7_POS_VS_NEG_NK_CELL_UP, GSE23695_CD57_POS_VS_NEG_NK_CELL_UP

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GSE22103_UNSTIM_VS_GMCSF_AND_IFNG_STIM_NEUTROPHIL_UP, GSE22103_UNSTIM_VS_GMCSF_AND_IFNG_STIM_NEUTROPHIL_UP
GSE22589_HEALTHY_VS_SIV_INFECTED_DC_UP, GSE22589_HEALTHY_VS_SIV_INFECTED_DC_UP
GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN, GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MYD88_KO_MACROPH
GSE42021_CD24INT_TREG_VS_CD24INT_TCONV_THYMUS_UP, GSE42021_CD24INT_TREG_VS_CD24INT_TCONV_THYMUS_UP
GSE8685_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP, GSE8685_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP
GSE34392_ST2_KO_VS_WT_DAY8_LCMV_EFFECTOR_CD8_TCELL_UP, GSE34392_ST2_KO_VS_WT_DAY8_LCMV_EFFECTOR_CD8_TCELL_UP
GSE17721_LPS_VS_CPG_12H_BMDC_UP, GSE17721_LPS_VS_CPG_12H_BMDC_UP
GSE17721_0.5H_VS_4H_PAM3CSK4_BMDC_UP, GSE17721_0.5H_VS_4H_PAM3CSK4_BMDC_UP
GSE13493_DP_VS_CD8POS_THYMOCYTE_DN, GSE13493_DP_VS_CD8POS_THYMOCYTE_DN
GSE11961_MARGINAL_ZONE_BCELL_VS_PLASMA_CELL_DAY7_DN, GSE11961_MARGINAL_ZONE_BCELL_VS_PLASMA_CELL_DAY7_DN
MORF_HEAB, MORF_HEAB
GSE40274_FOXP3_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATE
BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_TRANS, BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_TRANS
GSE43863_DAY6_EFF_VS_DAY150_MEM_TH1_CD4_TCELL_DN, GSE43863_DAY6_EFF_VS_DAY150_MEM_TH1_CD4_TCELL_DN
GSE17721_PAM3CSK4_VS_CPG_1H_BMDC_UP, GSE17721_PAM3CSK4_VS_CPG_1H_BMDC_UP
GSE14415_FOXP3_KO_NATURAL_TREG_VS_TCONV_DN, GSE14415_FOXP3_KO_NATURAL_TREG_VS_TCONV_DN
GSE21033_3H_VS_12H_POLYIC_STIM_DC_UP, GSE21033_3H_VS_12H_POLYIC_STIM_DC_UP
KEGG_NOTCH_SIGNALING_PATHWAY, KEGG_NOTCH_SIGNALING_PATHWAY
GAGCCTG_MIR484, GAGCCTG_MIR484
GSE17721_CTRL_VS_LPS_2H_BMDC_DN, GSE17721_CTRL_VS_LPS_2H_BMDC_DN
BIOCARTA_AKT_PATHWAY, BIOCARTA_AKT_PATHWAY
GSE36527_CD62L_HIGH_VS_CD62L_LOW_TREG_CD69_NEG_KLRG1_NEG_DN, GSE36527_CD62L_HIGH_VS_CD62L_LOW_TREG_CD69_NEG_KLRG1_NEG_DN
GO_CELLULAR_ALDEHYDE_METABOLIC_PROCESS, GO_CELLULAR_ALDEHYDE_METABOLIC_PROCESS
BILANGES_RAPAMYCIN_SENSITIVE_GENES, BILANGES_RAPAMYCIN_SENSITIVE_GENES
GO_AORTA_MORPHOGENESIS, GO_AORTA_MORPHOGENESIS
GO_PHOSPHATIDYLINOSITOL_PHOSPHATE_KINASE_ACTIVITY, GO_PHOSPHATIDYLINOSITOL_PHOSPHATE_KINASE_ACTIVITY
GO_MATURATION_OF_LSU_RRNA, GO_MATURATION_OF_LSU_RRNA
LANDIS_ERBB2_BREAST_TUMORS_65_UP, LANDIS_ERBB2_BREAST_TUMORS_65_UP
GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_DN, GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_DN
GO_GOLGI_CISTERNA_MEMBRANE, GO_GOLGI_CISTERNA_MEMBRANE
GO_SIALYLATION, GO_SIALYLATION
GO_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE, GO_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE
GO_PHOSPHATIDYLINOSITOL_3_PHOSPHATE_BINDING, GO_PHOSPHATIDYLINOSITOL_3_PHOSPHATE_BINDING
IWANAGA_CARCINOGENESIS_BY_KRAS_UP, IWANAGA_CARCINOGENESIS_BY_KRAS_UP
GO_PROTEIN_PHOSPHATASE_TYPE_2A_COMPLEX, GO_PROTEIN_PHOSPHATASE_TYPE_2A_COMPLEX
DUNNE_TARGETS_OF_AML1_MTG8_FUSION_DN, DUNNE_TARGETS_OF_AML1_MTG8_FUSION_DN
GO_CLATHRIN_COATED_VESICLE_MEMBRANE, GO_CLATHRIN_COATED_VESICLE_MEMBRANE
DAVICIONI_RHABDOMYOSARCOMA_PAX_FOXO1_FUSION_UP, DAVICIONI_RHABDOMYOSARCOMA_PAX_FOXO1_FUSION_UP
ELLWOOD_MYC_TARGETS_UP, ELLWOOD_MYC_TARGETS_UP
BUCKANOVICH_T_LYMPHOCYTE_HOMING_ON_TUMOR_UP, BUCKANOVICH_T_LYMPHOCYTE_HOMING_ON_TUMOR_UP
GO_AORTA_DEVELOPMENT, GO_AORTA_DEVELOPMENT
PID_INTEGRIN_A4B1_PATHWAY, PID_INTEGRIN_A4B1_PATHWAY
GO_PROTEIN_SELF_ASSOCIATION, GO_PROTEIN_SELF_ASSOCIATION
GO_POSITIVE_REGULATION_OF_RESPONSE_TO_CYTOKINE_STIMULUS, GO_POSITIVE_REGULATION_OF_RESPONSE_TO_CYTOKINE_STIMULUS
MATZUK_IMPLANTATION_AND_UTERINE, MATZUK_IMPLANTATION_AND_UTERINE
GO_CARBOHYDRATE_DERIVATIVE_TRANSPORTER_ACTIVITY, GO_CARBOHYDRATE_DERIVATIVE_TRANSPORTER_ACTIVITY
GO_DEVELOPMENTAL_MATURATION, GO_DEVELOPMENTAL_MATURATION
MODULE_560, MODULE_560
BIOCARTA_ECM_PATHWAY, BIOCARTA_ECM_PATHWAY
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