

_AND_IL27_STIM_MACROPHAGE_UP, GSE44955_MCSF_VS_MCSF_AND_IL27_STIM_MACROPHAGE_UP

GSE41867_LCMV_ARMSTRONG_VS_CLONE13_DAY15_EFFECTOR_CD8_TCELL_UP, GSE41867_LCMV_ARMSTRONG_VS_CLONE13_DAY15_EFFECTOR_CD8_TCELL_UP
GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_UP, GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_UP
EIF4E_UP, EIF4E_UP
MIR4493, MIR4493
VART_KSHV_INFECTION_ANGIOGENIC_MARKERS_UP, VART_KSHV_INFECTION_ANGIOGENIC_MARKERS_UP
GTTGNYNNRGNAAC_UNKNOWN, GTTGNYNNRGNAAC_UNKNOWN
KATSANOU_ELAVL1_TARGETS_UP, KATSANOU_ELAVL1_TARGETS_UP
GSE43863_NAIVE_VS_TFH_CD4_EFF_TCELL_D6_LCMV_DN, GSE43863_NAIVE_VS_TFH_CD4_EFF_TCELL_D6_LCMV_DN
GOMF_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_T_CELL_MEDIATED_IMMUNITY, GOBP_POSITIVE_REGULATION_OF_T_CELL_MEDIATED_IMMUNITY
NAKAMURA_ADIPOGENESIS_LATE_DN, NAKAMURA_ADIPOGENESIS_LATE_DN
MODULE_400, MODULE_400
GOBP_NEGATIVE_REGULATION_OF_MAST_CELL_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_MAST_CELL_ACTIVATION
QIU_PBMCMENACTRA_AGE_18_45YO_ANTI_DT_ANTIBODY_CORRELATION_PROFILE_3DY_DN, QIU_PBMCMENACTRA_AGE_18_45YO_ANTI_DT_ANTIBODY_CORRELATION_PROFILE_3DY_DN
LI_PBMCMENACTRA_AGE_18_45YO_ANTI_DT_ANTIBODY_CORRELATION_PROFILE_3DY_DN, LI_PBMCMENACTRA_AGE_18_45YO_ANTI_DT_ANTIBODY_CORRELATION_PROFILE_3DY_DN
HOFMANN_MYELODYSPLASTIC_SYNDROM_LOW_RISK_UP, HOFMANN_MYELODYSPLASTIC_SYNDROM_LOW_RISK_UP
SUMI_HNF4A_TARGETS, SUMI_HNF4A_TARGETS