

G_M_RAB10, GCM_RAB10

GCM_MYST2, GCM_MYST2
GCM_MLL, GCM_MLL
GCM_UBE2N, GCM_UBE2N
GCM_DFFA, GCM_DFFA
GCM_GSPT1, GCM_GSPT1
GCM_RAN, GCM_RAN
GCM_ZNF198, GCM_ZNF198
GSE13484_12H_UNSTIM_VS_YF17D_VACCINE_STIM_PBMCM_UP, GSE13484_12H_UNSTIM_VS_YF17D_VACCINE_STIM_PBMCM_UP
GCM_MAP4K4, GCM_MAP4K4
GSE13229_MATURE_VS_INTMATURE_NKCELL_DN, GSE13229_MATURE_VS_INTMATURE_NKCELL_DN
GCM_RBM8A, GCM_RBM8A
GSE21546_ELK1_KO_VS_SAP1A_KO_AND_ELK1_KO_ANTI_CD3_STIM_DP_THYMOCYTES_UP, GSE21546_ELK1_KO_VS_SAP1A_KO_AND_ELK1_KO_ANTI_CD3_STIM_DP_THYMOCYTES_UP
GCM_HBP1, GCM_HBP1
GSE17721_LPS_VS_GARDIQUIMOD_16H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_16H_BMDC_DN
GCM_PTK2, GCM_PTK2
GSE17721_CPG_VS_GARDIQUIMOD_8H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_8H_BMDC_DN
GCM_NCAM1, GCM_NCAM1
GCM_SUFU, GCM_SUFU
GCM_CRKL, GCM_CRKL
GO_ER_ASSOCIATED_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_ER_ASSOCIATED_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
MODULE_323, MODULE_323
GCM_BMPR2, GCM_BMPR2
GCM_ERBB2IP, GCM_ERBB2IP
ATACCTC_MIR202, ATACCTC_MIR202
GSE33292_WT_VS_TCF1_KO_DN3_THYMOCYTE_UP, GSE33292_WT_VS_TCF1_KO_DN3_THYMOCYTE_UP
GO_ERAD_PATHWAY, GO_ERAD_PATHWAY
GSE39556_CD8A_DC_VS_NK_CELL_MOUSE_3H_POST_POLYIC_INJ_DN, GSE39556_CD8A_DC_VS_NK_CELL_MOUSE_3H_POST_POLYIC_INJ_DN
GGGCATT_MIR365, GGCATT_MIR365
GSE29617_DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMCM_2008_DN, GSE29617_DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMCM_2008_DN
GSE4748_CTRL_VS_LPS_STIM_DC_3H_UP, GSE4748_CTRL_VS_LPS_STIM_DC_3H_UP
GSE8515_CTRL_VS_IL6_4H_STIM_MAC_UP, GSE8515_CTRL_VS_IL6_4H_STIM_MAC_UP
BYSTROEM_CORRELATED_WITH_IL5_DN, BYSTROEM_CORRELATED_WITH_IL5_DN
GSE7548_DAY7_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP, GSE7548_DAY7_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP
GCM_NUMA1, GCM_NUMA1
GCM_DLG1, GCM_DLG1
GCM_BAG5, GCM_BAG5
ACTGCAG_MIR173P, ACTGCAG_MIR173P
RAMASWAMY_METASTASIS_UP, RAMASWAMY_METASTASIS_UP
GCM_CSNK1A1, GCM_CSNK1A1
GCM_IL6ST, GCM_IL6ST
GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_6H_UP
ATGTTTC_MIR494, ATGTTTC_MIR494
GO_GOLGI_TO_VACUOLE_TRANSPORT, GO_GOLGI_TO_VACUOLE_TRANSPORT
BOYALT_LIVER_CANCER_SUBCLASS_G1_UP, BOYALT_LIVER_CANCER_SUBCLASS_G1_UP
GCM_CSNK1D, GCM_CSNK1D
GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_UP, GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_UP
GO_GDP_BINDING, GO_GDP_BINDING
AAAGACA_MIR511, AAAGACA_MIR511
GCM_DEAF1, GCM_DEAF1
GSE3982_EOSINOPHIL_VS_TH1_UP, GSE3982_EOSINOPHIL_VS_TH1_UP
GCTGAGT_MIR5125P, GCTGAGT_MIR5125P
GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP, GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP
GSE10500_ARTHRITIC_SYNOVIAL_FLUID_VS_HEALTHY_MACROPHAGE_DN, GSE10500_ARTHRITIC_SYNOVIAL_FLUID_VS_HEALTHY_MACROPHAGE_DN
GCM_RAP2A, GCM_RAP2A
MODULE_233, MODULE_233
CAATGCA_MIR33, CAATGCA_MIR33
GCM_FANCL, GCM_FANCL
JAK2_DN.V1_DN, JAK2_DN.V1_DN
AGTCAGC_MIR345, AGTCAGC_MIR345
GO_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI, GO_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI
GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN, GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN
GCM_SIRT2, GCM_SIRT2
GO_PROTEIN_K48_LINKED_DEUBIQUITINATION, GO_PROTEIN_K48_LINKED_DEUBIQUITINATION
GO_CELL_DIVISION_SITE, GO_CELL_DIVISION_SITE
GSE25123_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DN, GSE25123_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DN
BARRIER_COLON_CANCER_RECURRENCE_DN, BARRIER_COLON_CANCER_RECURRENCE_DN
GSE24210_IL35_TREATED_VS_UNTREATED_TCONV_CD4_TCELL_DN, GSE24210_IL35_TREATED_VS_UNTREATED_TCONV_CD4_TCELL_DN
CCACACA_MIR147, CCACACA_MIR147
GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMCM_DN, GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMCM_DN
GCM_PTPRD, GCM_PTPRD
GO_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT, GO_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT
ZIRN_TRETINOIN_RESPONSE_WT1_UP, ZIRN_TRETINOIN_RESPONSE_WT1_UP
GSE17974_0H_VS_0.5H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_0.5H_IN_VITRO_ACT_CD4_TCELL_DN
GTTATAT_MIR410, GTTATAT_MIR410
FXR_IR1_Q6, FXR_IR1_Q6
GO_REGULATION_OF_VACUOLE_ORGANIZATION, GO_REGULATION_OF_VACUOLE_ORGANIZATION
YRTCANNRCCG_UNKNOWN, YRTCANNRCCG_UNKNOWN
GO_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY, GO_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY
GO_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
chr1p22, chr1p22
WOTTON_RUNX_TARGETS_DN, WOTTON_RUNX_TARGETS_DN
chr14q22, chr14q22
GCM_MAP1B, GCM_MAP1B
GO_N_TERMINAL_PROTEIN_AMINO_ACID_MODIFICATION, GO_N_TERMINAL_PROTEIN_AMINO_ACID_MODIFICATION
GO_GOLGI_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT, GO_GOLGI_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT
GSE17721_PAM3CSK4_VS_GADIQUIMOD_4H_BMDC_UP, GSE17721_PAM3CSK4_VS_GADIQUIMOD_4H_BMDC_UP
RASHI_RESPONSE_TO_IONIZING_RADIATION_3, RASHI_RESPONSE_TO_IONIZING_RADIATION_3
GSE29615_CTRL_VS_DAY3_LAIV_IFLU_VACCINE_PBMCM_DN, GSE29615_CTRL_VS_DAY3_LAIV_IFLU_VACCINE_PBMCM_DN
GO_ACTIN_CYTOSKELETON_REORGANIZATION, GO_ACTIN_CYTOSKELETON_REORGANIZATION
GO_REGULATION_OF_ERAD_PATHWAY, GO_REGULATION_OF_ERAD_PATHWAY
GO_MORPHOGENESIS_OF_A_POLARIZED_EPITHELIUM, GO_MORPHOGENESIS_OF_A_POLARIZED_EPITHELIUM
GO_PROTEIN_NEDDYLATION, GO_PROTEIN_NEDDYLATION
GO_ASYMMETRIC_PROTEIN_LOCALIZATION, GO_ASYMMETRIC_PROTEIN_LOCALIZATION
GO_RETROMER_COMPLEX, GO_RETROMER_COMPLEX
GSE34156_TLR1_TLR2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_DN, GSE34156_TLR1_TLR2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_DN
FIGUEROA_AML_METHYLATION_CLUSTER_1_DN, FIGUEROA_AML_METHYLATION_CLUSTER_1_DN
KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_UV_IN_OLD, KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_UV_IN_OLD
AGGTGCA_MIR500, AGGTGCA_MIR500
AACGGTT_MIR451, AACGGTT_MIR451
GO_THIOLESTER_HYDROLASE_ACTIVITY, GO_THIOLESTER_HYDROLASE_ACTIVITY
ABDELMOHSEN_ELAVL4_TARGETS, ABDELMOHSEN_ELAVL4_TARGETS
EIF4E_DN, EIF4E_DN
REACTOME_GLYCEROPHOSPHOLIPID_BIOSYNTHESIS, REACTOME_GLYCEROPHOSPHOLIPID_BIOSYNTHESIS
GO_SCF_UBIQUITIN_LIGASE_COMPLEX, GO_SCF_UBIQUITIN_LIGASE_COMPLEX
GO_ESTABLISHMENT_OR_MAINTENANCE_OF_EPITHELIAL_CELL_APICAL_BASAL_POLARITY, GO_ESTABLISHMENT_OR_MAINTENANCE_OF_EPITHELIAL_CELL_APICAL_BASAL_POLARITY
GSE13411_NAIVE_VS_MEMORY_BCELL_UP, GSE13411_NAIVE_VS_MEMORY_BCELL_UP
GO_UBIQUITIN_UBIQUITIN_LIGASE_ACTIVITY, GO_UBIQUITIN_UBIQUITIN_LIGASE_ACTIVITY
chr19q12, chr19q12
GSE22611_NOD2_TRANSD_VS_CTRL_TRANSD_HEK293_MDP_STIM_2H_UP, GSE22611_NOD2_TRANSD_VS_CTRL_TRANSD_HEK293_MDP_STIM_2H_UP
GO_POLYOL_BIOSYNTHETIC_PROCESS, GO_POLYOL_BIOSYNTHETIC_PROCESS
GO_POSITIVE_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION, GO_POSITIVE_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION
APPIERTO_RESPONSE_TO_FENRETINIDE_UP, APPIERTO_RESPONSE_TO_FENRETINIDE_UP
GO_PROTEIN_DEMANNOSYLATION, GO_PROTEIN_DEMANNOSYLATION
GO_MICROVILLUS_ORGANIZATION, GO_MICROVILLUS_ORGANIZATION
RAO_BOUND_BY_SALL4_ISOFORM_A, RAO_BOUND_BY_SALL4_ISOFORM_A
MORF_ESR1, MORF_ESR1
GO_CELL_CORTEX_PART, GO_CELL_CORTEX_PART
FINETTI_BREAST_CANCERS_KINOME_BLUE, FINETTI_BREAST_CANCERS_KINOME_BLUE
REACTOME_IRAK2_MEDIATED_ACTIVATION_OF_TAK1_COMPLEX_UPON_TLR7_8_OR_9_STIMULATION, REACTOME_IRAK2_MEDIATED_ACTIVATION_OF_TAK1_COMPLEX_UPON_TLR7_8_OR_9_STIMULATION
GO_MYOSIN_V_BINDING, GO_MYOSIN_V_BINDING
chr12q14, chr12q14
chr13q34, chr13q34
GO_APICAL_PROTEIN_LOCALIZATION, GO_APICAL_PROTEIN_LOCALIZATION
GO_NEGATIVE_REGULATION_OF_NF_KAPPAB_IMPORT_INTO_NUCLEUS, GO_NEGATIVE_REGULATION_OF_NF_KAPPAB_IMPORT_INTO_NUCLEUS
HOWLIN_CITED1_TARGETS_1_DN, HOWLIN_CITED1_TARGETS_1_DN
GTAAAG_MIR302B, GTAAAG_MIR302B
GO_REGULATION_OF_TRANSPORTER_ACTIVITY, GO_REGULATION_OF_TRANSPORTER_ACTIVITY