CYTOSOLIC_CA2

MIR6749_3P, MIR6749_3P

MIR582_3P, MIR582_3P

GSE29615_CTRL_VS_LAIV_FLU_VACCINE_PBMIC_UP, GSE29615_CTRL_VS_LAIV_FLU_VACCINE_PBMIC_UP

GSE37505_NOD_VS_C57BL6_IRES_GFP_TREG_DN, GSE37505_NOD_VS_C57BL6_IRES_GFP_TREG_DN

ERWIN_COHEN_BLOOD_VACCINE_TC_83_AGE_23_48YO_VACCINATED_VS_CONTROL_2DY_DN, ERWIN_COHEN_BLOOD_VACCINE_TC_83_AGE_23_48YO_VACCINATED_VS_CONTROL_2DY_DN

MIR1185_5P, MIR1185_5P

MIR6126, MIR6126

MIR6887_3P, MIR6887_3P

GOMB_REGULATION_OF_JNK_CASCADE, GOMB_REGULATION_OF_JNK_CASCADE

GOMB_PLATELET_DEGRANULATION, GOMB_PLATELET_DEGRANULATION

MIR6811_5P, MIR6811_5P

MIR6511B_5P, MIR6511B_5P

LASTOWSKA_COAMPLIFIED_WITH_MYCN, LASTOWSKA_COAMPLIFIED_WITH_MYCN

MIR4317, MIR4317

MIR4487, MIR4487

GSE6269_FLU_VS_STREP_PNEUMO_INF_PBMIC_DN, GSE6269_FLU_VS_STREP_PNEUMO_INF_PBMIC_DN

MIR7113_3P, MIR7113_3P

GOMB_REGULATION_OF_MONONUCLEAR_CELL_MIGRATION, GOMB_REGULATION_OF_MONONUCLEAR_CELL_MIGRATION

HP_NEOPLASM_OF_THE_PITUITARY_GLAND, HP_NEOPLASM_OF_THE_PITUITARY_GLAND

GOMB_LYSOPHOSPHOLIPASE_ACTIVITY, GOMB_LYSOPHOSPHOLIPASE_ACTIVITY

MIR4522, MIR4522

KEGG_LONG_TERM_DEPRESSION, KEGG_LONG_TERM_DEPRESSION

GOMB_POSITIVE_REGULATION_OF_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY, GOMB_POSITIVE_REGULATION_OF_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY

KEGG_VASCULAR_SMOOTH_MUSCLE_CONTRACTION, KEGG_VASCULAR_SMOOTH_MUSCLE_CONTRACTION

GSE7509_FCCRIIB_VS_TNFA_IL1B_IL6_PGE_STIM_DC_DN, GSE7509_FCCRIIB_VS_TNFA_IL1B_IL6_PGE_STIM_DC_DN

GSE4305_ISR_VS_FLU_INF_INFANT_PBMIC_DN, GSE4305_ISR_VS_FLU_INF_INFANT_PBMIC_DN

VANDESULIS_COMMD1_TARGETS_GROUP_3_UP, VANDESULIS_COMMD1_TARGETS_GROUP_3_UP

GSE46242_TH1_VS_ANGERIC_TH1_CD4_TCELL_DN, GSE46242_TH1_VS_ANGERIC_TH1_CD4_TCELL_DN

GOMB_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION, GOMB_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION

GSE29615_CTRL_VS_DAY7_LAIV_FLU_VACCINE_PBMIC_UP, GSE29615_CTRL_VS_DAY7_LAIV_FLU_VACCINE_PBMIC_UP

HP_DISPROPORTIONATE_TALL_STATURE, HP_DISPROPORTIONATE_TALL_STATURE

MIR3162_5P, MIR3162_5P

HP_METATARSUS_VALGUS, HP_METATARSUS_VALGUS

REACTOME_OVARIAN_TUMOR_DOMAIN_PROTEASES, REACTOME_OVARIAN_TUMOR_DOMAIN_PROTEASES

REACTOME_P3K_AKT_SIGNALING_IN_CANCER, REACTOME_P3K_AKT_SIGNALING_IN_CANCER

HP_INCREASED_CIRCULATING_PROLACTIN_CONCENTRATION, HP_INCREASED_CIRCULATING_PROLACTIN_CONCENTRATION

REACTOME_NEGATIVE_REGULATION_OF_THE_P3K_AKT_NETWORK, REACTOME_NEGATIVE_REGULATION_OF_THE_P3K_AKT_NETWORK

KEGG_MELANOMA, KEGG_MELANOMA

MIR144_5P, MIR144_5P

PID_ERBB2_ERBB3_PATHWAY, PID_ERBB2_ERBB3_PATHWAY

LI_PBMIC_MENACTRA_AGE_18_45YO_CORRELATED_WITH_ANTL_DT_ANTIODY_3DY_POSITIVE, LI_PBMIC_MENACTRA_AGE_18_45YO_CORRELATED_WITH_ANTL_DT_ANTIODY_3DY_POSITIVE

REACTOME_NEPHRIN_FAMILY_INTERACTIONS, REACTOME_NEPHRIN_FAMILY_INTERACTIONS

GOMB_ANIMAL_ORGAN_FORMATION, GOMB_ANIMAL_ORGAN_FORMATION

MIR6827_5P, MIR6827_5P

PRMT5_TARGET_GENES, PRMT5_TARGET_GENES

BIOCARTA KERATINOCYTE_PATHWAY, BIOCARTA KERATINOCYTE_PATHWAY

TRAVAGLINI_LONG_TERM_EFFECTS_CELL_TRAVAGLINI_LONG_TERM_EFFECTS_CELL

DAVICION1_RHABDOMYOSARCOMA_PAX_FOXP1_FUSION_UP, DAVICION1_RHABDOMYOSARCOMA_PAX_FOXP1_FUSION_UP

GOMB_REGULATION_OF_LYMPHOCYTE_APOPTOTIC_PROCESS, GOMB_REGULATION_OF_LYMPHOCYTE_APOPTOTIC_PROCESS

MIR6515_5P, MIR6515_5P

BIOCARTA_RAC1_PATHWAY, BIOCARTA_RAC1_PATHWAY

WP_ESC_PLURIPOTENCY_PATHWAYS, WP_ESC_PLURIPOTENCY_PATHWAYS

GOMB_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOMB_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION

EV1L_I2, EV1L_I2

GOCC_LAMELLOPODIUM_MEMBRANE, GOCC_LAMELLOPODIUM_MEMBRANE

SABATES_COLORECTAL_ADENOMA_SIZE_UP, SABATES_COLORECTAL_ADENOMA_SIZE_UP

REACTOME_RHO_GTPASES_ACTIVATE_CIT, REACTOME_RHO_GTPASES_ACTIVATE_CIT

MIR4756_3P, MIR4756_3P

REACTOME_FORMATION_OF_FIBRIN_CLOT_CLOTTING_CASCADE, REACTOME_FORMATION_OF_FIBRIN_CLOT_CLOTTING_CASCADE

GOMB_NEGATIVE_REGULATION_OF_TRANSMEMBRANE_TRANSPORT, GOMB_NEGATIVE_REGULATION_OF_TRANSMEMBRANE_TRANSPORT

ZIRN_TRETINOIN_RESPONSE_WT1_UP, ZIRN_TRETINOIN_RESPONSE_WT1_UP

DESCARTES_FETAL_EYE_SMOOTH_MUSCLE_CELLS, DESCARTES_FETAL_EYE_SMOOTH_MUSCLE_CELLS

GOMB_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GOMB_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS

KEGG_MELANOGENESIS, KEGG_MELANOGENESIS

GOMB_ENERGY_RESERVE_METABOLIC_PROCESS, GOMB_ENERGY_RESERVE_METABOLIC_PROCESS

BIOCARTA_P38MAPK_PATHWAY, BIOCARTA_P38MAPK_PATHWAY

MIR4489, MIR4489

WP_P38_MAPK_SIGNALING_PATHWAY, WP_P38_MAPK_SIGNALING_PATHWAY

GOMB_REGULATION_OF_FATTY_ACID_BETA_OXIDATION, GOMB_REGULATION_OF_FATTY_ACID_BETA_OXIDATION

GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_SALMON_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_SALMON_DN

GOMB_POSITIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOMB_POSITIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY

VERNOCHET_ADIPOGENESIS, VERNOCHET_ADIPOGENESIS

GOMB_MEMORY, GOMB_MEMORY

GOMB_HISTIDINE_CATABOLIC_PROCESS, GOMB_HISTIDINE_CATABOLIC_PROCESS

REACTOME_HISTIDINE_CATABOLISM, REACTOME_HISTIDINE_CATABOLISM

MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_9, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_9

GOMB_EMBRYONIC_HEART_TUBE_MORPHOGENESIS, GOMB_EMBRYONIC_HEART_TUBE_MORPHOGENESIS

GOMB_EMBRYONIC_HEART_TUBE_DEVELOPMENT, GOMB_EMBRYONIC_HEART_TUBE_DEVELOPMENT

GOMB_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOMB_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION

GOMB_ACTIVATION_OF_PROTEIN_LOCALIZATION_TO_CENTROSOME, GOMB_ACTIVATION_OF_PROTEIN_LOCALIZATION_TO_CENTROSOME

GOMB_T_CELL_APOPTOTIC_PROCESS, GOMB_T_CELL_APOPTOTIC_PROCESS

REACTOME_ACTIVATION_OF_MATRIX_METALLOPROTEINASES, REACTOME_ACTIVATION_OF_MATRIX_METALLOPROTEINASES

GOMB_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GOMB_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS

HP_RIGHT_BUNDLE_BRANCH_BLOCK, HP_RIGHT_BUNDLE_BRANCH_BLOCK

MIR6501_5P, MIR6501_5P

MIR4749_3P, MIR4749_3P

LHX3_TARGET_GENES, LHX3_TARGET_GENES

GOCC_PLATELET_ALPHA_GRANULE, GOCC_PLATELET_ALPHA_GRANULE

REACTOME_RHO_GTPASES_ACTIVATE_PAKS, REACTOME_RHO_GTPASES_ACTIVATE_PAKS

GOMB_REGULATION_OF_LIPID_TRANSPORT, GOMB_REGULATION_OF_LIPID_TRANSPORT

WP_ASSOCIATION_BETWEEN_PHYSICOCHEMICAL_FEATURES_AND_TOXICITY_ASSOCIATED_PATHWAYS, WP_ASSOCIATION_BETWEEN_PHYSICOCHEMICAL_FEATURES_AND_TOXICITY_ASSOCIATED_PATHWAYS

ZHANG_ADIPOGENESIS_BY_BMP7, ZHANG_ADIPOGENESIS_BY_BMP7

GOMB_RESPONSE_TO_CHEMOKINE, GOMB_RESPONSE_TO_CHEMOKINE

HP_GASTROINTESTINAL_INFARCTIONS, HP_GASTROINTESTINAL_INFARCTIONS

HP_RESPRATORY_INSUFFICIENCY_DUE_TO_DEFECTIVE_CILIARY_CLEARANCE, HP_RESPRATORY_INSUFFICIENCY_DUE_TO_DEFECTIVE_CILIARY_CLEARANCE

VALK_AML_WITH_EV1L, VALK_AML_WITH_EV1L

GOMB_G_PROTEIN_BETA_GAMMA_SUBUNIT_COMPLEX_BINDING, GOMB_G_PROTEIN_BETA_GAMMA_SUBUNIT_COMPLEX_BINDING

GOMB_REGULATION_OF_ORGAN_GROWTH, GOMB_REGULATION_OF_ORGAN_GROWTH

MIR6724_5P, MIR6724_5P

GOMB_NEGATIVE_REGULATION_OF_MONONUCLEAR_CELL_MIGRATION, GOMB_NEGATIVE_REGULATION_OF_MONONUCLEAR_CELL_MIGRATION

MIR1277_3P, MIR1277_3P

WP_PURINERGIC_SIGNALING, WP_PURINERGIC_SIGNALING

chr9q21, chr9q21

GSE14413_UNSTIM_VS_IFNB_STIM_RAW264_CELLS_UP, GSE14413_UNSTIM_VS_IFNB_STIM_RAW264_CELLS_UP

GAUSSMANN_MLL_A4_FUSION_TARGETS_D_UP, GAUSSMANN_MLL_A4_FUSION_TARGETS_D_UP

GOMB_HISTIDINE_METABOLIC_PROCESS, GOMB_HISTIDINE_METABOLIC_PROCESS

GOMB_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_REGENERATION, GOMB_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_REGENERATION

GRADE_COLON_VS_RECTAL_CANCER_DN, GRADE_COLON_VS_RECTAL_CANCER_DN

GSE15750_WT_VS_TRAFK60_DAY10_EFF_CD8_TCELL_DN, GSE15750_WT_VS_TRAFK60_DAY10_EFF_CD8_TCELL_DN

GOMB_ICOSANOID_CATABOLIC_PROCESS, GOMB_ICOSANOID_CATABOLIC_PROCESS

BOYALUT_LIVER_CANCER_SUBLCLASS_G6_DN, BOYALUT_LIVER_CANCER_SUBLCLASS_G6_DN

GOMB_CARDIAC_NEURAL_CREST_CELL_DIFFERENTIATION_INVOLVED_IN_HEART_DEVELOPMENT, GOMB_CARDIAC_NEURAL_CREST_CELL_DIFFERENTIATION_INVOLVED_IN_HEART_DEVELOPMENT

HP_MYELOFIBROSIS, HP_MYELOFIBROSIS

HP_ABSENT_OUTER_DYNEIN_ARMS, HP_ABSENT_OUTER_DYNEIN_ARMS

WP_MFAP5_EFFECT_ON_PERMEABILITY_AND_MOTILITY_OF_ENDOTHELIAL_CELLS_VIA_CYTOSKELETON_REARRANGEMENT, WP_MFAP5_EFFECT_ON_PERMEABILITY_AND_MOTILITY_OF_ENDOTHELIAL_CELLS_VIA_CYTOSKELETON_REARRANGEMENT

GOMB_EMBRYONIC_CRANIAL_SKELETON_PROPHOGENESIS, GOMB_EMBRYONIC_CRANIAL_SKELETON_PROPHOGENESIS

GOMB_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS, GOMB_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS

GOMB_CARDIAC_NEURAL_CREST_CELL_DEVELOPMENT_INVOLVED_IN_OUTFLOW_TRACT_MORPHOGENESIS, GOMB_CARDIAC_NEURAL_CREST_CELL_DEVELOPMENT_INVOLVED_IN_OUTFLOW_TRACT_MORPHOGENESIS

GOMB_ACTIN_FILAMENT_NETWORK_FORMATION, GOMB_ACTIN_FILAMENT_NETWORK_FORMATION

BIOCARTA_BAD_PATHWAY, BIOCARTA_BAD_PATHWAY

GOMB_DEFINITIVE_HEMOPHOESIS, GOMB_DEFINITIVE_HEMOPHOESIS

GOMB_POSITIVE_REGULATION_OF_BONE_RESORPTION, GOMB_POSITIVE_REGULATION_OF_BONE_RESORPTION

ALONSO_METASTASIS_NEURAL_UP, ALONSO_METASTASIS_NEURAL_UP

GOMB_INNATE_IMMUNE_RESPONSE_IN_MUCOSA, GOMB_INNATE_IMMUNE_RESPONSE_IN_MUCOSA

GOMB_NEGATIVE_REGULATION_OF_LYMPHOCYTE_MIGRATION, GOMB_NEGATIVE_REGULATION_OF_LYMPHOCYTE_MIGRATION

SUZUKI_RESPONSE_TO_TSA, SUZUKI_RESPONSE_TO_TSA

MIRKELSEN_NPC_WITH_LCP_H3K27ME3, MIRKELSEN_NPC_WITH_LCP_H3K27ME3

GOMB_RESPONSE_TO_ESTROGEN, GOMB_RESPONSE_TO_ESTROGEN

GOMB_POSITIVE_REGULATION_OF_NEURON_PROJECTION_REGENERATION, GOMB_POSITIVE_REGULATION_OF_NEURON_PROJECTION_REGENERATION

GOMB_SYNAPSE_MATURATION, GOMB_SYNAPSE_MATURATION

GOMB_POSITIVE_REGULATION_OF_GLUCOSE_IMPORT, GOMB_POSITIVE_REGULATION_OF_GLUCOSE_IMPORT

GOMB_LEUKOTRIENE_B4_CATABOLIC_PROCESS, GOMB_LEUKOTRIENE_B4_CATABOLIC_PROCESS

GOMB_LEUKOTRIENE_B4_20_MONOOXYGENASE_ACTIVITY, GOMB_LEUKOTRIENE_B4_20_MONOOXYGENASE_ACTIVITY

HP_ABNORMALITY_OF_THE_INTERNAL_AUDITORY_CANAL, HP_ABNORMALITY_OF_THE_INTERNAL_AUDITORY_CANAL

GOMB_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_PROLIFERATION, GOMB_NEGATIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_PROLIFERATION

GOMB_LONG_CHAIN_FATTY_ACID_IMPORT_INTO_CELL, GOMB_LONG_CHAIN_FATTY_ACID_IMPORT_INTO_CELL

SASAKI_TARGETS_OF_1173_AND_T1G3_SASAKI_TARGETS_OF_1173_AND_T1G3

REACTOME_EICOSANIDS, REACTOME_EICOSANIDS

MODULE_557, MODULE_557

HP_CENTRAL_OPACIFICATION_OF_THE_CORNEA, HP_CENTRAL_OPACIFICATION_OF_THE_CORNEA

FAN_EMBRYONIC_CTX_IN_3_INTERNEURON, FAN_EMBRYONIC_CTX_IN_3_INTERNEURON

BIOCARTA_418B_PATHWAY, BIOCARTA_418B_PATHWAY

MIR454_3P, MIR454_3P

BIOCARTA_ERK5_PATHWAY, BIOCARTA_ERK5_PATHWAY

MIRKELSEN_ES_LCP_WITH_H3K4ME3, MIRKELSEN_ES_LCP_WITH_H3K4ME3

REACTOME_CREB1_PHOSPHORYLATION_THROUGH_NMDA_RECEPTOR_MEDIATED_ACTIVATION_OF_RAS_SIGNALING, REACTOME_CREB1_PHOSPHORYLATION_THROUGH_NMDA_RECEPTOR_MEDIATED_ACTIVATION_OF_RAS_SIGNALING

GOMB_POSITIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOMB_POSITIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION