

NEG_INTRATUMORAL_CD8_TCELL_UP, GSE43260_BTLA_POS_VS_NEG_INTRATUMORAL_CD8_TCELL_UP

GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_UP, GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_UP
GSE41176_WT_VS_TAK1_KO_ANT1JGM_STIM_BCELL_24H_UP, GSE41176_WT_VS_TAK1_KO_ANT1JGM_STIM_BCELL_24H_UP
GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_UP
GSE19401_PAM2CSK4_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN, GSE19401_PAM2CSK4_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN
GSE22935_WT_VS_MYD88_KO_MACROPHAGE_48H_MBOVIS_BCG_STIM_DN, GSE22935_WT_VS_MYD88_KO_MACROPHAGE_48H_MBOVIS_BCG_STIM_DN
GSE17721_POLYIC_VS_CPG_6H_BMDC_DN, GSE17721_POLYIC_VS_CPG_6H_BMDC_DN
GSE369_PRE_VS_POST_IL6_INJECTION_SOCS3_KO_LIVER_UP, GSE369_PRE_VS_POST_IL6_INJECTION_SOCS3_KO_LIVER_UP
GOBP_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN, GOBP_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN
GSE17322_CD103_POS_VS_CD11B_HIGH_LUNG_DC_UP, GSE17322_CD103_POS_VS_CD11B_HIGH_LUNG_DC_UP
GSE22886_IGM_MEMORY_BCELL_VS_BM_PLASMA_CELL_DN, GSE22886_IGM_MEMORY_BCELL_VS_BM_PLASMA_CELL_DN
GSE29617_CTRL_VS_DAY7_TIV_FLU_VACCINE_PBMIC_2008_DN, GSE29617_CTRL_VS_DAY7_TIV_FLU_VACCINE_PBMIC_2008_DN
GSE1460_CD4_THYMOCYTE_VS_THYMIC_STROMAL_CELL_DN, GSE1460_CD4_THYMOCYTE_VS_THYMIC_STROMAL_CELL_DN
GSE4142_GC_BCELL_VS_MEMORY_BCELL_UP, GSE4142_GC_BCELL_VS_MEMORY_BCELL_UP
GSE9239_CTRL_VS_TNF_INHIBITOR_TREATED_DC_UP, GSE9239_CTRL_VS_TNF_INHIBITOR_TREATED_DC_UP
GSE1460_INTRATHYMIC_T_PROGENITOR_VS_THYMIC_STROMAL_CELL_DN, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_THYMIC_STROMAL_CELL_DN
HAY_BONE_MARROW_DENDRITIC_CELL, HAY_BONE_MARROW_DENDRITIC_CELL
GSE13493_DP_VS_CD8POS_THYMOCYTE_DN, GSE13493_DP_VS_CD8POS_THYMOCYTE_DN
BROWNE_HCMV_INFECTION_48HR_UP, BROWNE_HCMV_INFECTION_48HR_UP
GSE360_L_DONOVANI_VS_M_TUBERCULOSIS_MAC_UP, GSE360_L_DONOVANI_VS_M_TUBERCULOSIS_MAC_UP
TGASTMAGC_NFE2_01, TGASTMAGC_NFE2_01
GSE360_L_DONOVANI_VS_L_MAJOR_DC_DN, GSE360_L_DONOVANI_VS_L_MAJOR_DC_DN
GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY, GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY
TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_10D_DN, TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_10D_DN
PELLICCIOTTA_HDAC_IN_ANTIGEN_PRESENTATION_UP, PELLICCIOTTA_HDAC_IN_ANTIGEN_PRESENTATION_UP
HP_FETAL_ULTRASOUND_SOFT_MARKER, HP_FETAL_ULTRASOUND_SOFT_MARKER
GSE360_CTRL_VS_L_MAJOR_MAC_DN, GSE360_CTRL_VS_L_MAJOR_MAC_DN
HP_RECURRENT_FRACTURES, HP_RECURRENT_FRACTURES
PAX8_TARGET_GENES, PAX8_TARGET_GENES
GOBP_RESPONSE_TO_PURINE_CONTAINING_COMPOUND, GOBP_RESPONSE_TO_PURINE_CONTAINING_COMPOUND
HP_PSYCHOSIS, HP_PSYCHOSIS
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_DN, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_DN
GOCC_LIPID_DROPLET, GOCC_LIPID_DROPLET
GOCC_CIS_GOLGI_NETWORK, GOCC_CIS_GOLGI_NETWORK
ZHANG_PROLIFERATING_VS_QUIESCENT, ZHANG_PROLIFERATING_VS_QUIESCENT
REACTOME_REGULATION_OF_INSULIN_LIKE_GROWTH_FACTOR_IGF_TRANSPORT_AND_UPTAKE_BY_INSULIN_LIKE_GROWTH_FACTOR_BINDING_PROTEINS_IGFBPS, REACTOME_REGULATION_OF_INSULIN_LIKE_GROWTH_FACTOR_IGF_TRANSPORT_AND_UPTAKE_BY_INSULIN_LIKE_GROWTH_FACTOR_BINDING_PROTEINS_IGFBPS
HP_THICKENED_NUCHAL_SKIN_FOLD, HP_THICKENED_NUCHAL_SKIN_FOLD
GOBP_RESPONSE_TO_CALCIIUM_ION, GOBP_RESPONSE_TO_CALCIIUM_ION
GOBP_RESPONSE_TO_ORGANOPHOSPHORUS, GOBP_RESPONSE_TO_ORGANOPHOSPHORUS
PEDERSEN_TARGETS_OF_611CTF_ISOFORM_OF_ERBB2, PEDERSEN_TARGETS_OF_611CTF_ISOFORM_OF_ERBB2
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN
GOBP_RESPONSE_TO_NUTRIENT, GOBP_RESPONSE_TO_NUTRIENT
HENDRICKS_SMARCA4_TARGETS_UP, HENDRICKS_SMARCA4_TARGETS_UP
HP_MILDLY_ELEVATED_CREATINE_KINASE, HP_MILDLY_ELEVATED_CREATINE_KINASE
HP_SCAPULAR_WINGING, HP_SCAPULAR_WINGING
HP_DECREASED_MUSCLE_MASS, HP_DECREASED_MUSCLE_MASS
BARRIER_CANCER_RELAPSE_NORMAL_SAMPLE_DN, BARRIER_CANCER_RELAPSE_NORMAL_SAMPLE_DN
LEE_LIVER_CANCER_ACOX1_UP, LEE_LIVER_CANCER_ACOX1_UP
MODULE_310, MODULE_310
GOBP_RESPONSE_TO_CAMP, GOBP_RESPONSE_TO_CAMP
PID_FRA_PATHWAY, PID_FRA_PATHWAY
GSE18148_CBBF_KO_VS_WT_TREG_UP, GSE18148_CBBF_KO_VS_WT_TREG_UP
MORF_CASP10, MORF_CASP10
HP_TURRICEPHALY, HP_TURRICEPHALY
HP_ACANTHOSIS_NIGRICANS, HP_ACANTHOSIS_NIGRICANS
WESTON_VEGFA_TARGETS, WESTON_VEGFA_TARGETS
HP_ABNORMAL_CIRCULATING_NUCLEOBASE_CONCENTRATION, HP_ABNORMAL_CIRCULATING_NUCLEOBASE_CONCENTRATION
TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_GNANULOCYTE_DN, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_GNANULOCYTE_DN
HP_DECREASED_HDL_CHOLESTEROL_CONCENTRATION, HP_DECREASED_HDL_CHOLESTEROL_CONCENTRATION
REACTOME_BINDING_AND_UPTAKE_OF_LIGANDS_BY_SCAVENGER_RECEPTORS, REACTOME_BINDING_AND_UPTAKE_OF_LIGANDS_BY_SCAVENGER_RECEPTORS
GOBP_NEGATIVE_REGULATION_OF_CILIIUM_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_CILIIUM_ASSEMBLY
HP_EEG_WITH_GENERALIZED_SLOW_ACTIVITY, HP_EEG_WITH_GENERALIZED_SLOW_ACTIVITY
GOBP_ER_ASSOCIATED_MISFOLDED_PROTEIN_CATABOLIC_PROCESS, GOBP_ER_ASSOCIATED_MISFOLDED_PROTEIN_CATABOLIC_PROCESS
GOBP_BILE_ACID_METABOLIC_PROCESS, GOBP_BILE_ACID_METABOLIC_PROCESS
HUL_MAPK14_TARGETS_UP, HUL_MAPK14_TARGETS_UP
REACTOME_COLLAGEN_FORMATION, REACTOME_COLLAGEN_FORMATION
GOBP_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE_BY_CIRCULATORY_RENIN_ANGIOTENSIN, GOBP_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE_BY_CIRCULATORY_RENIN_ANGIOTENSIN
HP_ABNORMAL_SHARPEY_FIBER_MORPHOLOGY, HP_ABNORMAL_SHARPEY_FIBER_MORPHOLOGY
HP_GRAMMAR_SPECIFIC_SPEECH_DISORDER, HP_GRAMMAR_SPECIFIC_SPEECH_DISORDER
HP_FRONTOTEMPORAL_CEREBRAL_ATROPHY, HP_FRONTOTEMPORAL_CEREBRAL_ATROPHY
LUND_SILENCED_BY_METHYLATION, LUND_SILENCED_BY_METHYLATION
GOBP_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE_BY_RENIN_ANGIOTENSIN, GOBP_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE_BY_RENIN_ANGIOTENSIN
REACTOME_METABOLISM_OF_ANGIOTENSINOGEN_TO_ANGIOTENSINS, REACTOME_METABOLISM_OF_ANGIOTENSINOGEN_TO_ANGIOTENSINS
GOBP_NEGATIVE_REGULATION_OF_INTERLEUKIN_1_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_INTERLEUKIN_1_PRODUCTION
TURASHVILI_BREAST_CARCINOMA_DUCTAL_VS_LOBULAR_UP, TURASHVILI_BREAST_CARCINOMA_DUCTAL_VS_LOBULAR_UP
RAMPON_ENRICHED_LEARNING_ENVIRONMENT_EARLY_UP, RAMPON_ENRICHED_LEARNING_ENVIRONMENT_EARLY_UP
GNF2_CYP2B6, GNF2_CYP2B6
GOBP_AMINO_SUGAR_BIOSYNTHETIC_PROCESS, GOBP_AMINO_SUGAR_BIOSYNTHETIC_PROCESS
HP_ABNORMAL_HDL_CHOLESTEROL_CONCENTRATION, HP_ABNORMAL_HDL_CHOLESTEROL_CONCENTRATION
HP_ESOPHAGEAL_STRICTURE, HP_ESOPHAGEAL_STRICTURE
GOBP_REGULATION_OF_PROTEIN_FOLDING, GOBP_REGULATION_OF_PROTEIN_FOLDING
HP_CHONDROCALCINOSIS, HP_CHONDROCALCINOSIS
FARMER_BREAST_CANCER_CLUSTER_7, FARMER_BREAST_CANCER_CLUSTER_7
GOBP_ACTIVATION_OF_ADENYLATE_CYCLASE_ACTIVITY, GOBP_ACTIVATION_OF_ADENYLATE_CYCLASE_ACTIVITY
HP_ASPIRATION, HP_ASPIRATION
HP_PHOTOSENSITIVE_MYOCLONIC_SEIZURE, HP_PHOTOSENSITIVE_MYOCLONIC_SEIZURE
TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_SUSTAINED_IN_MONOCYTE_UP, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_SUSTAINED_IN_MONOCYTE_UP
GOBP_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE_MEDIATED_BY_A_CHEMICAL_SIGNAL, GOBP_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE_MEDIATED_BY_A_CHEMICAL_SIGNAL
WEINMANN_ADAPTATION_TO_HYPOXIA_UP, WEINMANN_ADAPTATION_TO_HYPOXIA_UP
BIOCARTA_CYTOKINE_PATHWAY, BIOCARTA_CYTOKINE_PATHWAY
HP_APLASIA_OF_THE_MIDDLE_PHALANX_OF_THE_HAND, HP_APLASIA_OF_THE_MIDDLE_PHALANX_OF_THE_HAND
GOBP_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE_BY_HORMONE, GOBP_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE_BY_HORMONE
GOBP_NEGATIVE_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_PROLIFERATION, GOBP_NEGATIVE_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_PROLIFERATION
GOCC_CIS_GOLGI_NETWORK_MEMBRANE, GOCC_CIS_GOLGI_NETWORK_MEMBRANE
GOBP_GLUTAMATE_HOMEOSTASIS, GOBP_GLUTAMATE_HOMEOSTASIS
SASAI_TARGETS_OF_CXCR6_AND_PTCH1_DN, SASAI_TARGETS_OF_CXCR6_AND_PTCH1_DN
KORKOLA_CHORIOCARCINOMA, KORKOLA_CHORIOCARCINOMA
GOBP_NON_CANONICAL_WNT_SIGNALING_PATHWAY_VIA_MAPK_CASCADE, GOBP_NON_CANONICAL_WNT_SIGNALING_PATHWAY_VIA_MAPK_CASCADE
DESCARTES_FETAL_STOMACH_LYMPHATIC_ENDOTHELIAL_CELLS, DESCARTES_FETAL_STOMACH_LYMPHATIC_ENDOTHELIAL_CELLS
GOMF_PHOSPHOLIPASE_D_ACTIVITY, GOMF_PHOSPHOLIPASE_D_ACTIVITY
GOBP_REGULATION_OF_BILE_ACID_SECRETION, GOBP_REGULATION_OF_BILE_ACID_SECRETION
WP_EICOSANOID_METABOLISM_VIA_CYTOCHROME_P450_MONOOXYGENASES_CYP_PATHWAY, WP_EICOSANOID_METABOLISM_VIA_CYTOCHROME_P450_MONOOXYGENASES_CYP_PATHWAY