GSE19825_NAIVE_VS_DAY3_EFF_CD8_TCELL_DN, GSE19825_NAIVE_VS_DAY3_EFF_CD8_TCELL_DN
GSE19825_NAIVE_VS_IL2RALOW_DAY3_EFF_CD8_TCELL_DN, GSE19825_NAIVE_VS_IL2RALOW_DAY3_EFF_CD8_TCELL_DN
GSE37563_WT_VS_CTLA4_KO_CD4_TCELL_D4_POST_IMMUNIZATION_DN, GSE37563_WT_VS_CTLA4_KO_CD4_TCELL_D4_POST_IMMUNIZATION_DN
GSE15624_CTRL_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN, GSE15624_CTRL_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN
GSE14308_TH2_VS_NATURAL_TREG_UP, GSE14308_TH2_VS_NATURAL_TREG_UP
GSE46242_CTRL_VS_EGR2_DELETED_ANERGIC_TH1_CD4_TCELL_UP, GSE46242_CTRL_VS_EGR2_DELETED_ANERGIC_TH1_CD4_TCELL_UP
GSE36891_UNSTIM_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN, GSE36891_UNSTIM_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN
GSE40274_FOXP3_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GOBP_MYELOID_CELL_HOMEOSTASIS, GOBP_MYELOID_CELL_HOMEOSTASIS
GOBP_NEGATIVE_REGULATION_OF_AUTOPHAGY, GOBP_NEGATIVE_REGULATION_OF_AUTOPHAGY
HP_DECREASED_FACIAL_EXPRESSION, HP_DECREASED_FACIAL_EXPRESSION
GSE1566_WT_VS_EZH2_KO_LN_TCELL_DN, GSE1566_WT_VS_EZH2_KO_LN_TCELL_DN
WP_NEURODEGENERATION_WITH_BRAIN_IRON_ACCUMULATION_NBIA_SUBTYPES_PATHWAY, WP_NEURODEGENERATION_WITH_BRAIN_IRON_ACCUMULATION_NBIA_SUBTYPES_PATHWAY
AKT_UP_MTOR_DN.V1_DN, AKT_UP_MTOR_DN.V1_DN
GOBP_POSITIVE_REGULATION_OF_CELL_CYCLE_ARREST, GOBP_POSITIVE_REGULATION_OF_CELL_CYCLE_ARREST
GSE43863_DAY6_EFF_VS_DAY150_MEM_TFH_CD4_TCELL_DN, GSE43863_DAY6_EFF_VS_DAY150_MEM_TFH_CD4_TCELL_DN
RPA1_TARGET_GENES, RPA1_TARGET_GENES
WP_AMPACTIVATED_PROTEIN_KINASE_AMPK_SIGNALING, WP_AMPACTIVATED_PROTEIN_KINASE_AMPK_SIGNALING
GOBP_FIBROBLAST_PROLIFERATION, GOBP_FIBROBLAST_PROLIFERATION
CHEN_HOXA5_TARGETS_9HR_DN, CHEN_HOXA5_TARGETS_9HR_DN
GSE360_L_MAJOR_VS_M_TUBERCULOSIS_MAC_UP, GSE360_L_MAJOR_VS_M_TUBERCULOSIS_MAC_UP
GOMF_DEACETYLASE_ACTIVITY, GOMF_DEACETYLASE_ACTIVITY
REACTOME_REGULATION_OF_TP53_ACTIVITY_THROUGH_ACETYLATION, REACTOME_REGULATION_OF_TP53_ACTIVITY_THROUGH_ACETYLATION
GOBP_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II, GOBP_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II
GOCC_PHAGOCYTIC_CUP, GOCC_PHAGOCYTIC_CUP
GOBP_ANOIKIS, GOBP_ANOIKIS
GOBP_NEGATIVE_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS
GOBP_REGULATION_OF_EXTRACELLULAR_MATRIX_ORGANIZATION, GOBP_REGULATION_OF_EXTRACELLULAR_MATRIX_ORGANIZATION
GOMF_O_ACYLTRANSFERASE_ACTIVITY, GOMF_O_ACYLTRANSFERASE_ACTIVITY
HP_MICROSCOPIC_HEMATURIA, HP_MICROSCOPIC_HEMATURIA
HOFFMANN_SMALL_PRE_BII_TO_IMMATURE_B_LYMPHOCYTE_UP, HOFFMANN_SMALL_PRE_BII_TO_IMMATURE_B_LYMPHOCYTE_UP
GOBP_POSITIVE_REGULATION_OF_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II, GOBP_POSITIVE_REGULATION_OF_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II
HP_ABNORMAL_EYELID_PHYSIOLOGY, HP_ABNORMAL_EYELID_PHYSIOLOGY
GSE45365_WT_VS_IFNAR_KO_CD8A_DC_MCMV_INFECTION_DN, GSE45365_WT_VS_IFNAR_KO_CD8A_DC_MCMV_INFECTION_DN
GOMF_PHOSPHATIDYLINOSITOL_3_5_BISPHOSPHATE_BINDING, GOMF_PHOSPHATIDYLINOSITOL_3_5_BISPHOSPHATE_BINDING
MATZUK_SPERMATOGONIA, MATZUK_SPERMATOGONIA
PID_HDAC_CLASSIII_PATHWAY, PID_HDAC_CLASSIII_PATHWAY
GOBP_POSITIVE_REGULATION_OF_GLIAL_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_GLIAL_CELL_DIFFERENTIATION
chr16q22, chr16q22
PATEL_SKIN_OF_BODY_ZOSTAVAX_AGE_70_93YO_VZV_CHALLENGE_3DY_DN, PATEL_SKIN_OF_BODY_ZOSTAVAX_AGE_70_93YO_VZV_CHALLENGE_3DY_DN
GOBP_NEGATIVE_REGULATION_OF_SMALL_MOLECULE_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_SMALL_MOLECULE_METABOLIC_PROCESS
HP_ABNORMALITY_OF_THE_DISTAL_PHALANGES_OF_THE_TOES, HP_ABNORMALITY_OF_THE_DISTAL_PHALANGES_OF_THE_TOES
MIR4732_5P, MIR4732_5P
REACTOME_RUNX2_REGULATES_BONE_DEVELOPMENT, REACTOME_RUNX2_REGULATES_BONE_DEVELOPMENT
HP_HEMIMEGALENCEPHALY, HP_HEMIMEGALENCEPHALY
HP_MUSCLE_STIFFNESS, HP_MUSCLE_STIFFNESS
GOBP_POSITIVE_REGULATION_OF_PHAGOCYTOSIS_ENGULFMENT, GOBP_POSITIVE_REGULATION_OF_PHAGOCYTOSIS_ENGULFMENT
HMX1_01, HMX1_01
GOBP_PROTEIN_HOMOTETRAMERIZATION, GOBP_PROTEIN_HOMOTETRAMERIZATION
PAX3_01, PAX3_01
HP_POLYGENIC_INHERITANCE, HP_POLYGENIC_INHERITANCE
GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_REGULATORY_REGION_DNA_BINDING, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_REGULATORY_REGION_DNA_BINDING
GOMF_METHYL_CPG_BINDING, GOMF_METHYL_CPG_BINDING
GOMF_GROWTH_FACTOR_RECEPTOR_BINDING, GOMF_GROWTH_FACTOR_RECEPTOR_BINDING
GOMF_LRR_DOMAIN_BINDING, GOMF_LRR_DOMAIN_BINDING
MIR6804_5P, MIR6804_5P
GOMF_PEPTIDE_ALPHA_N_ACETYLTRANSFERASE_ACTIVITY, GOMF_PEPTIDE_ALPHA_N_ACETYLTRANSFERASE_ACTIVITY
GOBP_N_TERMINAL_PROTEIN_AMINO_ACID_ACETYLATION, GOBP_N_TERMINAL_PROTEIN_AMINO_ACID_ACETYLATION
GOBP_AORTIC_VALVE_MORPHOGENESIS, GOBP_AORTIC_VALVE_MORPHOGENESIS
GOBP_NEGATIVE_REGULATION_OF_ATP_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_ATP_METABOLIC_PROCESS
GOBP_PHOSPHATIDYLCHOLINE_ACYL_CHAIN_REMODELING, GOBP_PHOSPHATIDYLCHOLINE_ACYL_CHAIN_REMODELING
GOBP_NEGATIVE_REGULATION_OF_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II, GOBP_NEGATIVE_REGULATION_OF_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II
MIR6865_3P, MIR6865_3P
HOLLEMAN_DAUNORUBICIN_ALL_DN, HOLLEMAN_DAUNORUBICIN_ALL_DN
GRATIAS_RETINOBLASTOMA_16Q24, GRATIAS_RETINOBLASTOMA_16Q24
GOBP_NEGATIVE_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS
HP_APLASIA_INVOLVING_BONES_OF_THE_LOWER_LIMBS, HP_APLASIA_INVOLVING_BONES_OF_THE_LOWER_LIMBS
MIR10396B_5P, MIR10396B_5P
BUSSLINGER_GASTRIC_NECK_CELLS, BUSSLINGER_GASTRIC_NECK_CELLS
GOBP_NEGATIVE_REGULATION_OF_GLYCOLYTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_GLYCOLYTIC_PROCESS
GOBP_BICARBONATE_TRANSPORT, GOBP_BICARBONATE_TRANSPORT
GOBP_THYMIC_T_CELL_SELECTION, GOBP_THYMIC_T_CELL_SELECTION
GOBP_POSITIVE_REGULATION_OF_MICROGLIAL_CELL_ACTIVATION, GOBP_POSITIVE_REGULATION_OF_MICROGLIAL_CELL_ACTIVATION
HP_VISUAL_HALLUCINATIONS, HP_VISUAL_HALLUCINATIONS
HP_EVERTED_UPPER_LIP_VERMILION, HP_EVERTED_UPPER_LIP_VERMILION
GOBP_REGULATION_OF_DNA_LIGATION, GOBP_REGULATION_OF_DNA_LIGATION
GOBP_NEGATIVE_REGULATION_OF_RELAXATION_OF_MUSCLE, GOBP_NEGATIVE_REGULATION_OF_RELAXATION_OF_MUSCLE
GOBP_REGULATION_OF_MACROPHAGE_DIFFERENTIATION, GOBP_REGULATION_OF_MACROPHAGE_DIFFERENTIATION
HP_MYOPATHIC_FACIES, HP_MYOPATHIC_FACIES
BIOCARTA_ALK_PATHWAY, BIOCARTA_ALK_PATHWAY
HP_OVERGROWTH_OF_EXTERNAL_GENITALIA, HP_OVERGROWTH_OF_EXTERNAL_GENITALIA
MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_25, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_25
HP_ACUTE_PANCREATITIS, HP_ACUTE_PANCREATITIS
GOBP_TANGENTIAL_MIGRATION_FROM_THE_SUBVENTRICULAR_ZONE_TO_THE_OLFACTORY_BULB, GOBP_TANGENTIAL_MIGRATION_FROM_THE_SUBVENTRICULAR_ZONE_TO_THE_OLFACTORY_BULB
HP_CLEFT_HARD_PALATE, HP_CLEFT_HARD_PALATE
DING_LUNG_CANCER_MUTATED_SIGNIFICANTLY, DING_LUNG_CANCER_MUTATED_SIGNIFICANTLY
GOBP_CARDIAC_VASCULAR_SMOOTH_MUSCLE_CELL_DIFFERENTIATION, GOBP_CARDIAC_VASCULAR_SMOOTH_MUSCLE_CELL_DIFFERENTIATION
GOBP_REGULATION_OF_RELAXATION_OF_MUSCLE, GOBP_REGULATION_OF_RELAXATION_OF_MUSCLE
HP_CALCINOSIS, HP_CALCINOSIS
GOBP_CARBON_CATABOLITE_REGULATION_OF_TRANSCRIPTION, GOBP_CARBON_CATABOLITE_REGULATION_OF_TRANSCRIPTION
GOBP_POSITIVE_REGULATION_OF_DNA_LIGATION, GOBP_POSITIVE_REGULATION_OF_DNA_LIGATION
FOURATI_BLOOD_TWINRIX_AGE_65_81Y0_RESPONDERS_VS_POOR_RESPONDERS_TRAINING_SET_0DY_NETWORK_INFERENCE_UP, FOURATI_BLOOD_TWINRIX_AGE_65_81Y0_RESPONDERS_VS_POOR_RESPONDERS_TRAINING_SET_0DY_NE
GOMF_SOLUTE_BICARBONATE_SYMPORTER_ACTIVITY, GOMF_SOLUTE_BICARBONATE_SYMPORTER_ACTIVITY
TSUTSUMI_FBXW8_TARGETS, TSUTSUMI_FBXW8_TARGETS
REACTOME_INTERLEUKIN_36_PATHWAY, REACTOME_INTERLEUKIN_36_PATHWAY
WP_GENES_TARGETED_BY_MIRNAS_IN_ADIPOCYTES, WP_GENES_TARGETED_BY_MIRNAS_IN_ADIPOCYTES
GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_CHEMOTAXIS, GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_CHEMOTAXIS
GOBP_CELLULAR_RESPONSE_TO_LAMINAR_FLUID_SHEAR_STRESS, GOBP_CELLULAR_RESPONSE_TO_LAMINAR_FLUID_SHEAR_STRESS
GOBP_DORSAL_AORTA_DEVELOPMENT, GOBP_DORSAL_AORTA_DEVELOPMENT
XU_HGF_TARGETS_REPRESSED_BY_AKT1_UP, XU_HGF_TARGETS_REPRESSED_BY_AKT1_UP
GOBP_REGULATION_OF_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY
SMIRNOV_CIRCULATING_ENDOTHELIOCYTES_IN_CANCER_DN, SMIRNOV_CIRCULATING_ENDOTHELIOCYTES_IN_CANCER_DN
GOBP_REGULATION_OF_TRANSCRIPTION_BY_GLUCOSE, GOBP_REGULATION_OF_TRANSCRIPTION_BY_GLUCOSE
GOMF_ALPHA_2A_ADRENERGIC_RECEPTOR_BINDING, GOMF_ALPHA_2A_ADRENERGIC_RECEPTOR_BINDING
REACTOME_RUNX2_REGULATES_CHONDROCYTE_MATURATION, REACTOME_RUNX2_REGULATES_CHONDROCYTE_MATURATION
GOBP_TACHYKININ_RECEPTOR_SIGNALING_PATHWAY, GOBP_TACHYKININ_RECEPTOR_SIGNALING_PATHWAY
GOBP_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_HEART_DEVELOPMENT, GOBP_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_HEART_D
GOBP_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_BETA_SIGNALING_PATHWAY, GOBP_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_BETA_SIGNALING_PATHWAY
REACTOME_PHOSPHATE_BOND_HYDROLYSIS_BY_NTPDASE_PROTEINS, REACTOME_PHOSPHATE_BOND_HYDROLYSIS_BY_NTPDASE_PROTEINS
GOBP_PRIMITIVE_STREAK_FORMATION, GOBP_PRIMITIVE_STREAK_FORMATION
GOMF_NEUREXIN_FAMILY_PROTEIN_BINDING, GOMF_NEUREXIN_FAMILY_PROTEIN_BINDING
REACTOME_PLASMA_LIPOPROTEIN_REMODELING, REACTOME_PLASMA_LIPOPROTEIN_REMODELING
GOBP_MACROPHAGE_COLONY_STIMULATING_FACTOR_SIGNALING_PATHWAY, GOBP_MACROPHAGE_COLONY_STIMULATING_FACTOR_SIGNALING_PATHWAY
GOBP_POSITIVE_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
HP_PRECOCIOUS_PUBERTY_IN_FEMALES, HP_PRECOCIOUS_PUBERTY_IN_FEMALES
GOBP_REGULATION_OF_THE_FORCE_OF_HEART_CONTRACTION_BY_CHEMICAL_SIGNAL, GOBP_REGULATION_OF_THE_FORCE_OF_HEART_CONTRACTION_BY_CHEMICAL_SIGNAL
GOBP_POSITIVE_REGULATION_OF_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY
GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_HEART_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INV
HP_DECREASED_CIRCULATING_COPPER_CONCENTRATION, HP_DECREASED_CIRCULATING_COPPER_CONCENTRATION
GOBP_REGULATION_OF_NON_MEMBRANE_SPANNING_PROTEIN_TYROSINE_KINASE_ACTIVITY, GOBP_REGULATION_OF_NON_MEMBRANE_SPANNING_PROTEIN_TYROSINE_KINASE_ACTIVITY
GOBP_DORSAL_AORTA_MORPHOGENESIS, GOBP_DORSAL_AORTA_MORPHOGENESIS
GOBP_ADENYLATE_CYCLASE_INHIBITING_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY, GOBP_ADENYLATE_CYCLASE_INHIBITING_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY
YAMASHITA_LIVER_CANCER_WITH_EPCAM_DN, YAMASHITA_LIVER_CANCER_WITH_EPCAM_DN

IL2RAHIGH_DAY3_EFF_CD8_TCELL_DN, GSE19825_NAIVE_VS_IL2RAHIGH_DAY3_EFF_CD8_TCELL_DN