

GSE13484_UNSTIM_VS_YF1D7_VACCINE_STIM_PPMC_UP, GSE13484_UNSTIM_VS_YF1D7_VACCINE_STIM_PPMC_UP
GSE9988_ANTL_TREMI_VS_ANTL_TREMI_AND_LPS_MONOCYTE_UP, GSE9988_ANTL_TREMI_VS_ANTL_TREMI_AND_LPS_MONOCYTE_UP
GSE9988_ANTL_TREMI_VS_CTRL_TREATED_MONOCYTES_DN, GSE9988_ANTL_TREMI_VS_CTRL_TREATED_MONOCYTES_DN
GSE9988_ANTL_TREMI_AND_LPS_VS_CTRL_TREATED_MONOCYTES_DN, GSE9988_ANTL_TREMI_AND_LPS_VS_CTRL_TREATED_MONOCYTES_DN
GSE9988_LOW_LPS_VS_CTRL_TREATED_MONOCYTE_DN, GSE9988_LOW_LPS_VS_CTRL_TREATED_MONOCYTE_DN
GSE9988_LOW_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN, GSE9988_LOW_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN
GSE21360_PRIMARY_VS_TERTIARY_MEMORY_CD8_TCELL_UP, GSE21360_PRIMARY_VS_TERTIARY_MEMORY_CD8_TCELL_UP
GSE7590_DC_VS_MONOCYTE_WITH_FCCGRIB_STIM_UP, GSE7590_DC_VS_MONOCYTE_WITH_FCCGRIB_STIM_UP
GSE4606_UNSTIM_VS_CD40L_IL2_IL5_3DAY_IRFAMID_SORTED_BCELL_UP, GSE4606_UNSTIM_VS_CD40L_IL2_IL5_3DAY_STIMULATED_IRFAMID_SORTED_BCELL_UP
GSE9988_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN, GSE9988_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN
GSE9988_ANTL_TREMI_AND_LPS_VS_VEHICLE_TREATED_MONOCYTES_DN, GSE9988_ANTL_TREMI_AND_LPS_VS_VEHICLE_TREATED_MONOCYTES_DN
GSE23695_CD57_POS_VS_NEG_NK_CELL_UP, GSE23695_CD57_POS_VS_NEG_NK_CELL_UP
GSE11864_CSF1_FNG_VS_CSF1_PAM3CYS_IN_MAC_UP, GSE11864_CSF1_FNG_VS_CSF1_PAM3CYS_IN_MAC_UP
GSE21240_HEALTHY_VS_ARTHRITIC_MOUSE_CD4_TCELL_DN, GSE21240_HEALTHY_VS_ARTHRITIC_MOUSE_CD4_TCELL_DN
GSE16755_CTRL_VS_IPNA_TREATED_MAC_UP, GSE16755_CTRL_VS_IPNA_TREATED_MAC_UP
GSE9988_ANTL_TREMI_VS_LPS_MONOCYTE_UP, GSE9988_ANTL_TREMI_VS_LPS_MONOCYTE_UP
GSE369_PRE_VS_POST_IL6_INJECTION_S0C3_KO_LIVER_DN, GSE369_PRE_VS_POST_IL6_INJECTION_S0C3_KO_LIVER_DN
GSE37416_CTRL_VS_12H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_12H_F_TULARENSIS_LVS_NEUTROPHIL_UP
GSE36527_CD69_NKG_VS_POS_TREG_CD62L_10S_KIR6G_NKG_DN, GSE36527_CD69_NKG_VS_POS_TREG_CD62L_10S_KIR6G_NKG_DN
GSE17721_PAMCSK4_VS_CPG_4H_BMDC_UP, GSE17721_PAMCSK4_VS_CPG_4H_BMDC_UP
GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_DN, GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_DN
GSE43863_TH1_VS_THF_MEMORY_CD4_TCELL_DN, GSE43863_TH1_VS_THF_MEMORY_CD4_TCELL_DN
MORF_TERF2IP, MORF_TERF2IP
GSE1930_STIM_VS_STIM_AND_IL12_48H_CD8_T_CELL_UP, GSE1930_STIM_VS_STIM_AND_IL12_48H_CD8_T_CELL_UP
GSE3585_UNTREATED_VS_IFNG_STIM_MACROPHAGE_DN, GSE3585_UNTREATED_VS_IFNG_STIM_MACROPHAGE_DN
GSE15330_WT_VS_IKAROS_KO_HSC_DN, GSE15330_WT_VS_IKAROS_KO_HSC_DN
GSE41176_WT_VS_TAK1_KO_ANTL_ICM_STIM_BCELL_24H_DN, GSE41176_WT_VS_TAK1_KO_ANTL_ICM_STIM_BCELL_24H_DN
GSE41176_UNSTIM_VS_ANTL_ICM_STIM_BCELL_3H_DN, GSE41176_UNSTIM_VS_ANTL_ICM_STIM_BCELL_3H_DN
GSE2050_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP, GSE2050_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP
GSE37533_UNTREATED_VS_PIRAZOLONE_TREATED_CD4_TCELL_PPARGI_AND_FOXO3_TRANSDUCED_DN, GSE37533_UNTREATED_VS_PIRAZOLONE_TREATED_CD4_TCELL_PPARGI_AND_FOXO3_TRANSDUCED_DN
GSE37416_CTRL_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_UP
GSE34392_ST2_KO_VS_WT_DAYS_1CMV_EFFECTOR_CD8_TCELL_UP, GSE34392_ST2_KO_VS_WT_DAYS_1CMV_EFFECTOR_CD8_TCELL_UP
GSE22601_IMMATURE_CD4_SINGLE_POSITIVE_VS_CD8_SINGLE_POSITIVE_THYMOCTYE_UP, GSE22601_IMMATURE_CD4_SINGLE_POSITIVE_VS_CD8_SINGLE_POSITIVE_THYMOCTYE_UP
GSE7348_UNSTIM_VS_LPS_STIM_MACROPHAGE_UP, GSE7348_UNSTIM_VS_LPS_STIM_MACROPHAGE_UP
GSE17721_POLYIC_VS_PAMCSK4_16H_BMDC_UP, GSE17721_POLYIC_VS_PAMCSK4_16H_BMDC_UP
GSE17721_LPS_VS_POLYIC_4H_BMDC_DN, GSE17721_LPS_VS_POLYIC_4H_BMDC_DN
GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_DN, GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_DN
GSE38681_WT_VS_LYL1_KO_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_UP, GSE38681_WT_VS_LYL1_KO_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_UP
GSE2706_UNSTIM_VS_2H_LPS_AND_R848_DC_UP, GSE2706_UNSTIM_VS_2H_LPS_AND_R848_DC_UP
GSE9960_HEALTHY_VS_GRAM_POS_SEPSIS_PPMC_UP, GSE9960_HEALTHY_VS_GRAM_POS_SEPSIS_PPMC_UP
GSE7548_NAIVE_VS_DAY17_PCC_IMMUNIZATION_CD4_TCELL_UP, GSE7548_NAIVE_VS_DAY17_PCC_IMMUNIZATION_CD4_TCELL_UP
GSE17301_IPNA2_VS_IPNA5_STIM_AC2D3_ACT_CD8_TCELL_UP, GSE17301_IPNA2_VS_IPNA5_STIM_AC2D3_ACT_CD8_TCELL_UP
GSE17721_CTRL_VS_PAMCSK4_4H_BMDC_UP, GSE17721_CTRL_VS_PAMCSK4_4H_BMDC_UP
GSE2706_UNSTIM_VS_2H_LPS_DC_UP, GSE2706_UNSTIM_VS_2H_LPS_DC_UP
MORF_BECN1, MORF_BECN1
GSE17580_TREG_VS_TIEFF_DN, GSE17580_TREG_VS_TIEFF_DN
GSE15330_WT_VS_IKAROS_KO_GRANULOCYTE_MONOCYTE_PROGENITOR_DN, GSE15330_WT_VS_IKAROS_KO_GRANULOCYTE_MONOCYTE_PROGENITOR_DN
GSE26343_WT_VS_NFAT5_KO_MACROPHAGE_LPS_STIM_UP, GSE26343_WT_VS_NFAT5_KO_MACROPHAGE_LPS_STIM_UP
SCHERER_PPMC_APSV_WETVAX_AGE_18_32YO_50_TO_60DY_DN, SCHERER_PPMC_APSV_WETVAX_AGE_18_32YO_50_TO_60DY_DN
GSE360_CTRL_VS_M_TUBERCULOSIS_MAC_UP, GSE360_CTRL_VS_M_TUBERCULOSIS_MAC_UP
GSE6674_PL2_3_VS_ANTL_ICM_AND_CPG_STIM_BCELL_UP, GSE6674_PL2_3_VS_ANTL_ICM_AND_CPG_STIM_BCELL_UP
GSE2598_IL4_VS_IL4_AND_ROSGLITAZONE_STIM_MACROPHAGE_DAY10_DN, GSE2598_IL4_VS_IL4_AND_ROSGLITAZONE_STIM_MACROPHAGE_DAY10_DN
GSE17721_LPS_VS_POLYIC_24H_BMDC_DN, GSE17721_LPS_VS_POLYIC_24H_BMDC_DN
GSE22611_UNSTIM_VS_2H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP, GSE22611_UNSTIM_VS_2H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP
DAZARD_RESPONSE_TO_UV_SCC_DN, DAZARD_RESPONSE_TO_UV_SCC_DN
GSE12198_CTRL_VS_HIGH_IL2_STIM_NK_CELL_DN, GSE12198_CTRL_VS_HIGH_IL2_STIM_NK_CELL_DN
GSE4027_POS_AND_1EFT_TRANSDUCED_VS_CTRL_CD4_TCELL_DN, GSE4027_POS_AND_1EFT_TRANSDUCED_VS_CTRL_CD4_TCELL_DN
GSE33425_CD61_HIGH_VS_NEG_CD8_TCELL_UP, GSE33425_CD61_HIGH_VS_NEG_CD8_TCELL_UP
GSE3039_CD4_TCELL_VS_ALPHAALPHA_CD8_TCELL_UP, GSE3039_CD4_TCELL_VS_ALPHAALPHA_CD8_TCELL_UP
GSE17721_POLYIC_VS_GARDIQUIMOD_16H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_16H_BMDC_UP
GSE1930_NAIVE_VS_24H_IN_VITRO_STIM_IL12_CD8_TCELL_UP, GSE1930_NAIVE_VS_24H_IN_VITRO_STIM_IL12_CD8_TCELL_UP
GSE1930_NAIVE_VS_24H_IN_VITRO_STIM_CD8_TCELL_UP, GSE1930_NAIVE_VS_24H_IN_VITRO_STIM_CD8_TCELL_UP
GSE21670_IL6_VS_TCIF_AND_IL6_TREATED_CD4_TCELL_UP, GSE21670_IL6_VS_TCIF_AND_IL6_TREATED_CD4_TCELL_UP
GSE1930_NAIVE_VS_24H_IN_VITRO_STIM_INFAB_CD8_TCELL_UP, GSE1930_NAIVE_VS_24H_IN_VITRO_STIM_INFAB_CD8_TCELL_UP
GSE17721_POLYIC_VS_GARDIQUIMOD_24H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_24H_BMDC_UP
GOBP_VESICLE_MEDIATED_TRANSPORT_TO_THE_PLASMA_MEMBRANE, GOBP_VESICLE_MEDIATED_TRANSPORT_TO_THE_PLASMA_MEMBRANE
GSE4606_UNSTIM_VS_CD40L_IL2_IL5_DAYS3_STIMULATED_BCELL_UP, GSE4606_UNSTIM_VS_CD40L_IL2_IL5_DAYS3_STIMULATED_BCELL_UP
MIR267_3P, MIR267_3P
REACTOME_CELLULAR_RESPONSE_TO_HEAT_STRESS, REACTOME_CELLULAR_RESPONSE_TO_HEAT_STRESS
MORF_CSNK1D, MORF_CSNK1D
GSE37332_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_UP, GSE37332_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_UP
GSE6259_FUT3L_INDUCED_3D1_POS_DC_VS_CD8_TCELL_UP, GSE6259_FUT3L_INDUCED_3D1_POS_DC_VS_CD8_TCELL_UP
GOCC_HISTONE_METHYLTRANSFERASE_COMPLEX, GOCC_HISTONE_METHYLTRANSFERASE_COMPLEX
GSE35685_CD4POS_CD38NEG_VS_CD34POS_CD10POS_BONE_MARROW_UP, GSE35685_CD4POS_CD38NEG_VS_CD34POS_CD10POS_BONE_MARROW_UP
GSE15330_LYMPHOID_MULTIPOTENT_VS_PRO_BCELL_UP, GSE15330_LYMPHOID_MULTIPOTENT_VS_PRO_BCELL_UP
REACTOME_TRANSPORT_OF_MATURE_MRNAS_DERIVED_FROM_INTRONLESS_TRANSCRIPTS, REACTOME_TRANSPORT_OF_MATURE_MRNAS_DERIVED_FROM_INTRONLESS_TRANSCRIPTS
REACTOME_REGULATION_OF_HSF1_MEDIATED_HEAT_SHOCK_RESPONSE, REACTOME_REGULATION_OF_HSF1_MEDIATED_HEAT_SHOCK_RESPONSE
RODRIGUES_NTN1_TARGETS_DN, RODRIGUES_NTN1_TARGETS_DN
REACTOME_SUMOYLATION_OF_RNA_BINDING_PROTEINS, REACTOME_SUMOYLATION_OF_RNA_BINDING_PROTEINS
MIR935, MIR935
HP_PARKINSONISM, HP_PARKINSONISM
GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_UP, GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_UP
BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_TRANS, BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_TRANS
MIR423_5P, MIR423_5P
REACTOME_TRANSPORT_OF_THE_SLRP_DEPENDANT_MATURE_MRNA, REACTOME_TRANSPORT_OF_THE_SLRP_DEPENDANT_MATURE_MRNA
MIR939_3P, MIR939_3P
RASHL_RESPONSE_TO_IONIZING_RADIATION_5, RASHL_RESPONSE_TO_IONIZING_RADIATION_5
MIR3184_5P, MIR3184_5P
GOME_NUCLEAR_RECEPTOR_COACTIVATOR_ACTIVITY, GOME_NUCLEAR_RECEPTOR_COACTIVATOR_ACTIVITY
YTAAYNGCT_UNKNOWN, YTAAYNGCT_UNKNOWN
AACYNNNNTTCCS_UNKNOWN, AACYNNNNTTCCS_UNKNOWN
HP_HALLUCINATIONS, HP_HALLUCINATIONS
GOBP_POSITIVE_REGULATION_OF_AUTOPHAGY, GOBP_POSITIVE_REGULATION_OF_AUTOPHAGY
GOME_GDP_BINDING_GOME_GDP_BINDING
GSE16385_IL4_VS_ROSGLITAZONE_STIM_MACROPHAGE_DN, GSE16385_IL4_VS_ROSGLITAZONE_STIM_MACROPHAGE_DN
MIR3652, MIR3652
GSE14308_TH1_VS_NAIVE_CD4_TCELL_DN, GSE14308_TH1_VS_NAIVE_CD4_TCELL_DN
GOME_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY, GOME_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY
HP_BRADYKININ, HP_BRADYKININ
MIR4718, MIR4718
GOBP_HISTONE_H3_ACETYLATION, GOBP_HISTONE_H3_ACETYLATION
GSE13306_RA_VS_UNTREATED_TREG_DN, GSE13306_RA_VS_UNTREATED_TREG_DN
HP_CONGENITAL_CONTRACTURE, HP_CONGENITAL_CONTRACTURE
MIR6829_3P, MIR6829_3P
MIR6784_3P, MIR6784_3P
MIR6791_3P, MIR6791_3P
ATGTAGC_MIR221_MIR222, ATGTAGC_MIR221_MIR222
PID_RAC1_PATHWAY, PID_RAC1_PATHWAY
MODULE_87, MODULE_87
GOBP_REGULATION_OF_VACUOLE_ORGANIZATION, GOBP_REGULATION_OF_VACUOLE_ORGANIZATION
MODULE_86, MODULE_86
HP_INAPPROPRIATE_BEHAVIOR, HP_INAPPROPRIATE_BEHAVIOR
GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_PLASMA_MEMBRANE, GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_PLASMA_MEMBRANE
GOME_SIGNALING_ADAPTOR_ACTIVITY, GOME_SIGNALING_ADAPTOR_ACTIVITY
GOME_N_ACETYLTRANSFERASE_ACTIVITY, GOME_N_ACETYLTRANSFERASE_ACTIVITY
HP_ABNORMAL_FACIAL_EXPRESSION, HP_ABNORMAL_FACIAL_EXPRESSION
HP_POSTURAL_INSTABILITY, HP_POSTURAL_INSTABILITY
HP_DECREASED_FACIAL_EXPRESSION, HP_DECREASED_FACIAL_EXPRESSION
GOBP_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY, GOBP_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY
MIR4802_3P, MIR4802_3P
BIOCARTA_TNFR1_PATHWAY, BIOCARTA_TNFR1_PATHWAY
MIR212_5P, MIR212_5P
MIR3159, MIR3159
PID_AR_PATHWAY, PID_AR_PATHWAY
HP_PROMINENT_DIGIT_PAD, HP_PROMINENT_DIGIT_PAD
PID_HIV_AIEF_PATHWAY, PID_HIV_AIEF_PATHWAY
HP_DISPROPORTIONATE_TALL_STATURE, HP_DISPROPORTIONATE_TALL_STATURE
MIR6826_3P, MIR6826_3P
GOBP_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBLY, GOBP_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBLY
MIR6889_3P, MIR6889_3P
REACTOME_REGULATION_OF_TP53_ACTIVITY_THROUGH_ACETYLATION, REACTOME_REGULATION_OF_TP53_ACTIVITY_THROUGH_ACETYLATION
HP_ARTHROGRIPOSY_MULTIPLEX_CONGENITA, HP_ARTHROGRIPOSY_MULTIPLEX_CONGENITA
REACTOME_SLC_TRANSPORTER_DISORDERS, REACTOME_SLC_TRANSPORTER_DISORDERS
MIR2158_5P, MIR2158_5P
HP_RESTING_TREMOR, HP_RESTING_TREMOR
WP_CHEMOKINE_SIGNALING_PATHWAY, WP_CHEMOKINE_SIGNALING_PATHWAY
L1L1_THYROID_CANCER_CLUSTER_1, L1L1_THYROID_CANCER_CLUSTER_1
REACTOME_ONCOGENIC_MAPK_SIGNALING, REACTOME_ONCOGENIC_MAPK_SIGNALING
MIR4725_5P, MIR4725_5P
GSE27786_NKCELL_VS_NEUTROPHIL_DN, GSE27786_NKCELL_VS_NEUTROPHIL_DN
PID_P38_ALPHA_BETA_PATHWAY, PID_P38_ALPHA_BETA_PATHWAY
GOBP_ENDOCYTIC_RECYCLING, GOBP_ENDOCYTIC_RECYCLING
GOME_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY, GOME_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY
GOME_N_ACETYLTRANSFERASE_ACTIVITY, GOME_N_ACETYLTRANSFERASE_ACTIVITY
MIR5704, MIR5704
PID_TNF_PATHWAY, PID_TNF_PATHWAY
GOBP_GOLGI_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT, GOBP_GOLGI_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT
DAVIES_MULTIPLE_MYELOMA_VS_MGUS_DN, DAVIES_MULTIPLE_MYELOMA_VS_MGUS_DN
REACTOME_TGF_BETA_RECEPTOR_SIGNALING_ACTIVATES_SMAADS, REACTOME_TGF_BETA_RECEPTOR_SIGNALING_ACTIVATES_SMAADS
GOME_SIGNALING_RECEPTOR_COMPLEX_ADAPTOR_ACTIVITY, GOME_SIGNALING_RECEPTOR_COMPLEX_ADAPTOR_ACTIVITY
PID_TGFBF_PATHWAY, PID_TGFBF_PATHWAY
REACTOME_SIGNALING_TO_ERKS, REACTOME_SIGNALING_TO_ERKS
PID_FAS_PATHWAY, PID_FAS_PATHWAY
WP_TNF_RELATED_WEAK_INDUCER_OF_APOPTOSIS_TWEAK_SIGNALING_PATHWAY, WP_TNF_RELATED_WEAK_INDUCER_OF_APOPTOSIS_TWEAK_SIGNALING_PATHWAY
GSE43863_TH1_VS_LY6C_INT_CXCR5POS_MEMORY_CD4_TCELL_DN, GSE43863_TH1_VS_LY6C_INT_CXCR5POS_MEMORY_CD4_TCELL_DN
BIOCARTA_DEATH_PATHWAY, BIOCARTA_DEATH_PATHWAY
GOBP_AUTOPHAGOSOME_MATURATION, GOBP_AUTOPHAGOSOME_MATURATION
GOBP_NEGATIVE_REGULATION_OF_INNATE_IMMUNE_RESPONSE, GOBP_NEGATIVE_REGULATION_OF_INNATE_IMMUNE_RESPONSE
GENTILE_UV_RESPONSE_CLUSTER_D5, GENTILE_UV_RESPONSE_CLUSTER_D5
HP_HYPOMIMIC_FACE, HP_HYPOMIMIC_FACE
HP_NEURONAL_LOSS_IN_CENTRAL_NERVOUS_SYSTEM, HP_NEURONAL_LOSS_IN_CENTRAL_NERVOUS_SYSTEM
GOBP_STRESS_FIBER_ASSEMBLY, GOBP_STRESS_FIBER_ASSEMBLY
CN2_MRN, CN2_MRN
HP_ABNORMAL_SPEECH_PROSODY, HP_ABNORMAL_SPEECH_PROSODY
BCAT100_UPI_V1, BCAT100_UPI_V1
HP_NASOGASTRIC_TUBE_FEEDING_IN_INFANCY, HP_NASOGASTRIC_TUBE_FEEDING_IN_INFANCY
MIR4749_3P, MIR4749_3P
MIR3174, MIR3174
MYLLKANGAS_AMPLIFICATION_HOT_SPOT_23, MYLLKANGAS_AMPLIFICATION_HOT_SPOT_23
GOME_PROTEIN_SERINE_PHOSPHATASE_ACTIVITY, GOME_PROTEIN_SERINE_PHOSPHATASE_ACTIVITY
GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_UP, GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_UP
GOBP_NEGATIVE_REGULATION_OF_MAPK_CASCADE, GOBP_NEGATIVE_REGULATION_OF_MAPK_CASCADE
ZNF746_TARGET_GENES, ZNF746_TARGET_GENES
GOCC_SYNAPTIC_VESICLE_MEMBRANE, GOCC_SYNAPTIC_VESICLE_MEMBRANE
HP_ASTROCYTOSIS, HP_ASTROCYTOSIS
REACTOME_DOWNREGULATION_OF_TGF_BETA_RECEPTOR_SIGNALING, REACTOME_DOWNREGULATION_OF_TGF_BETA_RECEPTOR_SIGNALING
CHNG_MULTIPLE_MYELOMA_HYPERPLOID_DN, CHNG_MULTIPLE_MYELOMA_HYPERPLOID_DN
REACTOME_MAPK_TARGETS_NUCLEAR_EVENTS_MEDIATED_BY_MAP_KINASES, REACTOME_MAPK_TARGETS_NUCLEAR_EVENTS_MEDIATED_BY_MAP_KINASES
GOCC_GCC, MIR296, GOCC_GCC, MIR296
GOBP_RETROGRADE_TRANSPORT_ENDOSOME_TO_PLASMA_MEMBRANE, GOBP_RETROGRADE_TRANSPORT_ENDOSOME_TO_PLASMA_MEMBRANE
GOBP_HISTONE_DEUBIQUITINATION, GOBP_HISTONE_DEUBIQUITINATION
REACTOME_RAF_INDEPENDENT_MAPK1_3_ACTIVATION, REACTOME_RAF_INDEPENDENT_MAPK1_3_ACTIVATION
BIOCARTA_TOLL_PATHWAY, BIOCARTA_TOLL_PATHWAY
GSE1569_NONSUPPRESSIVE_TCELL_VS_ACTIVATED_TREG_UP, GSE1569_NONSUPPRESSIVE_TCELL_VS_ACTIVATED_TREG_UP
GOCC_SAGA_TYPE_COMPLEX, GOCC_SAGA_TYPE_COMPLEX
PID_AVB3_INTEGRIN_PATHWAY, PID_AVB3_INTEGRIN_PATHWAY
REACTOME_BETA_CATENIN_PHOSPHORYLATION_CASCADE, REACTOME_BETA_CATENIN_PHOSPHORYLATION_CASCADE
HP_HAND_CLENCHING, HP_HAND_CLENCHING
HP_CEREBRAL_INCLUSION_BODIES, HP_CEREBRAL_INCLUSION_BODIES
GOME_MYOSIN_BINDING_GOME_MYOSIN_BINDING
REACTOME_TNFR1_INDUCED_PROAPOPTOTIC_SIGNALING, REACTOME_TNFR1_INDUCED_PROAPOPTOTIC_SIGNALING
GOCC_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE, GOCC_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE
GOBP_POSITIVE_REGULATION_OF_ERBB_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_ERBB_SIGNALING_PATHWAY
PID_ALK1_PATHWAY, PID_ALK1_PATHWAY
GOBP_CELLULAR_RESPONSE_TO_MECHANICAL_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_MECHANICAL_STIMULUS
BIOCARTA_TALL1_PATHWAY, BIOCARTA_TALL1_PATHWAY
BIOCARTA_STRESS_PATHWAY, BIOCARTA_STRESS_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GOBP_NEGATIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
GOBP_PROTEIN_LOCALIZATION_TO_ENDOSOME, GOBP_PROTEIN_LOCALIZATION_TO_ENDOSOME
HP_INAPPROPRIATE_LAUGHTER, HP_INAPPROPRIATE_LAUGHTER
HP_DIMINISHED_MOVEMENT, HP_DIMINISHED_MOVEMENT
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MEMBRANE, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MEMBRANE
MAGRANGEAS_MULTIPLE_MYELOMA_JGG_VS_JGA_DN, MAGRANGEAS_MULTIPLE_MYELOMA_JGG_VS_JGA_DN
GOBP_PEPITIDYL_TYROSINE_DEPHOSPHORYLATION, GOBP_PEPITIDYL_TYROSINE_DEPHOSPHORYLATION
HP_HAPPY_DEMANOR, HP_HAPPY_DEMANOR
KYNG_DNA_DAMAGE_BY_GAMMA_AND_UV_RADIATION, KYNG_DNA_DAMAGE_BY_GAMMA_AND_UV_RADIATION
HP_VISUAL_HALLUCINATIONS, HP_VISUAL_HALLUCINATIONS
MIR1193, MIR1193
REACTOME_NEGATIVE_REGULATION_OF_FLT3, REACTOME_NEGATIVE_REGULATION_OF_FLT3
GOME_PROTEIN_TYROSINE_PHOSPHATASE_ACTIVITY, GOME_PROTEIN_TYROSINE_PHOSPHATASE_ACTIVITY
GOBP_DENDRITE_EXTENSION, GOBP_DENDRITE_EXTENSION
GRADE_COLO_N_VS_RECTAL_CANCER_DN, GRADE_COLO_N_VS_RECTAL_CANCER_DN
GOBP_POSITIVE_REGULATION_OF_ALPHA_BETA_T_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_ALPHA_BETA_T_CELL_PROLIFERATION
MIR8055, MIR8055
MIR227_3P, MIR227_3P
GOBP_TOLL_LIKE_RECEPTOR_3_SIGNALING_PATHWAY, GOBP_TOLL_LIKE_RECEPTOR_3_SIGNALING_PATHWAY
GOME_PHOSPHOLIPID_TRANSFER_ACTIVITY, GOME_PHOSPHOLIPID_TRANSFER_ACTIVITY
HP_PINEAL_CYST, HP_PINEAL_CYST
BIOCARTA_CCR5_PATHWAY, BIOCARTA_CCR5_PATHWAY
REACTOME_DISASSEMBLY_OF_THE_DESTRUCTION_COMPLEX_AND_RECRUITMENT_OF_AXIN_TO_THE_MEMBRANE, REACTOME_DISASSEMBLY_OF_THE_DESTRUCTION_COMPLEX_AND_RECRUITMENT_OF_AXIN_TO_THE_MEMBRANE
GOBP_REGULATION_OF_LYSOSOMAL_LUMEN_PH, GOBP_REGULATION_OF_LYSOSOMAL_LUMEN_PH
REACTOME_CHEMOKINE_RECEPTORS_BIND_CHEMOKINES, REACTOME_CHEMOKINE_RECEPTORS_BIND_CHEMOKINES
NIKOLSKY_BREAST_CANCER_22Q13_AMPlicON, NIKOLSKY_BREAST_CANCER_22Q13_AMPlicON
GOBP_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS, GOBP_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS
MIR4763_5P, MIR4763_5P
REACTOME_SIGNALING_TO_RAS, REACTOME_SIGNALING_TO_RAS
GOBP_SEQUESTERING_OF_TRIGLYCERIDE, GOBP_SEQUESTERING_OF_TRIGLYCERIDE
GOME_MITOGEN_ACTIVATED_PROTEIN_KINASE_BINDING, GOME_MITOGEN_ACTIVATED_PROTEIN_KINASE_BINDING
GOME_G_PROTEIN_ALPHA_SUBUNIT_BINDING, GOME_G_PROTEIN_ALPHA_SUBUNIT_BINDING
KEGG_RIG_1_LIKE_RECEPTOR_SIGNALING_PATHWAY, KEGG_RIG_1_LIKE_RECEPTOR_SIGNALING_PATHWAY
WP_HYPOTHESIZED_PATHWAYS_IN_PATHOGENESIS_OF_CARDIOVASCULAR_DISEASE, WP_HYPOTHESIZED_PATHWAYS_IN_PATHOGENESIS_OF_CARDIOVASCULAR_DISEASE
MORI_PLASMA_CELL_DN, MORI_PLASMA_CELL_DN
MIR2114_3P, MIR2114_3P
GOBP_ACTIVATION_OF_CYSINE_TYPE_ENDOPEPTIDASE_ACTIVITY, GOBP_ACTIVATION_OF_CYSINE_TYPE_ENDOPEPTIDASE_ACTIVITY
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY
HP_CONGENITAL_NYSTAGMUS, HP_CONGENITAL_NYSTAGMUS
HP_PARKINSONISM_WITH_FAVORABLE_RESPONSE_TO_DOPAMINERGIC_MEDICATION, HP_PARKINSONISM_WITH_FAVORABLE_RESPONSE_TO_DOPAMINERGIC_MEDICATION
GOBP_VISUAL_BEHAVIOR, GOBP_VISUAL_BEHAVIOR
HP_PULMONARY_ARTERY_DILATATION, HP_PULMONARY_ARTERY_DILATATION
GENTLES_LEUKEMIC_STEM_CELL_UP, GENTLES_LEUKEMIC_STEM_CELL_UP
HOFMANN_MYELODYSPLASTIC_SYNDROM_RISK_DN, HOFMANN_MYELODYSPLASTIC_SYNDROM_RISK_DN
GOBP_T_CELL_MIGRATION, GOBP_T_CELL_MIGRATION
MIR472_3P, MIR472_3P
YRCCAKNNNGCCG_UNKNOWN, YRCCAKNNNGCCG_UNKNOWN
GOCC_BORC_COMPLEX, GOCC_BORC_COMPLEX
REACTOME_SIGNALING_BY_MODERATE_KINASE_ACTIVITY_BRAF_MUTANTS, REACTOME_SIGNALING_BY_MODERATE_KINASE_ACTIVITY_BRAF_MUTANTS
GOBP_POSITIVE_REGULATION_OF_VACUOLE_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_VACUOLE_ORGANIZATION
REACTOME_PRRAPK_EVENTS, REACTOME_PRRAPK_EVENTS
GOBP_BETA_CATENIN_DESTRUCTION_COMPLEX_DISASSEMBLY, GOBP_BETA_CATENIN_DESTRUCTION_COMPLEX_DISASSEMBLY
GOBP_REGULATION_OF_ORGAN_GROWTH, GOBP_REGULATION_OF_ORGAN_GROWTH
HP_PULMONARY_HEMORRHAGE, HP_PULMONARY_HEMORRHAGE
HP_MOTOR_APHASIA, HP_MOTOR_APHASIA
TTCGTT_MIR91, TTCGTT_MIR91
HP_CUTANEOUS_MELANOMA, HP_CUTANEOUS_MELANOMA
MIR550A_3P, MIR550A_3P
BIOCARTA_NFKB_PATHWAY, BIOCARTA_NFKB_PATHWAY
REACTOME_ERK_MAPK_TARGETS, REACTOME_ERK_MAPK_TARGETS
GOBP_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE, GOBP_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE
GOBP_T_CELL_APOPTOTIC_PROCESS, GOBP_T_CELL_APOPTOTIC_PROCESS
REACTOME_PTK6_REGULATES_RTKS_AND_THEIR_EFFECTORS_AKT1_AND_DOK1, REACTOME_PTK6_REGULATES_RTKS_AND_THEIR_EFFECTORS_AKT1_AND_DOK1
MODULE_524, MODULE_524
BIOCARTA_CTCE_PATHWAY, BIOCARTA_CTCE_PATHWAY
HP_VITILIGO, HP_VITILIGO
BACOLOD_RESISTANCE_TO_ALKYLATING_AGENTS_UP, BACOLOD_RESISTANCE_TO_ALKYLATING_AGENTS_UP
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_PERIPHERY, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_PERIPHERY
GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT, GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT
GOBP_REGULATION_OF_GOLGI_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT, GOBP_REGULATION_OF_GOLGI_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT
GOBP_REGULATION_OF_BONE_REMODELING, GOBP_REGULATION_OF_BONE_REMODELING
CHIRGA, CHIRGA
MIR1180_5P, MIR1180_5P
REACTOME_NEUROTRANSMITTER_RELEASE_CYCLE, REACTOME_NEUROTRANSMITTER_RELEASE_CYCLE
REACTOME_TRAIL_SIGNALING, REACTOME_TRAIL_SIGNALING
GOBP_RESPONSE_TO_OXYGEN_GLUCOSE_DEPRIVATION, GOBP_RESPONSE_TO_OXYGEN_GLUCOSE_DEPRIVATION
HP_PROMINENT_NASOLABIAL_FOLD, HP_PROMINENT_NASOLABIAL_FOLD
GOBP_NEGATIVE_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT
WP_CANNABINOID_RECEPTOR_SIGNALING, WP_CANNABINOID_RECEPTOR_SIGNALING
KIM_LIVER_CANCER_POOR_SURVIVAL_UP, KIM_LIVER_CANCER_POOR_SURVIVAL_UP
BIOCARTA_EGFR_SMRTK_PATHWAY, BIOCARTA_EGFR_SMRTK_PATHWAY
HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_ICG_VACCINE_AGE_18_45YO_56D_TOP_100_DEG_AFTER_IN_VITRO_RE_STIMULATION_UP, HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_ICG_VACCINE_AGE_18_45YO_56D_TOP_100_DEG_AFTER_IN_VITRO_RE_STIMULATION_UP
GOBP_INACTIVATION_OF_MAPK_ACTIVITY, GOBP_INACTIVATION_OF_MAPK_ACTIVITY
GOBP_REGULATION_OF_LIPID_STORAGE, GOBP_REGULATION_OF_LIPID_STORAGE
GOBP_LYSOSOMAL_LUMEN_ACIDIFICATION, GOBP_LYSOSOMAL_LUMEN_ACIDIFICATION
GOCC_SCAR_COMPLEX, GOCC_SCAR_COMPLEX
GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CENTROSOME, GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CENTROSOME