

GSE4260_BTLL_POS_VS_NEG_INTRATUMORAL_CD4_TCELL_UP, GSE4260_BTLL_POS_VS_NEG_INTRATUMORAL_CD8_TCELL_UP
GSE2286_NAIVE_CD4_TCELL_VS_NKCELL_DN, GSE2286_NAIVE_CD4_TCELL_VS_NKCELL_DN
GSE1460_CORD_VS_ADULT_BLOOD_NAIVE_CD4_TCELL_DN, GSE1460_CORD_VS_ADULT_BLOOD_NAIVE_CD4_TCELL_DN
GSE1460_CORD_VS_ADULT_BLOOD_NAIVE_CD4_TCELL_DN, GSE1460_CORD_VS_ADULT_BLOOD_NAIVE_CD4_TCELL_DN
GSE1025_MYELOID_VS_LUPUS_MYELOID_DN, GSE1025_MYELOID_VS_LUPUS_MYELOID_DN
GSE1974_0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_DN, GSE1974_0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_DN
GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_UP, GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_UP
GSE1460_CD4_THYMOCYTE_VS_THYMIC_STROMAL_CELL_UP, GSE1460_CD4_THYMOCYTE_VS_THYMIC_STROMAL_CELL_UP
GSE2203_UNTREATED_VS_TREATED_MIFEN_DN, GSE2203_UNTREATED_VS_TREATED_MIFEN_DN
GSE26495_NAIVE_VS_PD1HIGH_CD8_TCELL_DN, GSE26495_NAIVE_VS_PD1HIGH_CD8_TCELL_DN
ZHENG_CORD_BLOOD_C9_GNANULOCYTE_MACROPHAGE_PROGENITOR, ZHENG_CORD_BLOOD_C9_GNANULOCYTE_MACROPHAGE_PROGENITOR
GSE1460_INTRATHYMIC_T_PROGENITOR_VS_CD4_THYMOCYTE_DN, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_CD4_THYMOCYTE_DN
GSE3982_CTRL_VS_LPS_48H_DC_DN, GSE3982_CTRL_VS_LPS_48H_DC_DN
GSE460_1_DONOVAN_VS_M_TUBERCULOSIS_MAC_UP, GSE460_1_DONOVAN_VS_M_TUBERCULOSIS_MAC_UP
GSE2103_CTRL_VS_POLYIC_STIM_DC_4H_UP, GSE2103_CTRL_VS_POLYIC_STIM_DC_4H_UP
GOBP_REGULATION_OF_ORGANELLE_ASSEMBLY, GOBP_REGULATION_OF_ORGANELLE_ASSEMBLY
GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_UP, GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_UP
GOBP_RESPONSE_TO_IONIZING_RADIATION, GOBP_RESPONSE_TO_IONIZING_RADIATION
GSE20715_WT_VS_IL18_KO_LUNG_UP, GSE20715_WT_VS_IL18_KO_LUNG_UP
GSE37465_C57BL6_VS_NOD_FOXP3_FUSION_GFP_TREG_DN, GSE37465_C57BL6_VS_NOD_FOXP3_FUSION_GFP_TREG_DN
WP_TGFBETA_SIGNALING_PATHWAY, WP_TGFBETA_SIGNALING_PATHWAY
GSE20152_SPHK1_KO_VS_WT_HTNFA_OVERXPRESS_ANKLE_UP, GSE20152_SPHK1_KO_VS_WT_HTNFA_OVERXPRESS_ANKLE_UP
GSE1460_DP_VS_CD4_THYMOCYTE_DN, GSE1460_DP_VS_CD4_THYMOCYTE_DN
MIR379_3P, MIR411_3P, MIR379_3P, MIR411_3P
GCM1_RBM4A, GCM1_RBM4A
GSE17721_CTRL_VS_PAMC5K4_12H_BMDC_DN, GSE17721_CTRL_VS_PAMC5K4_12H_BMDC_DN
GSE17721_CTRL_VS_POLYIC_1H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_1H_BMDC_DN
IZADPANAH_STEM_CELL_ADIPOSE_VS_BONE_UP, IZADPANAH_STEM_CELL_ADIPOSE_VS_BONE_UP
MIR5697, MIR5697
MIR5530, MIR5530
GSE37201_HEMATOPOIETIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_UP, GSE37201_HEMATOPOIETIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_UP
REACTOME_TOLL_LIKE_RECEPTOR_9_TLR9_CASCADE, REACTOME_TOLL_LIKE_RECEPTOR_9_TLR9_CASCADE
DE_YYL_TARGETS_DN, DE_YYL_TARGETS_DN
HP_HEPATOSPLENOMEGALY, HP_HEPATOSPLENOMEGALY
GSE40274_CTRL_VS_FOXP3_AND_IRF4_TRANSUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_CTRL_VS_FOXP3_AND_IRF4_TRANSUCED_ACTIVATED_CD4_TCELL_UP
GOBP_DNA_DEPENDENT_ATPASE_ACTIVITY, GOBP_DNA_DEPENDENT_ATPASE_ACTIVITY
ZNFS84_TARGET_GENES, ZNFS84_TARGET_GENES
SENES_HDAC2_TARGETS_UP, SENES_HDAC2_TARGETS_UP
PAX8_TARGET_GENES, PAX8_TARGET_GENES
GOBP_REGULATION_OF_ADAPTIVE_IMMUNE_RESPONSE, GOBP_REGULATION_OF_ADAPTIVE_IMMUNE_RESPONSE
GSE2826_WT_VS_XID_BCELL_DN, GSE2826_WT_VS_XID_BCELL_DN
WP_ECTODERM_DIFFERENTIATION, WP_ECTODERM_DIFFERENTIATION
GOBP_PROTEIN_K48_LINKED_UBIQUITINATION, GOBP_PROTEIN_K48_LINKED_UBIQUITINATION
WP_TNF_ALPHA_SIGNALING_PATHWAY, WP_TNF_ALPHA_SIGNALING_PATHWAY
MIR628_3P, MIR628_3P
GSE26783_ANTL_MIR33_VS_CTRL_ATHEROSCLEROSIS_MACROPHAGE_DN, GSE26783_ANTL_MIR33_VS_CTRL_ATHEROSCLEROSIS_MACROPHAGE_DN
CGGAARNGCGNG, UNKNOWN, CGGAARNGCGNG, UNKNOWN
KEGG_COLORECTAL_CANCER, KEGG_COLORECTAL_CANCER
HP_ABNORMAL_THORACIC_SPINE_MORPHOLOGY, HP_ABNORMAL_THORACIC_SPINE_MORPHOLOGY
MIR8082, MIR8082
HP_XEROSTOMIA_HP_XEROSTOMIA
MODULE_427, MODULE_427
BROWNE_HCMV_INFECTION_12HR_DN, BROWNE_HCMV_INFECTION_12HR_DN
GALE_APL_WITH_FLT3_MUTATED_UP, GALE_APL_WITH_FLT3_MUTATED_UP
HP_ABNORMAL_CIRCULATING_PROTEINOGENIC_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_PROTEINOGENIC_AMINO_ACID_CONCENTRATION
HP_ABNORMAL_CIRCULATING_CHOLESTEROL_CONCENTRATION, HP_ABNORMAL_CIRCULATING_CHOLESTEROL_CONCENTRATION
PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_4, PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_4
REACTOME_INTERLEUKIN_17_SIGNALING, REACTOME_INTERLEUKIN_17_SIGNALING
MIR6838_3P, MIR6838_3P
GOBP_POSITIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION
TRAVAGLINI_LUNG_VASCULAR_SMOOTH_MUSCLE_CELL, TRAVAGLINI_LUNG_VASCULAR_SMOOTH_MUSCLE_CELL
HP_AMYOTROPHIC_LATERAL_SCLEROSIS, HP_AMYOTROPHIC_LATERAL_SCLEROSIS
AATGCG, MIR136, AATGCG, MIR136
HP_FATIGABLE_WEAKNESS_OF_BULBAR_MUSCLES, HP_FATIGABLE_WEAKNESS_OF_BULBAR_MUSCLES
GOCC_ACTOMYOSIN, GOCC_ACTOMYOSIN
MAPK3_TARGET_GENES, MAPK3_TARGET_GENES
GOBP_FIBROBLAST_PROLIFERATION, GOBP_FIBROBLAST_PROLIFERATION
CHEN_LIVER_METABOLISM_QTL_CS, CHEN_LIVER_METABOLISM_QTL_CS
WP_REGULATION_OF_ACTIN_CYTOSKELETON, WP_REGULATION_OF_ACTIN_CYTOSKELETON
GOCC_ACTIN_FILAMENT_BUNDLE, GOCC_ACTIN_FILAMENT_BUNDLE
HERNANDEZ_ABERRANT_MITOSIS_BY_DOCETACEL_2NM_UP, HERNANDEZ_ABERRANT_MITOSIS_BY_DOCETACEL_2NM_UP
MIR179, MIR3179
GOBP_CORTICAL_CYTOSKELETON_ORGANIZATION, GOBP_CORTICAL_CYTOSKELETON_ORGANIZATION
GOBP_POSITIVE_REGULATION_OF_ORGANELLE_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_ORGANELLE_ASSEMBLY
KEGG_VEGF_SIGNALING_PATHWAY, KEGG_VEGF_SIGNALING_PATHWAY
GOBP_POSITIVE_REGULATION_OF_ADAPTIVE_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_ADAPTIVE_IMMUNE_RESPONSE
MIR586_3P, MIR586_3P
HP_DEMYELINATING_PERIPHERAL_NEUROPATHY, HP_DEMYELINATING_PERIPHERAL_NEUROPATHY
PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_1, PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_1
REACTOME_NUCLEAR_EVENTS_KINASE_AND_TRANSCRIPTION_FACTOR_ACTIVATION, REACTOME_NUCLEAR_EVENTS_KINASE_AND_TRANSCRIPTION_FACTOR_ACTIVATION
CCATCCA_MIR432, CCATCCA_MIR432
DESCARTES_FETAL_EYE_VASCULAR_ENDOTHELIAL_CELLS, DESCARTES_FETAL_EYE_VASCULAR_ENDOTHELIAL_CELLS
HP_FATIGABLE_WEAKNESS_OF_RESPIRATORY_MUSCLES, HP_FATIGABLE_WEAKNESS_OF_RESPIRATORY_MUSCLES
GOBP_RNA_DESTABILIZATION, GOBP_RNA_DESTABILIZATION
HP_MULTIPLE_JOINT_CONTRACTURES, HP_MULTIPLE_JOINT_CONTRACTURES
LUL_THYROID_CANCER_CLUSTER_1, LUL_THYROID_CANCER_CLUSTER_1
MENON_FETAL_KIDNEY_3_STROMAL_CELLS, MENON_FETAL_KIDNEY_3_STROMAL_CELLS
CTAACAC_MIR2957, CTAACAC_MIR2957
GSE4606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_JRF4MID_BCELL_UP, GSE4606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_JRF4MID_BCELL_UP
PID_RHOA_PATHWAY, PID_RHOA_PATHWAY
KEGG_FC_EPSILON_RL_SIGNALING_PATHWAY, KEGG_FC_EPSILON_RL_SIGNALING_PATHWAY
GOBP_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS, GOBP_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS
HP_ABNORMAL_OVARIAN_PHYSIOLOGY, HP_ABNORMAL_OVARIAN_PHYSIOLOGY
GOBP_MITOGEN_ACTIVATED_PROTEIN_KINASE_BINDING, GOBP_MITOGEN_ACTIVATED_PROTEIN_KINASE_BINDING
MIR3654, MIR3654
GCAAGAC_MIR431, GCAAGAC_MIR431
MIR6826_5P, MIR6826_5P
GSE22025_UNTREATED_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_DN, GSE22025_UNTREATED_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_DN
EPERTT_CE_HSC_LSC, EPERTT_CE_HSC_LSC
GENTILE_UV_RESPONSE_CLUSTER_D5, GENTILE_UV_RESPONSE_CLUSTER_D5
GOBP_REGULATION_OF_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GOBP_REGULATION_OF_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GOBP_NEGATIVE_REGULATION_OF_RNA_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_RNA_CATABOLIC_PROCESS
HP_FAILURE_TO_THRIVE_IN_INFANCY, HP_FAILURE_TO_THRIVE_IN_INFANCY
MIR446_3P, MIR446_3P
MIR3144_3P, MIR3144_3P
KHETCHOUMIAN_TRIM24_TARGETS_UP, KHETCHOUMIAN_TRIM24_TARGETS_UP
NATSUME_RESPONSE_TO_INTERFERON_BETA_UP, NATSUME_RESPONSE_TO_INTERFERON_BETA_UP
GSE4887_WT_VS_GFI1_KO_PDC_UP, GSE4887_WT_VS_GFI1_KO_PDC_UP
GOCC_NUCLEAR_INNER_MEMBRANE, GOCC_NUCLEAR_INNER_MEMBRANE
GTCCTACC_MIR379, GTCCTACC_MIR379
HP_NORMOCYTIC_ANEMIA, HP_NORMOCYTIC_ANEMIA
GOCC_CUL2_RING_UBIQUITIN_LIGASE_COMPLEX, GOCC_CUL2_RING_UBIQUITIN_LIGASE_COMPLEX
CHEBOTAYE_GR_TARGETS_UP, CHEBOTAYE_GR_TARGETS_UP
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS, BLUE_DN
TENEDINI_MEGAKARYOCYTE_MARKERS, TENEDINI_MEGAKARYOCYTE_MARKERS
JOSEPH_RESPONSE_TO_SODIUM_BUTYRATE_DN, JOSEPH_RESPONSE_TO_SODIUM_BUTYRATE_DN
GOBP_B_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_B_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE
SAMOES_TARGETS_OF_KHSV_MIRNAS_DN, SAMOES_TARGETS_OF_KHSV_MIRNAS_DN
GOBP_RESPONSE_TO_X_RAY, GOBP_RESPONSE_TO_X_RAY
GRABARCZYK_BCL11B_TARGETS_DN, GRABARCZYK_BCL11B_TARGETS_DN
METTL14_TARGET_GENES, METTL14_TARGET_GENES
WP_CARDIAC_HYPERTROPIC_RESPONSE, WP_CARDIAC_HYPERTROPIC_RESPONSE
MIR435_3P, MIR435_3P
MIR7945_3P, MIR7945_3P
HP_SHORT_CHIN, HP_SHORT_CHIN
HOLLEMAN_VINCISTRINE_RESISTANCE_B_ALL_UP, HOLLEMAN_VINCISTRINE_RESISTANCE_B_ALL_UP
HP_ABNORMALITY_OF_PRIMARY_TEETH, HP_ABNORMALITY_OF_PRIMARY_TEETH
HP_CARDIOVASCULAR_CALCIFICATION, HP_CARDIOVASCULAR_CALCIFICATION
GOBP_HEAT_GENERATION, GOBP_HEAT_GENERATION
GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN, GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN
GOME_SCAFFOLD_PROTEIN_BINDING, GOME_SCAFFOLD_PROTEIN_BINDING
GOBP_PRL_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II, GOBP_PRL_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II
MIR3115, MIR3115
GOBP_POSITIVE_REGULATION_OF_FIBROBLAST_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_FIBROBLAST_PROLIFERATION
GOBP_CALCINEURIN_MEDIATED_SIGNALING, GOBP_CALCINEURIN_MEDIATED_SIGNALING
MIR449C_3P, MIR449C_3P
HP_HYPERCHOLESTEROLEMIA, HP_HYPERCHOLESTEROLEMIA
GOBP_STRIATED_MUSCLE_CELL_DEVELOPMENT, GOBP_STRIATED_MUSCLE_CELL_DEVELOPMENT
GOBP_COLLAGEN_BIOSYNTHETIC_PROCESS, GOBP_COLLAGEN_BIOSYNTHETIC_PROCESS
REACTOME_MAPK_TARGETS_NUCLEAR_EVENTS_MEDIATED_BY_MAP_KINASES, REACTOME_MAPK_TARGETS_NUCLEAR_EVENTS_MEDIATED_BY_MAP_KINASES
GOBP_AMINOGLYCAN_CATABOLIC_PROCESS, GOBP_AMINOGLYCAN_CATABOLIC_PROCESS
VANTVEER_BREAST_CANCER_BRCA1_UP, VANTVEER_BREAST_CANCER_BRCA1_UP
GOBP_POSITIVE_REGULATION_OF_GLIONECROSIS, GOBP_POSITIVE_REGULATION_OF_GLIONECROSIS
MIR6_3P, MIR6_3P
WP_GENES_INVOLVED_IN_MALE_INFERTILITY, WP_GENES_INVOLVED_IN_MALE_INFERTILITY
MATTHEWS_API_TARGETS, MATTHEWS_API_TARGETS
REACTOME_NGF_STIMULATED_TRANSCRIPTION, REACTOME_NGF_STIMULATED_TRANSCRIPTION
GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT, GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT
HP_ABNORMALITIES_OF_PLACENTA_OR_UMBILICAL_CORD, HP_ABNORMALITIES_OF_PLACENTA_OR_UMBILICAL_CORD
GOBP_MUSCLE_CELL_DEVELOPMENT, GOBP_MUSCLE_CELL_DEVELOPMENT
LIU_BREAST_CANCER_LIU_BREAST_CANCER
GAO_STOMACH_24W_C5_PUTATIVE_PIT_CELL_PROGENITOR, GAO_STOMACH_24W_C5_PUTATIVE_PIT_CELL_PROGENITOR
GSE360_L1_MAJOR_VS_T_GONDILODC_UP, GSE360_L1_MAJOR_VS_T_GONDILODC_UP
WP_HOPISTATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_AUTOPHAGY, WP_HOPISTATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_AUTOPHAGY
GOBP_POSITIVE_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS
GOBP_REGULATION_OF_BEHAVIOR, GOBP_REGULATION_OF_BEHAVIOR
GOBP_EXOCRINE_SYSTEM_DEVELOPMENT, GOBP_EXOCRINE_SYSTEM_DEVELOPMENT
REACTOME_UPTAKE_AND_ACTIONS_OF_BACTERIAL_TOXINS, REACTOME_UPTAKE_AND_ACTIONS_OF_BACTERIAL_TOXINS
WP_ALLOGRAFT_REJECTION, WP_ALLOGRAFT_REJECTION
MARCHINI TRABECTEDIN_RESISTANCE_DN, MARCHINI TRABECTEDIN_RESISTANCE_DN
HOXD1_TARGET_GENES, HOXD1_TARGET_GENES
HERNANDEZ_ABERRANT_MITOSIS_BY_DOCETACEL_2NM_DN, HERNANDEZ_ABERRANT_MITOSIS_BY_DOCETACEL_2NM_DN
HP_ASTROCYTOSIS, HP_ASTROCYTOSIS
BCAT_G05748_UP, BCAT_G05748_UP
HP_DECREASED_SKULL_OSSIFICATION, HP_DECREASED_SKULL_OSSIFICATION
HP_IRREGULAR_MENSTRUATION, HP_IRREGULAR_MENSTRUATION
GOBP_RESPONSE_TO_CORTICOSTEROID, GOBP_RESPONSE_TO_CORTICOSTEROID
GOBP_CELLULAR_RESPONSE_TO_DEXAMETHASONE_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_DEXAMETHASONE_STIMULUS
HP_YOUNG_ADULT_ONSET, HP_YOUNG_ADULT_ONSET
AGCGCAG, MIR91, AGCGCAG, MIR91
HP_PAIN_INSENSITIVITY, HP_PAIN_INSENSITIVITY
GOBP_HETEROTYPIC_CELL_CELL_ADHESION, GOBP_HETEROTYPIC_CELL_CELL_ADHESION
BROWNE_HCMV_INFECTION_4HR_UP, BROWNE_HCMV_INFECTION_4HR_UP
GOBP_IMMUNOGLOBULIN_PRODUCTION_INVOLVED_IN_IMMUNOGLOBULIN_MEDIATED_IMMUNE_RESPONSE, GOBP_IMMUNOGLOBULIN_PRODUCTION_INVOLVED_IN_IMMUNOGLOBULIN_MEDIATED_IMMUNE_RESPONSE
KAMIKUBO_MYELOID_MN_NETWORK, KAMIKUBO_MYELOID_MN_NETWORK
LI_PROSTATE_CANCER_EPIGENETIC, LI_PROSTATE_CANCER_EPIGENETIC
BCAT100_UP_VI_UP, BCAT100_UP_VI_UP
GOBP_POSITIVE_REGULATION_OF_T_CELL_MEDIATED_IMMUNITY, GOBP_POSITIVE_REGULATION_OF_T_CELL_MEDIATED_IMMUNITY
GOBP_2_OXOGUTARATE_METABOLIC_PROCESS, GOBP_2_OXOGUTARATE_METABOLIC_PROCESS
GOBP_TOLL_LIKE_RECEPTOR_9_SIGNALING_PATHWAY, GOBP_TOLL_LIKE_RECEPTOR_9_SIGNALING_PATHWAY
GOBP_NEUROINFLAMMATORY_RESPONSE, GOBP_NEUROINFLAMMATORY_RESPONSE
DUNNE_TARGETS_OF_AML1_MTG8_FUSION_DN, DUNNE_TARGETS_OF_AML1_MTG8_FUSION_DN
GOBP_TRANSEPTHELIAL_TRANSPORT, GOBP_TRANSEPTHELIAL_TRANSPORT
GOBP_REGULATION_OF_HEAT_GENERATION, GOBP_REGULATION_OF_HEAT_GENERATION
MIR219A_2_3P, MIR219A_2_3P
LIEN_BREAST_CARCINOMA_METAPLASTIC, LIEN_BREAST_CARCINOMA_METAPLASTIC
SA_MMP_CYTOKINE_CONNECTION, SA_MMP_CYTOKINE_CONNECTION
GOBP_SALIVARY_GLAND_DEVELOPMENT, GOBP_SALIVARY_GLAND_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_HEAT_GENERATION, GOBP_POSITIVE_REGULATION_OF_HEAT_GENERATION
WP_UREA_CYCLE_AND_ASSOCIATED_PATHWAYS, WP_UREA_CYCLE_AND_ASSOCIATED_PATHWAYS
GOBP_RESPONSE_TO_ARSENIC_CONTAINING_SUBSTANCE, GOBP_RESPONSE_TO_ARSENIC_CONTAINING_SUBSTANCE
HP_BROAD_FACE, HP_BROAD_FACE
GOBP_POSITIVE_REGULATION_OF_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY
HP_CONGENITAL_SENSORINEURAL_HEARING_IMPAIRMENT, HP_CONGENITAL_SENSORINEURAL_HEARING_IMPAIRMENT
HP_LARGE_BEAKED_NOSE, HP_LARGE_BEAKED_NOSE
HP_ABNORMAL_PELVIS_BONE_MORPHOLOGY, HP_ABNORMAL_PELVIS_BONE_MORPHOLOGY
HP_KAYNAUD_PHENOMENON, HP_KAYNAUD_PHENOMENON
GOBP_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION, GOBP_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION
MAGRANGEAS_MULTIPLE_MYELOMA_JGG_VS_JGA_DN, MAGRANGEAS_MULTIPLE_MYELOMA_JGG_VS_JGA_DN
LIU_TUMOR_ANGIOGENESIS_UP, LIU_TUMOR_ANGIOGENESIS_UP
WEINMANN_ADAPTATION_TO_HYPOXIA_DN, WEINMANN_ADAPTATION_TO_HYPOXIA_DN
GOBP_VENTRICULAR_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION, GOBP_VENTRICULAR_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION
TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_MONOCYTE_DN, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_MONOCYTE_DN
REACTOME_ERK_MAPK_TARGETS, REACTOME_ERK_MAPK_TARGETS
GOBP_POSITIVE_REGULATION_OF_HORMONE_SECRETION, GOBP_POSITIVE_REGULATION_OF_HORMONE_SECRETION
HP_SKIN_PLAQUE, HP_SKIN_PLAQUE
PID_HDAC_CLASSIII_PATHWAY, PID_HDAC_CLASSIII_PATHWAY
GOBP_REGULATION_OF_ISOTYPE_SWITCHING, GOBP_REGULATION_OF_ISOTYPE_SWITCHING
GOBP_SEQUESTERING_OF_TRIGLYCERIDE, GOBP_SEQUESTERING_OF_TRIGLYCERIDE
GOBP_NEGATIVE_REGULATION_OF_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GOBP_NEGATIVE_REGULATION_OF_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GOBP_POSITIVE_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GOBP_POSITIVE_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS
HP_SEVERE_FAILURE_TO_THRIVE, HP_SEVERE_FAILURE_TO_THRIVE
GOBP_INOSITOL_PHOSPHATE_MEDIATING_SIGNALING, GOBP_INOSITOL_PHOSPHATE_MEDIATING_SIGNALING
GOBP_SMOOTH_MUSCLE_CONTRACTION, GOBP_SMOOTH_MUSCLE_CONTRACTION
GOBP_REGULATION_OF_MEMBRANE_REPOLARIZATION, GOBP_REGULATION_OF_MEMBRANE_REPOLARIZATION
GOBP_DNA_LIGATION, GOBP_DNA_LIGATION
GOBP_REGULATION_OF_PROTEIN_FOLDING, GOBP_REGULATION_OF_PROTEIN_FOLDING
GUTIERREZ_WALDENSTROMS_MACROGLOBULINEMIA_2, GUTIERREZ_WALDENSTROMS_MACROGLOBULINEMIA_2
HP_THORACIC_KYPHOSCOLIOSIS, HP_THORACIC_KYPHOSCOLIOSIS
GOBP_CARDIAC_CELL_DEVELOPMENT, GOBP_CARDIAC_CELL_DEVELOPMENT
HP_FEMALE_HYPOGONADISM, HP_FEMALE_HYPOGONADISM
GOBP_REGULATION_OF_WATER_LOSS_VIA_SKIN, GOBP_REGULATION_OF_WATER_LOSS_VIA_SKIN
WP_STRIATED_MUSCLE_CONTRACTION_PATHWAY, WP_STRIATED_MUSCLE_CONTRACTION_PATHWAY
HU_FETAL_RETINA_AMACRINE, HU_FETAL_RETINA_AMACRINE
MIR6753_5P, MIR6753_5P
GSE1347_WT_VS_ZFX_KO_BCELL_ANTILGCM_STIM_12H_UP, GSE1347_WT_VS_ZFX_KO_BCELL_ANTILGCM_STIM_12H_UP
HP_REDUCED_SUBCUTANEOUS_ADIPOSE_TISSUE, HP_REDUCED_SUBCUTANEOUS_ADIPOSE_TISSUE
BIOCARTA_IL12_PATHWAY, BIOCARTA_IL12_PATHWAY
HP_BLUE_RIDES, HP_BLUE_RIDES
GOBP_MYTUBE_CELL_DEVELOPMENT, GOBP_MYTUBE_CELL_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_BLOOD_PRESSURE, GOBP_POSITIVE_REGULATION_OF_BLOOD_PRESSURE
GOBP_SINGLE_STRAND_BREAK_REPAIR, GOBP_SINGLE_STRAND_BREAK_REPAIR
GOBP_NEGATIVE_REGULATION_OF_NEURONINFLAMMATORY_RESPONSE, GOBP_NEGATIVE_REGULATION_OF_NEURONINFLAMMATORY_RESPONSE
GOBP_BRANCHING_INVOLVED_IN_SALIVARY_GLAND_MORPHOGENESIS, GOBP_BRANCHING_INVOLVED_IN_SALIVARY_GLAND_MORPHOGENESIS
GOBP_REGULATION_OF_VENTRICULAR_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION, GOBP_REGULATION_OF_VENTRICULAR_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION
GOBP_CARDIAC_MUSCLE_FIBER_DEVELOPMENT, GOBP_CARDIAC_MUSCLE_FIBER_DEVELOPMENT
LEE_LIVER_CANCER_HEPATOBLAST, LEE_LIVER_CANCER_HEPATOBLAST
MODULE_375, MODULE_375
GOBP_REGULATION_OF_PODOSOME_ASSEMBLY, GOBP_REGULATION_OF_PODOSOME_ASSEMBLY
HP_IRREGULAR_VERTEBRAL_ENDPLATES, HP_IRREGULAR_VERTEBRAL_ENDPLATES
PID_P8_GAMMA_DELTA_PATHWAY, PID_P8_GAMMA_DELTA_PATHWAY
KM_ALL_DISORDERS_OILCODENDROCYTE_NUMBER_CORR_DN, KM_ALL_DISORDERS_OILCODENDROCYTE_NUMBER_CORR_DN
REACTOME_SIGNAL_AMPLIFICATION, REACTOME_SIGNAL_AMPLIFICATION
MIR1298_3P, MIR1298_3P
GOBP_REGULATION_OF_BLOOD_PRESSURE, GOBP_REGULATION_OF_BLOOD_PRESSURE
GOBP_PEPITIDYL_LYSINE_DEACETYLATION, GOBP_PEPITIDYL_LYSINE_DEACETYLATION
GOBP_POSITIVE_REGULATION_OF_TRANSFORMING_GROWTH_FACTOR_BETA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_TRANSFORMING_GROWTH_FACTOR_BETA_PRODUCTION
DEMAGALHAES_AGING_DN, DEMAGALHAES_AGING_DN
GOBP_GANGLIOSIDE_METABOLIC_PROCESS, GOBP_GANGLIOSIDE_METABOLIC_PROCESS
PID_ANTHRAX_PATHWAY, PID_ANTHRAX_PATHWAY
GOBP_SEMI_LUNAR_VALVE_DEVELOPMENT, GOBP_SEMI_LUNAR_VALVE_DEVELOPMENT
XU_RESPONSE_TO_TREITONIN_DN, XU_RESPONSE_TO_TREITONIN_DN
GOCC_CUL4A_RING_E3_UBIQUITIN_LIGASE_COMPLEX, GOCC_CUL4A_RING_E3_UBIQUITIN_LIGASE_COMPLEX
WP_HAIR_FOLLICLE_DEVELOPMENT_ORGANOGENESIS_PART_2_OF_3, WP_HAIR_FOLLICLE_DEVELOPMENT_ORGANOGENESIS_PART_2_OF_3
GOBP_REGULATION_OF_EXECUTION_PHASE_OF_APOPTOSIS, GOBP_REGULATION_OF_EXECUTION_PHASE_OF_APOPTOSIS
GOBP_PULMONARY_VALVE_DEVELOPMENT, GOBP_PULMONARY_VALVE_DEVELOPMENT
GOBP_NEGATIVE_REGULATION_OF_LIPID_STORAGE, GOBP_NEGATIVE_REGULATION_OF_LIPID_STORAGE
HP_ABNORMAL_CIRCULATING_GLUTAMINE_FAMILY_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_GLUTAMINE_FAMILY_AMINO_ACID_CONCENTRATION
WANG_LSD1_TARGETS_UP, WANG_LSD1_TARGETS_UP
HP_CHRONIC_ORAL_CANDIDIASIS, HP_CHRONIC_ORAL_CANDIDIASIS
GOBP_HYALURONAN_CATABOLIC_PROCESS, GOBP_HYALURONAN_CATABOLIC_PROCESS
REACTOME_ADP_SIGNALING_THROUGH_P2Y_PURINOCEPTOR_1, REACTOME_ADP_SIGNALING_THROUGH_P2Y_PURINOCEPTOR_1
GOBP_ASPARTATE_FAMILY_AMINO_ACID_CATABOLIC_PROCESS, GOBP_ASPARTATE_FAMILY_AMINO_ACID_CATABOLIC_PROCESS
GOBP_CELLULAR_RESPONSE_TO_MURAMYLDIPEPTIDE, GOBP_CELLULAR_RESPONSE_TO_MURAMYLDIPEPTIDE
GOBP_POSITIVE_REGULATION_OF_NERVOUS_SYSTEM_PROCESS, GOBP_POSITIVE_REGULATION_OF_NERVOUS_SYSTEM_PROCESS
GOBP_SPHINGOMYELIN_BIOSYNTHETIC_PROCESS, GOBP_SPHINGOMYELIN_BIOSYNTHETIC_PROCESS
HP_ANKYLOGLOSSIA, HP_ANKYLOGLOSSIA
GOBP_IMMUNE_RESPONSE_TO_TUMOR_CELL, GOBP_IMMUNE_RESPONSE_TO_TUMOR_CELL
MASRL_RESISTANCE_TO_TAMOXIFEN_AND_AROMATASE_INHIBITORS_DN, MASRL_RESISTANCE_TO_TAMOXIFEN_AND_AROMATASE_INHIBITORS_DN
MODULE_289, MODULE_289
GOBP_NEGATIVE_REGULATION_OF_CELL_JUNCTION_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_CELL_JUNCTION_ASSEMBLY
LEE_SPL_THYMOCYTE, LEE_SPL_THYMOCYTE
GOBP_REGULATION_OF_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION, GOBP_REGULATION_OF_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION
GOBP_SARCOMERE_ORGANIZATION, GOBP_SARCOMERE_ORGANIZATION
WP_CONSTITUTIVE_ANDROSTANE_RECEPTOR_PATHWAY, WP_CONSTITUTIVE_ANDROSTANE_RECEPTOR_PATHWAY
HP_WIDE_CRANIAL_SUTURES, HP_WIDE_CRANIAL_SUTURES
MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_17, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_17
HP_ABNORMALITY_OF_SECONDARY_SEXUAL_HAIR, HP_ABNORMALITY_OF_SECONDARY_SEXUAL_HAIR
GOME_TRNA_SPECIFIC_ADENOSINE_DEAMINASE_ACTIVITY, GOME_TRNA_SPECIFIC_ADENOSINE_DEAMINASE_ACTIVITY
GOBP_ASTROCYTE_DEVELOPMENT, GOBP_ASTROCYTE_DEVELOPMENT
GOBP_SMOOTH_MUSCLE_TISSUE_DEVELOPMENT, GOBP_SMOOTH_MUSCLE_TISSUE_DEVELOPMENT
GOBP_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION, GOBP_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION
SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOXES_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOXES_UP
REACTOME_SYNTHESIS_OF_PIP3_AT_THE_GOLGI_MEMBRANE, REACTOME_SYNTHESIS_OF_PIP3_AT_THE_GOLGI_MEMBRANE
HP_CENTRAL_APNIA, HP_CENTRAL_APNIA
HP_HYPOPLASIA_OF_THE_UTERUS, HP_HYPOPLASIA_OF_THE_UTERUS
HP_APLASIA_HYPOPLASIA_OF_THE_COLON, HP_APLASIA_HYPOPLASIA_OF_THE_COLON
GOBP_CAMP_CATABOLIC_PROCESS, GOBP_CAMP_CATABOLIC_PROCESS
HP_COMPENSATED_HYPOTHYROIDISM, HP_COMPENSATED_HYPOTHYROIDISM
FAN_EMBRYONIC_CTX_EX_1_EXCITATORY_NEURON, FAN_EMBRYONIC_CTX_EX_1_EXCITATORY_NEURON
GOCC_CYTOSKELETON_OF_PRESYNAPTIC_ACTIVE_ZONE, GOCC_CYTOSKELETON_OF_PRESYNAPTIC_ACTIVE_ZONE
HP_MITRAL_STENOSIS, HP_MITRAL_STENOSIS
DESCARTES_FETAL_THYMUS_THYMOCYTES, DESCARTES_FETAL_THYMUS_THYMOCYTES
GOBP_GASTRO_INTESTINAL_SYSTEM_SMOOTH_MUSCLE_CONTRACTION, GOBP_GASTRO_INTESTINAL_SYSTEM_SMOOTH_MUSCLE_CONTRACTION
GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY, GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY
GOBP_REGULATION_OF_CENTROMERE_COMPLEX_ASSEMBLY, GOBP_REGULATION_OF_CENTROMERE_COMPLEX_ASSEMBLY
WP_PREGNANE_X_RECEPTOR_PATHWAY, WP_PREGNANE_X_RECEPTOR_PATHWAY
HP_ABNORMALITY_OF_THE_VERTEBRAL_ENDPLATES, HP_ABNORMALITY_OF_THE_VERTEBRAL_ENDPLATES
GOBP_POSITIVE_REGULATION_OF_BLOOD_CIRCULATION, GOBP_POSITIVE_REGULATION_OF_BLOOD_CIRCULATION
WP_GANGLIO_SPHINGOLIPID_METABOLISM, WP_GANGLIO_SPHINGOLIPID_METABOLISM
GOBP_NOREPINEPHRINE_METABOLIC_PROCESS, GOBP_NOREPINEPHRINE_METABOLIC_PROCESS
GOBP_RESPONSE_TO_SALT, GOBP_RESPONSE_TO_SALT
GAUSSMANN_MLL_AH_FUSION_TARGETS_C_DN, GAUSSMANN_MLL_AH_FUSION_TARGETS_C_DN
GOBP_RESPONSE_TO_MUSCLE_INACTIVITY, GOBP_RESPONSE_TO_MUSCLE_INACTIVITY
GOBP_POSITIVE_REGULATION_OF_T_CELL_MEDIATED_CYTOTOXICITY, GOBP_POSITIVE_REGULATION_OF_T_CELL_MEDIATED_CYTOTOXICITY
GOBP_NEGATIVE_REGULATION_OF_FEEDING_BEHAVIOR, GOBP_NEGATIVE_REGULATION_OF_FEEDING_BEHAVIOR
HP_AORTIC_VALVE_CALCIFICATION, HP_AORTIC_VALVE_CALCIFICATION
GOBP_GASTRIC_MOTILITY, GOBP_GASTRIC_MOTILITY
HP_ELEVATED_GAMMA_GLUTAMYLTRANSFERASE_LEVEL, HP_ELEVATED_GAMMA_GLUTAMYLTRANSFERASE_LEVEL

NAIVE_CD4_TCELL_CORD_BLOOD_UP, GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_UP