

CANCER_20Q13_AMPLIFICATION_DN, GINESTIER_BREAST_CANCER_20Q13_AMPLIFICATION_DN

GSE6269_HEALTHY_VS_STAPH_AUREUS_INF_PBMG_UP, GSE6269_HEALTHY_VS_STAPH_AUREUS_INF_PBMG_UP
GSE13738_RESTING_VS_TCR_ACTIVATED_CD4_TCELL_UP, GSE13738_RESTING_VS_TCR_ACTIVATED_CD4_TCELL_UP
GOBP_REGULATION_OF_RAS_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_RAS_PROTEIN_SIGNAL_TRANSDUCTION
GSE19888_ADENOSINE_A3R_INH_VS_TCELL_MEMBRANES_ACT_MAST_CELL_UP, GSE19888_ADENOSINE_A3R_INH_VS_TCELL_MEMBRANES_ACT_MAST_CELL_UP
ACACTGG_MIR199A_MIR199B, ACACTGG_MIR199A_MIR199B
GSE17721_CTRL_VS_POLYIC_8H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_8H_BMDC_DN
HP_NON_MOTOR_SEIZURE, HP_NON_MOTOR_SEIZURE
GSE12963_ENV_NEF_VS_ENV_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_UP, GSE12963_ENV_NEF_VS_ENV_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_UP
MIR330_5P, MIR330_5P
GSE26495_PD1HIGH_VS_PD1LOW_CD8_TCELL_UP, GSE26495_PD1HIGH_VS_PD1LOW_CD8_TCELL_UP
GOBP_POSITIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC, GOBP_POSITIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC
MIR6753_3P, MIR6753_3P
MIR7107_3P, MIR7107_3P
MIR10525_3P, MIR10525_3P
MIR6727_3P, MIR6727_3P
SQSTM1_TARGET_GENES, SQSTM1_TARGET_GENES
MIR7855_5P, MIR7855_5P
GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_UP, GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_UP
GSE19941_IL10_KO_VS_IL10_KO_AND_NFKBP50_KO_UNSTIM_MACROPHAGE_UP, GSE19941_IL10_KO_VS_IL10_KO_AND_NFKBP50_KO_UNSTIM_MACROPHAGE_UP
HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_LOWER_LIMBS, HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_LOWER_LIMBS
GOBP_ENERGY_HOMEOSTASIS, GOBP_ENERGY_HOMEOSTASIS
GSE18281_PERIMEDULLARY_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_DN, GSE18281_PERIMEDULLARY_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_DN
REACTOME_MET_ACTIVATES_RAP1_AND_RAC1, REACTOME_MET_ACTIVATES_RAP1_AND_RAC1
PID_NECTIN_PATHWAY, PID_NECTIN_PATHWAY
GOMF_VINCULIN_BINDING, GOMF_VINCULIN_BINDING
GSE32901_TH17_EMRIKED_VS_TH17_NEG_CD4_TCELL_UP, GSE32901_TH17_EMRIKED_VS_TH17_NEG_CD4_TCELL_UP
GOBP_CORPUS_CALLOSUM_DEVELOPMENT, GOBP_CORPUS_CALLOSUM_DEVELOPMENT
GOBP_DIACYLGLYCEROL_METABOLIC_PROCESS, GOBP_DIACYLGLYCEROL_METABOLIC_PROCESS
GOCC_GATOR2_COMPLEX, GOCC_GATOR2_COMPLEX
GOMF_NITRIC_OXIDE_SYNTHASE_BINDING, GOMF_NITRIC_OXIDE_SYNTHASE_BINDING
GOBP_CHROMATIN_MEDIATED_MAINTENANCE_OF_TRANSCRIPTION, GOBP_CHROMATIN_MEDIATED_MAINTENANCE_OF_TRANSCRIPTION
GCM_PRKCG, GCM_PRKCG
SUZUKI_RESPONSE_TO_TSA_AND_DECITABINE_1B, SUZUKI_RESPONSE_TO_TSA_AND_DECITABINE_1B
XU_HGF_TARGETS_INDUCED_BY_AKT1_48HR_UP, XU_HGF_TARGETS_INDUCED_BY_AKT1_48HR_UP