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GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_DN, GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_DN
GSE17974 2H VS 72H UNTREATED IN VITRO CD4 TCELL DN, GSE17974 2H VS 72H UNTREATED IN VITRO CD4 TCELL DN
GSE13738_RESTING_VS_TCR_ACTIVATED_CD4_TCELL_DN, GSE13738_RESTING_VS_TCR_ACTIVATED_CD4_TCELL_DN
GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_LOWDOSE_STIM_DC_UP, GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_LOWDOSE_STIM_DC_UP
GSE13738_RESTING_VS_BYSTANDER_ACTIVATED_CD4_TCELL_DN, GSE13738_RESTING_VS_BYSTANDER_ACTIVATED_CD4_TCELL_DN
GSE1460_DP_VS_CD4_THYMOCYTE_UP, GSE1460_DP_VS_CD4_THYMOCYTE_UP
GSE13738_TCR_VS_BYSTANDER_ACTIVATED_CD4_TCELL_UP, GSE13738_TCR_VS_BYSTANDER_ACTIVATED_CD4_TCELL_UP
GSE10240_IL22_VS_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN, GSE10240_IL22_VS_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN
GSE37301_PRO_BCELL_VS_RAG2_KO_NK_CELL_DN, GSE37301_PRO_BCELL_VS_RAG2_KO_NK_CELL_DN
WHITFIELD CELL CYCLE G1 S, WHITFIELD CELL CYCLE G1 S
MIR7161_5P, MIR7161_5P
GSE11864_CSF1_VS_CSF1_IFNG_IN_MAC_DN, GSE11864_CSF1_VS_CSF1_IFNG_IN_MAC_DN
GSE25087_TREG_VS_TCONV_FETUS_UP, GSE25087_TREG_VS_TCONV_FETUS_UP
GSE32986_UNSTIM_VS_GMCSF_STIM_DC_UP, GSE32986_UNSTIM_VS_GMCSF_STIM_DC_UP
GSE25088_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DN, GSE25088_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DN
GSE20727_CTRL_VS_DNFB_ALLERGEN_TREATED_DC_UP, GSE20727_CTRL_VS_DNFB_ALLERGEN_TREATED_DC_UP
GSE4142_NAIVE_BCELL_VS_PLASMA_CELL_DN, GSE4142_NAIVE_BCELL_VS_PLASMA_CELL_DN
ZNF547_TARGET_GENES, ZNF547_TARGET_GENES
MIR548U, MIR548U
BROWNE_HCMV_INFECTION_18HR_UP, BROWNE_HCMV_INFECTION_18HR_UP
GSE2706_R848_VS_LPS_8H_STIM_DC_DN, GSE2706_R848_VS_LPS_8H_STIM_DC_DN
WP_SIGNALING_PATHWAYS_IN_GLIOBLASTOMA, WP_SIGNALING_PATHWAYS_IN_GLIOBLASTOMA
GSE34156_TLR1_TLR2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_DN, GSE34156_TLR1_TLR2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_24H_TREATED_MC
GSE24574_BCL6_LOW_TFH_VS_NAIVE_CD4_TCELL_DN, GSE24574_BCL6_LOW_TFH_VS_NAIVE_CD4_TCELL_DN
MARTORIATI_MDM4_TARGETS_NEUROEPITHELIUM_UP, MARTORIATI_MDM4_TARGETS_NEUROEPITHELIUM_UP
KEGG_PANCREATIC_CANCER, KEGG_PANCREATIC_CANCER
GSE17721 CPG VS GARDIQUIMOD 6H BMDC UP, GSE17721 CPG VS GARDIQUIMOD 6H BMDC UP
GSE46242_CTRL_VS_EGR2_DELETED_ANERGIC_TH1_CD4_TCELL_DN, GSE46242_CTRL_VS_EGR2_DELETED_ANERGIC_TH1_CD4_TCELL_DN
GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_DN, GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_DN
ATF2_UP.V1_DN, ATF2_UP.V1_DN
GSE40441_NRP1_POS_INDUCED_TREG_VS_NRP1_NEG_NATURAL_TREG_DN, GSE40441_NRP1_POS_INDUCED_TREG_VS_NRP1_NEG_NATURAL_TREG_DN
WP_PANCREATIC_ADENOCARCINOMA_PATHWAY, WP_PANCREATIC_ADENOCARCINOMA_PATHWAY
GOMF_ENDONUCLEASE_ACTIVITY, GOMF_ENDONUCLEASE_ACTIVITY
GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_DN, GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_DN
KEGG_APOPTOSIS, KEGG_APOPTOSIS
MIR6792 5P, MIR6792 5P
AKT_UP_MTOR_DN.V1_DN, AKT_UP_MTOR_DN.V1_DN
GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS, GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS
WAMUNYOKOLI_OVARIAN_CANCER_GRADES_1_2_UP, WAMUNYOKOLI_OVARIAN_CANCER_GRADES_1_2_UP
MIR4764_3P, MIR4764_3P
KEGG SMALL CELL LUNG CANCER, KEGG SMALL CELL LUNG CANCER
MIR3167, MIR3167
HP_ANAL_ATRESIA, HP_ANAL_ATRESIA
WP STRUCTURAL PATHWAY OF INTERLEUKIN 1 IL1, WP STRUCTURAL PATHWAY OF INTERLEUKIN 1 IL1
GSE13306_TREG_VS_TCONV_SPLEEN_UP, GSE13306_TREG_VS_TCONV_SPLEEN_UP
GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNOGLOBULINS, GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNOGLOBULINS
HP ABNORMAL BLOOD INORGANIC CATION CONCENTRATION, HP ABNORMAL BLOOD INORGANIC CATION CONCENTRATION
GOBP_NEGATIVE_REGULATION_OF_DNA_RECOMBINATION, GOBP_NEGATIVE_REGULATION_OF_DNA_RECOMBINATION
LINDSTEDT_DENDRITIC_CELL_MATURATION_C, LINDSTEDT_DENDRITIC_CELL_MATURATION_C
MIR6855_5P, MIR6855_5P
SATO_SILENCED_EPIGENETICALLY_IN_PANCREATIC_CANCER, SATO_SILENCED_EPIGENETICALLY_IN_PANCREATIC_CANCER
MIR146B_3P, MIR146B_3P
GSE43955_1H_VS_42H_ACT_CD4_TCELL_DN, GSE43955_1H_VS_42H_ACT_CD4_TCELL_DN
GOBP_B_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_B_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE
HP_ABNORMAL_GROWTH_HORMONE_LEVEL, HP_ABNORMAL_GROWTH_HORMONE_LEVEL
GOBP_NEGATIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION, GOBP_NEGATIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_
WP_MIRNAS_INVOLVED_IN_DNA_DAMAGE_RESPONSE, WP_MIRNAS_INVOLVED_IN_DNA_DAMAGE_RESPONSE
HP_RHIZOMELIA, HP_RHIZOMELIA
PID_IL6_7_PATHWAY, PID_IL6_7_PATHWAY
HP_DISPROPORTIONATE_SHORT_LIMB_SHORT_STATURE, HP_DISPROPORTIONATE_SHORT_LIMB_SHORT_STATURE
GOMF_GROWTH_FACTOR_RECEPTOR_BINDING, GOMF_GROWTH_FACTOR_RECEPTOR_BINDING
REACTOME_SIGNALING_BY_LEPTIN, REACTOME_SIGNALING_BY_LEPTIN
MIR648, MIR648
MIR4294, MIR4294
RAY_TARGETS_OF_P210_BCR_ABL_FUSION_UP, RAY_TARGETS_OF_P210_BCR_ABL_FUSION_UP
REACTOME_RUNX2_REGULATES_BONE_DEVELOPMENT, REACTOME_RUNX2_REGULATES_BONE_DEVELOPMENT
LEIN_CEREBELLUM_MARKERS, LEIN_CEREBELLUM_MARKERS
GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGM_BCELL_DN, GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGM_BCELL_DN
MODULE_360, MODULE_360
MODULE_545, MODULE_545
GOBP_TOLL_LIKE_RECEPTOR_9_SIGNALING_PATHWAY, GOBP_TOLL_LIKE_RECEPTOR_9_SIGNALING_PATHWAY
GOMF_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY_TRANSPOSING_C_C_BONDS, GOMF_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY_TRANSPOSING_C_C_BONDS
GOCC_EXTRINSIC_COMPONENT_OF_ENDOSOME_MEMBRANE, GOCC_EXTRINSIC_COMPONENT_OF_ENDOSOME_MEMBRANE
SIMBULAN_PARP1_TARGETS_UP, SIMBULAN_PARP1_TARGETS_UP
DAZARD_UV_RESPONSE_CLUSTER_G4, DAZARD_UV_RESPONSE_CLUSTER_G4
GOBP_ASYMMETRIC_CELL_DIVISION, GOBP_ASYMMETRIC_CELL_DIVISION
MODULE_511, MODULE_511
LA_MEN1_TARGETS, LA_MEN1_TARGETS
HP ABNORMAL MEGAKARYOCYTE MORPHOLOGY, HP ABNORMAL MEGAKARYOCYTE MORPHOLOGY
HAY_BONE_MARROW_CD34_POS_MULTILIN, HAY_BONE_MARROW_CD34_POS_MULTILIN
GNF2 CCNA1, GNF2 CCNA1
REACTOME_RUNX2_REGULATES_OSTEOBLAST_DIFFERENTIATION, REACTOME_RUNX2_REGULATES_OSTEOBLAST_DIFFERENTIATION
LE_SKI_TARGETS_DN, LE_SKI_TARGETS_DN
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RDLAN HIGHDOSE STIM DC UP, GSE32986 UNSTIM VS CURDLAN HIGHDOSE STIM DC UP

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GOBP_INTERLEUKIN_7_MEDIATED_SIGNALING_PATHWAY, GOBP_INTERLEUKIN_7_MEDIATED_SIGNALING_PATHWAY
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REACTOME EGFR INTERACTS WITH PHOSPHOLIPASE C GAMMA, REACTOME EGFR INTERACTS WITH PHOSPHOLIPASE C GAMMA

REACTOME THE RETINOID CYCLE IN CONES DAYLIGHT VISION, REACTOME THE RETINOID CYCLE IN CONES DAYLIGHT VISION

HP_ABNORMALITY_OF_ENDOCRINE_PANCREAS_PHYSIOLOGY, HP_ABNORMALITY_OF_ENDOCRINE_PANCREAS_PHYSIOLOGY

GOBP_OLIGOPEPTIDE_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_OLIGOPEPTIDE_IMPORT_ACROSS_PLASMA_MEMBRANE

REACTOME_INTERLEUKIN_6_FAMILY_SIGNALING, REACTOME_INTERLEUKIN_6_FAMILY_SIGNALING

MAHADEVAN_GIST_MORPHOLOGICAL_SWITCH, MAHADEVAN_GIST_MORPHOLOGICAL_SWITCH

HP_ABNORMAL_VAS_DEFERENS_MORPHOLOGY, HP_ABNORMAL_VAS_DEFERENS_MORPHOLOGY

GOMF_INTERLEUKIN_12_RECEPTOR_BINDING, GOMF_INTERLEUKIN_12_RECEPTOR_BINDING

GOBP_BRANCHIOMOTOR_NEURON_AXON_GUIDANCE, GOBP_BRANCHIOMOTOR_NEURON_AXON_GUIDANCE

GOBP_NEGATIVE_REGULATION_OF_BLOOD_CIRCULATION, GOBP_NEGATIVE_REGULATION_OF_BLOOD_CIRCULATION

HP_FETAL_POLYURIA, HP_FETAL_POLYURIA

HP_HYPOCHLOREMIA, HP_HYPOCHLOREMIA

HP_ACHOLIC_STOOLS, HP_ACHOLIC_STOOLS

HP_KETOACIDOSIS, HP_KETOACIDOSIS

PID_ERBB_NETWORK_PATHWAY, PID_ERBB_NETWORK_PATHWAY

GNF2 MLF1, GNF2 MLF1 chr11q24, chr11q24 GOBP_ADENYLATE_CYCLASE_INHIBITING_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY, GOBP_ADENYLATE_CYCLASE_INHIBITING_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY chr3q28, chr3q28

REACTOME_REGULATION_OF_GENE_EXPRESSION_IN_EARLY_PANCREATIC_PRECURSOR_CELLS, REACTOME_REGULATION_OF_GENE_EXPRESSION_IN_EARLY_PANCREATIC_PRECURSOR_

GOBP_ESTABLISHMENT_OR_MAINTENANCE_OF_MONOPOLAR_CELL_POLARITY, GOBP_ESTABLISHMENT_OR_MAINTENANCE_OF_MONOPOLAR_CELL_POLARITY