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GSE25890_CTRL_VS_IL33_IL7_TREATED_NUOCYTES_DN, GSE25890_CTRL_VS_IL33_IL7_TREATED_NUOCYTES_
GSE20198 UNTREATED VS IL12 TREATED ACT CD4 TCELL UP, GSE20198 UNTREATED VS IL12 TREATED
GSE14413_UNSTIM_VS_IFNB_STIM_NIH3T3_CELLS_DN, GSE14413_UNSTIM_VS_IFNB_STIM_NIH3T3_CELLS_DN
GSE2128 C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_DN, GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCY
PID ERBB1 DOWNSTREAM PATHWAY, PID ERBB1 DOWNSTREAM PATHWAY
GSE5099_MONOCYTE_VS_ALTERNATIVE_M2_MACROPHAGE_DN, GSE5099_MONOCYTE_VS_ALTERNATIVE_N
GSE17721_LPS_VS_CPG_0.5H_BMDC_UP, GSE17721_LPS_VS_CPG_0.5H_BMDC_UP
PID MET PATHWAY, PID MET PATHWAY
BHATI_G2M_ARREST_BY_2METHOXYESTRADIOL_DN, BHATI_G2M_ARREST_BY_2METHOXYESTRADIOL_DN
GO_LYASE_ACTIVITY, GO_LYASE_ACTIVITY
GO ENDOSOME_ORGANIZATION, GO_ENDOSOME_ORGANIZATION
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_
GO CELLULAR RESPONSE TO DSRNA, GO CELLULAR RESPONSE TO DSRNA
KEGG_MTOR_SIGNALING_PATHWAY, KEGG_MTOR_SIGNALING_PATHWAY
GSE12003_4D_VS_8D_CULTURE_BM_PROGENITOR_DN, GSE12003_4D_VS_8D_CULTURE_BM_PROGENITOR_DN
GO_MYELOID_CELL_DEVELOPMENT, GO_MYELOID_CELL_DEVELOPMENT
DELLA_RESPONSE_TO_TSA_AND_BUTYRATE, DELLA_RESPONSE_TO_TSA_AND_BUTYRATE
PID THROMBIN PAR1 PATHWAY, PID THROMBIN PAR1 PATHWAY
chr1p31, chr1p31
REACTOME_CD28_CO_STIMULATION, REACTOME_CD28_CO_STIMULATION
GO MULTI ORGANISM ORGANELLE ORGANIZATION, GO MULTI ORGANISM ORGANELLE ORGANIZATION
HSF_Q6, HSF_Q6
ZHU_SKIL_TARGETS_UP, ZHU_SKIL_TARGETS_UP
MANTOVANI NFKB TARGETS UP, MANTOVANI NFKB TARGETS UP
KUMAR_AUTOPHAGY_NETWORK, KUMAR_AUTOPHAGY_NETWORK
PID ER NONGENOMIC PATHWAY, PID ER NONGENOMIC PATHWAY
GSE37319 WT_VS_RC3H1 KO_CD44LOW_CD8_TCELL_DN, GSE37319 WT_VS_RC3H1 KO_CD44LOW_CD8_TCEL
GO DEAMINASE ACTIVITY, GO DEAMINASE ACTIVITY
GO_CARBON_CARBON_LYASE_ACTIVITY, GO_CARBON_CARBON_LYASE_ACTIVITY
CHESLER_BRAIN_HIGHEST_EXPRESSION, CHESLER_BRAIN_HIGHEST_EXPRESSION
GO PROTEIN INSERTION INTO MEMBRANE, GO PROTEIN INSERTION INTO MEMBRANE
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