

HEATSHOCK_AND_LPS_STIM_MEF_DN, GSE16266_LPS_VS_HEATSHOCK_AND_LPS_STIM_MEF_DN

- GSE19198_CTRL_VS_IL21_TREATED_TCELL_6H_UP, GSE19198_CTRL_VS_IL21_TREATED_TCELL_6H_UP
- GSE17721_PAM3CSK4_VS_GADIQUIMOD_24H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_24H_BMDC_DN
- GSE19198_6H_VS_24H_IL21_TREATED_TCELL_DN, GSE19198_6H_VS_24H_IL21_TREATED_TCELL_DN
- GSE13738_TCR_VS_BYSTANDER_ACTIVATED_CD4_TCELL_DN, GSE13738_TCR_VS_BYSTANDER_ACTIVATED_CD4_TCELL_DN
- GSE2197_IMMUNOSUPPRESSIVE_DNA_VS_UNTREATED_IN_DC_UP, GSE2197_IMMUNOSUPPRESSIVE_DNA_VS_UNTREATED_IN_DC_UP
- MIR520F_3P, MIR520F_3P
- GOMF_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY
- GSE22342_CD11C_HIGH_VS_LOW_DECIDUAL_MACROPHAGES_UP, GSE22342_CD11C_HIGH_VS_LOW_DECIDUAL_MACROPHAGES_UP
- MIR767_3P, MIR767_3P
- MIR600, MIR600
- MIR7158_3P, MIR7158_3P
- GOMF_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY, GOMF_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY
- BMI1_DN_MEL18_DN.V1_UP, BMI1_DN_MEL18_DN.V1_UP
- GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_6H_ACT_CD4_TCELL_UP, GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_6H_ACT_CD4_TCELL_UP
- GHANDHI_DIRECT_IRRADIATION_UP, GHANDHI_DIRECT_IRRADIATION_UP
- ROPERO_HDAC2_TARGETS, ROPERO_HDAC2_TARGETS
- WINZEN_DEGRADED_VIA_KHSRP, WINZEN_DEGRADED_VIA_KHSRP
- MIR4750_3P, MIR4750_3P
- MIR632, MIR632
- GOBP_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
- GOBP_OUTFLOW_TRACT_MORPHOGENESIS, GOBP_OUTFLOW_TRACT_MORPHOGENESIS
- MARTIN_NFKB_TARGETS_UP, MARTIN_NFKB_TARGETS_UP
- SEKI_INFLAMMATORY_RESPONSE_LPS_DN, SEKI_INFLAMMATORY_RESPONSE_LPS_DN
- MIR6801_3P, MIR6801_3P
- MIR325, MIR325
- HENDRICKS_SMARCA4_TARGETS_DN, HENDRICKS_SMARCA4_TARGETS_DN
- GOBP_CARDIAC_SEPTUM_MORPHOGENESIS, GOBP_CARDIAC_SEPTUM_MORPHOGENESIS
- REACTOME_NICOTINAMIDE_SALVAGING, REACTOME_NICOTINAMIDE_SALVAGING
- GOBP_POSITIVE_REGULATION_OF_GLUCOSE_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_GLUCOSE_METABOLIC_PROCESS
- MIR616_3P, MIR616_3P
- GOBP_HEAD_MORPHOGENESIS, GOBP_HEAD_MORPHOGENESIS
- GOBP_MELANOCYTE_DIFFERENTIATION, GOBP_MELANOCYTE_DIFFERENTIATION
- GSE7596_AKT_TRANSD_VS_CTRL_CD4_TCONV_WITH_TGFB_DN, GSE7596_AKT_TRANSD_VS_CTRL_CD4_TCONV_WITH_TGFB_DN
- GOBP_OUTFLOW_TRACT_SEPTUM_MORPHOGENESIS, GOBP_OUTFLOW_TRACT_SEPTUM_MORPHOGENESIS
- HP_DISTURBANCE_OF_FACIAL_EXPRESSION, HP_DISTURBANCE_OF_FACIAL_EXPRESSION
- GOBP_REGULATION_OF_CALCIIUM_MEDIATED_SIGNALING, GOBP_REGULATION_OF_CALCIIUM_MEDIATED_SIGNALING
- MIR346, MIR346
- GOBP_REGULATION_OF_PEPTIDYL_CYSSTEINE_S_NITROSYLATION, GOBP_REGULATION_OF_PEPTIDYL_CYSSTEINE_S_NITROSYLATION
- GOMF_LIPASE_ACTIVATOR_ACTIVITY, GOMF_LIPASE_ACTIVATOR_ACTIVITY
- GOBP_CRANIAL_NERVE_MORPHOGENESIS, GOBP_CRANIAL_NERVE_MORPHOGENESIS
- chr7p14, chr7p14
- BRUNEAU_SEPTATION_ATRIAL, BRUNEAU_SEPTATION_ATRIAL
- HP_APLASIA_HYPOPLASIA_OF_THE_2ND_METACARPAL, HP_APLASIA_HYPOPLASIA_OF_THE_2ND_METACARPAL
- GOCC_DENDRITIC_SPINE_MEMBRANE, GOCC_DENDRITIC_SPINE_MEMBRANE
- HP_ABNORMALITY_OF_THE_2ND_METACARPAL, HP_ABNORMALITY_OF_THE_2ND_METACARPAL
- HP_SPLIT_FOOT, HP_SPLIT_FOOT
- GOBP_MAMMARY_GLAND_FORMATION, GOBP_MAMMARY_GLAND_FORMATION