_KLRG1HIGH_EFF_CD8_TCELL_DN, GSE10239_MEMORY_VS_KLRG1HIGH_EFF_CD8_TCELL_DN

MODULE_71, MODULE_71

LEE_LIVER_CANCER_MYC_TGFA_UP, LEE_LIVER_CANCER_MYC_TGFA_UP

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GSE10239_MEMORY_VS_KLRG1INT_EFF_CD8_TCELL_DN, GSE10239_MEMORY_VS_KLRG1INT_EFF_CD8_TCELL_DN
GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP
GSE7768_OVA_ALONE_VS_OVA_WITH_MPL_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP, GSE7768_OVA_ALONE_VS_OVA_WITH_MPL_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP
GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_TREATED_CD4_TCELL_UP
GSE8835_HEALTHY_VS_CLL_CD8_TCELL_UP, GSE8835_HEALTHY_VS_CLL_CD8_TCELL_UP
GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_IL12_CD8_TCELL_DN, GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_IL12_CD8_TCELL_DN
GSE21670_STAT3_KO_VS_WT_CD4_TCELL_IL6_TREATED_UP, GSE21670_STAT3_KO_VS_WT_CD4_TCELL_IL6_TREATED_UP
GSE19923_E2A_KO_VS_E2A_AND_HEB_KO_DP_THYMOCYTE_UP, GSE19923_E2A_KO_VS_E2A_AND_HEB_KO_DP_THYMOCYTE_UP
GSE46025_WT_VS_FOXO1_KO_KLRG1_LOW_CD8_EFFECTOR_TCELL_DN, GSE46025_WT_VS_FOXO1_KO_KLRG1_LOW_CD8_EFFECTOR_TCELL_DN
GSE7460_TREG_VS_TCONV_ACT_UP, GSE7460_TREG_VS_TCONV_ACT_UP
GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_DN, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_DN
GSE13484_UNSTIM_VS_3H_YF17D_VACCINE_STIM_PBMC_DN, GSE13484_UNSTIM_VS_3H_YF17D_VACCINE_STIM_PBMC_DN
GSE7460_WT_VS_FOXP3_HET_ACT_WITH_TGFB_TCONV_DN, GSE7460_WT_VS_FOXP3_HET_ACT_WITH_TGFB_TCONV_DN
GSE17721_LPS_VS_PAM3CSK4_16H_BMDC_UP, GSE17721_LPS_VS_PAM3CSK4_16H_BMDC_UP
GSE26669_CD4_VS_CD8_TCELL_IN_MLR_COSTIM_BLOCK_UP, GSE26669_CD4_VS_CD8_TCELL_IN_MLR_COSTIM_BLOCK_UP
GSE21379 WT VS SAP KO CD4 TCELL DN, GSE21379 WT VS SAP KO CD4 TCELL DN
GSE20366_CD103_KLRG1_DP_VS_DN_TREG_DN, GSE20366_CD103_KLRG1_DP_VS_DN_TREG_DN
GSE32034_LY6C_HIGH_VS_LOW_ROSIGLIZATONE_TREATED_MONOCYTE_DN, GSE32034_LY6C_HIGH_VS_LOW_ROSIGLIZATONE_TREATED_MONOCYTE_DN
GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN, GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN
GSE37605_TREG_VS_TCONV_C57BL6_FOXP3_FUSION_GFP_DN, GSE37605_TREG_VS_TCONV_C57BL6_FOXP3_FUSION_GFP_DN
GO_POST_GOLGI_VESICLE_MEDIATED_TRANSPORT, GO_POST_GOLGI_VESICLE_MEDIATED_TRANSPORT
GSE23114_WT_VS_SLE2C1_MOUSE_PERITONEAL_CAVITY_B1A_BCELL_UP, GSE23114_WT_VS_SLE2C1_MOUSE_PERITONEAL_CAVITY_B1A_BCELL_UP
GO_PROTEIN_TETRAMERIZATION, GO_PROTEIN_TETRAMERIZATION
GSE32255_WT_UNSTIM_VS_IMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_DN, GSE32255_WT_UNSTIM_VS_IMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_DN
GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_UP, GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_UP
chr14q24, chr14q24
GO_PHOSPHATIDYLINOSITOL_BINDING, GO_PHOSPHATIDYLINOSITOL_BINDING
GO_PROTEIN_HOMOTETRAMERIZATION, GO_PROTEIN_HOMOTETRAMERIZATION
GNF2_MAP2K3, GNF2_MAP2K3
chr6q25, chr6q25
SESTO_RESPONSE_TO_UV_C8, SESTO_RESPONSE_TO_UV_C8
GO REGULATION OF TOR SIGNALING, GO REGULATION OF TOR SIGNALING
GO_NUCLEAR_ENVELOPE_REASSEMBLY, GO_NUCLEAR_ENVELOPE_REASSEMBLY
GNF2_TAL1, GNF2_TAL1
GSE21927_SPLENIC_C26GM_TUMOROUS_VS_4T1_TUMOR_MONOCYTES_UP, GSE21927_SPLENIC_C26GM_TUMOROUS_VS_4T1_TUMOR_MONOCYTES_UP
REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATION_MEDIATED_BY_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATED_PHOSPHORYLATION_AND_ACTIVATED_PHOSPHORYLATION_AND_ACTIVATED_PHOSPHORYLATION_AND_ACTIVATED_PHOSPHORYLATION_AND_ACTIVATED_PHOSPHORYLATION_AND_ACTIVATED_PHOSPHORYLATION_AND_ACTIVATED_PHOSPHORYLATION_AND_ACTIVATED_PHOSPHORYLATION_AND_ACTIVATED_PHOSPHORYLATION_AND_ACTI
GSE28237_EARLY_VS_LATE_GC_BCELL_UP, GSE28237_EARLY_VS_LATE_GC_BCELL_UP
GO_TBP_CLASS_PROTEIN_BINDING, GO_TBP_CLASS_PROTEIN_BINDING
GO_MITOTIC_SISTER_CHROMATID_COHESION, GO_MITOTIC_SISTER_CHROMATID_COHESION
GO_CYTOKINETIC_PROCESS, GO_CYTOKINETIC_PROCESS
GO_ADIPOSE_TISSUE_DEVELOPMENT, GO_ADIPOSE_TISSUE_DEVELOPMENT
GO_POSITIVE_REGULATION_OF_CYTOKINESIS, GO_POSITIVE_REGULATION_OF_CYTOKINESIS
GO_RAC_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_ACTIVITY, GO_RAC_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_ACTIVITY
GO_NEGATIVE_REGULATION_OF_ERBB_SIGNALING_PATHWAY, GO_NEGATIVE_REGULATION_OF_ERBB_SIGNALING_PATHWAY
ZEMBUTSU_SENSITIVITY_TO_FLUOROURACIL, ZEMBUTSU_SENSITIVITY_TO_FLUOROURACIL
GO_REGULATION_OF_RECEPTOR_ACTIVITY, GO_REGULATION_OF_RECEPTOR_ACTIVITY
VISALA_RESPONSE_TO_HEAT_SHOCK_AND_AGING_UP, VISALA_RESPONSE_TO_HEAT_SHOCK_AND_AGING_UP
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GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP