

GSE9006_TYPE_1_VS_TYPE_2_DIABETES_PBM_AT_DX_UP, GSE9006_TYPE_1_VS_TYPE_2_DIABETES_PBM_AT_DX_UP
GSE16450_CTRL_VS_IFNA_6H_STIM_MATURE_NEURON_CELL_LINE_DN, GSE16450_CTRL_VS_IFNA_6H_STIM_MATURE_NEURON_CELL_LINE_DN
GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBM_1MONTH_POST_DX_UP, GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBM_1MONTH_POST_DX_UP
GSE9006_TYPE_1_DIABETES_AT_DX_VS_1MONTH_POST_DX_PBMC_UP, GSE9006_TYPE_1_DIABETES_AT_DX_VS_1MONTH_POST_DX_PBMC_UP
GSE9601_UNTREATED_VS_P13K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_UP, GSE9601_UNTREATED_VS_P13K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_UP
GSE45365_WT_VS_IFNAR_KO_CD8A_DC_MCMV_INFECTION_UP, GSE45365_WT_VS_IFNAR_KO_CD8A_DC_MCMV_INFECTION_UP
GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_STAT3_KO_CD4_TCELL_DN, GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_STAT3_KO_CD4_TCELL_DN
HALLMARK_DNA_REPAIR, HALLMARK_DNA_REPAIR
GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_PDC_DN, GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_PDC_DN
GSE21670_TGFB_VS_TGFB_AND_IL6_TREATED_CD4_TCELL_DN, GSE21670_TGFB_VS_TGFB_AND_IL6_TREATED_CD4_TCELL_DN
GSE9006_TYPE_1_DIABETES_AT_DX_VS_4MONTH_POST_DX_PBMC_UP, GSE9006_TYPE_1_DIABETES_AT_DX_VS_4MONTH_POST_DX_PBMC_UP
GNF2_STAT6, GNF2_STAT6
GSE17721_0.5H_VS_4H_PAM3CSK4_BMDC_UP, GSE17721_0.5H_VS_4H_PAM3CSK4_BMDC_UP
GSE22886_NAIVE_CD4_TCELL_VS_12H_ACT_TH1_UP, GSE22886_NAIVE_CD4_TCELL_VS_12H_ACT_TH1_UP
GSE16522_ANTL_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_DN, GSE16522_ANTL_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_DN
GSE20715_WT_VS_TLR4_KO_48H_OZONE_LUNG_DN, GSE20715_WT_VS_TLR4_KO_48H_OZONE_LUNG_DN
GSE360_CTRL_VS_T_GONDIL_MAC_UP, GSE360_CTRL_VS_T_GONDIL_MAC_UP
GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBM_AT_DX_UP, GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBM_AT_DX_UP
GSE22886_TCELL_VS_BCELL_NAIVE_UP, GSE22886_TCELL_VS_BCELL_NAIVE_UP
MODULE_257, MODULE_257
GSE25123_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_UP, GSE25123_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_UP
GSE7831_UNSTIM_VS_CPG_STIM_PDC_1H_UP, GSE7831_UNSTIM_VS_CPG_STIM_PDC_1H_UP
GSE6269_HEALTHY_VS_FLU_INF_PBMC_DN, GSE6269_HEALTHY_VS_FLU_INF_PBMC_DN
GSE22886_CD4_TCELL_VS_BCELL_NAIVE_UP, GSE22886_CD4_TCELL_VS_BCELL_NAIVE_UP
GSE22886_NAIVE_TCELL_VS_DC_UP, GSE22886_NAIVE_TCELL_VS_DC_UP
GSE17721_ALL_VS_24H_PAM3CSK4_BMDC_UP, GSE17721_ALL_VS_24H_PAM3CSK4_BMDC_UP
GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_MONOCYTE_UP, GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_MONOCYTE_UP
GNF2_MYD88, GNF2_MYD88
GSE1432_CTRL_VS_IFNG_1H_MICROGLIA_UP, GSE1432_CTRL_VS_IFNG_1H_MICROGLIA_UP
JOHNSTONE_PARVB_TARGETS_2_UP, JOHNSTONE_PARVB_TARGETS_2_UP
GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_DN, GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_DN
GO_CORE_PROMOTER_BINDING, GO_CORE_PROMOTER_BINDING
GSE22886_IGG_IGA_MEMORY_BCELL_VS_BM_PLASMA_CELL_DN, GSE22886_IGG_IGA_MEMORY_BCELL_VS_BM_PLASMA_CELL_DN
REACTOME_RNA_POL_II_PRE_TRANSCRIPTION_EVENTS, REACTOME_RNA_POL_II_PRE_TRANSCRIPTION_EVENTS
GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_MAC_UP, GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_MAC_UP
REACTOME_FORMATION_OF_RNA_POL_II_ELONGATION_COMPLEX, REACTOME_FORMATION_OF_RNA_POL_II_ELONGATION_COMPLEX
GSE17721_CTRL_VS_GARDIQUIMOD_1H_BMDC_UP, GSE17721_CTRL_VS_GARDIQUIMOD_1H_BMDC_UP
GNF2_CD97, GNF2_CD97
GSE43863_NAIVE_VS_Ly6C_LOW_CXCR5NEG_CD4_EFF_TCELL_D6_LCMV_DN, GSE43863_NAIVE_VS_Ly6C_LOW_CXCR5NEG_CD4_EFF_TCELL_D6_LCMV_DN
GSE10273_HIGH_IL7_VS_HIGH_IL7_AND_IRF4_IN_IRF4_8_NULL_PRE_BCELL_DN, GSE10273_HIGH_IL7_VS_HIGH_IL7_AND_IRF4_IN_IRF4_8_NULL_PRE_BCELL_DN
PENG_RAPAMYCIN_RESPONSE_UP, PENG_RAPAMYCIN_RESPONSE_UP
PECE_MAMMARY_STEM_CELL_UP, PECE_MAMMARY_STEM_CELL_UP
GSE3039_ALPHAALPHA_CD8_TCELL_VS_B2_BCELL_UP, GSE3039_ALPHAALPHA_CD8_TCELL_VS_B2_BCELL_UP
GSE18804_SPLEEN_MACROPHAGE_VS_COLON_TUMORAL_MACROPHAGE_UP, GSE18804_SPLEEN_MACROPHAGE_VS_COLON_TUMORAL_MACROPHAGE_UP
GSE1460_CD4_THYMOCYTE_VS_THYMIC_STROMAL_CELL_UP, GSE1460_CD4_THYMOCYTE_VS_THYMIC_STROMAL_CELL_UP
WOOD_EBV_EBNA1_TARGETS_UP, WOOD_EBV_EBNA1_TARGETS_UP
GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_24H_DN, GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_24H_DN
UEDA_CENTRAL_CLOCK, UEDA_CENTRAL_CLOCK
GSE17721_PAM3CSK4_VS_GADIQUIMOD_0.5H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_0.5H_BMDC_DN
GO_HEAT_SHOCK_PROTEIN_BINDING, GO_HEAT_SHOCK_PROTEIN_BINDING
GO_PURINE_NUCLEOSIDE_BIOSYNTHETIC_PROCESS, GO_PURINE_NUCLEOSIDE_BIOSYNTHETIC_PROCESS
GO_RAS_PROTEIN_SIGNAL_TRANSDUCTION, GO_RAS_PROTEIN_SIGNAL_TRANSDUCTION
GSE1448_CTRL_VS_ANTL_VALPHA2_DP_THYMOCYTE_UP, GSE1448_CTRL_VS_ANTL_VALPHA2_DP_THYMOCYTE_UP
GSE32533_MIR17_KO_VS_MIR17_OVEREXPRESS_ACT_CD4_TCELL_UP, GSE32533_MIR17_KO_VS_MIR17_OVEREXPRESS_ACT_CD4_TCELL_UP
GCM_BECD1, GCM_BECD1
GO_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_COMPLEX, GO_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_COMPLEX
KEGG_RENAL_CELL_CARCINOMA, KEGG_RENAL_CELL_CARCINOMA
GO_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX, GO_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX
GSE8921_3H_VS_24H_TLR1_2_STIM_MONOCYTE_DN, GSE8921_3H_VS_24H_TLR1_2_STIM_MONOCYTE_DN
GSE20500_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP, GSE20500_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP
GNF2_CBFb, GNF2_CBFb
GSE27291_0H_VS_6H_STIM_GAMMADelta_TCELL_UP, GSE27291_0H_VS_6H_STIM_GAMMADelta_TCELL_UP
NADLER_OBESITY_UP, NADLER_OBESITY_UP
GSE360_L_MAJOR_VS_M_TUBERCULOSIS_DC_UP, GSE360_L_MAJOR_VS_M_TUBERCULOSIS_DC_UP
GSE27670_BLIMP1_VS_LMP1_TRANSDUCED_GC_BCELL_UP, GSE27670_BLIMP1_VS_LMP1_TRANSDUCED_GC_BCELL_UP
GO_PURINE_NUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS, GO_PURINE_NUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS
WIELAND_UP_BY_HBV_INFECTION, WIELAND_UP_BY_HBV_INFECTION
GSE27786_BCELL_VS_CD4_TCELL_DN, GSE27786_BCELL_VS_CD4_TCELL_DN
SESTO_RESPONSE_TO_UV_C7, SESTO_RESPONSE_TO_UV_C7
GSE360_CTRL_VS_M_TUBERCULOSIS_MAC_DN, GSE360_CTRL_VS_M_TUBERCULOSIS_MAC_DN
REACTOME_RNA_POL_II_TRANSCRIPTION_PRE_INITIATION_AND_PROMOTER_OPENING, REACTOME_RNA_POL_II_TRANSCRIPTION_PRE_INITIATION_AND_PROMOTER_OPENING
GO_RIBONUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS, GO_RIBONUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS
LOPEZ_TRANSLATION_VIA_FN1_SIGNALING, LOPEZ_TRANSLATION_VIA_FN1_SIGNALING
MARTINEZ_RESPONSE_TO TRABECTEDIN_UP, MARTINEZ_RESPONSE_TO TRABECTEDIN_UP
GNF2_INPP5D, GNF2_INPP5D
HOLLEMAN_PREDNISOLONE_RESISTANCE_B_ALL_UP, HOLLEMAN_PREDNISOLONE_RESISTANCE_B_ALL_UP
GSE18281_CORTICAL_VS_MEDULLARY_THYMOCYTE_UP, GSE18281_CORTICAL_VS_MEDULLARY_THYMOCYTE_UP
ACAGGT_MIR10A_MIR10B, ACAGGT_MIR10A_MIR10B
LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_SMALL_VS_HUGE_DN, LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_SMALL_VS_HUGE_DN
DELPUCH_FOXO3_TARGETS_UP, DELPUCH_FOXO3_TARGETS_UP
GO_POSITIVE_REGULATION_OF_VIRAL_TRANSCRIPTION, GO_POSITIVE_REGULATION_OF_VIRAL_TRANSCRIPTION
GSE17721_CTRL_VS_LPS_0.5H_BMDC_DN, GSE17721_CTRL_VS_LPS_0.5H_BMDC_DN
GO_PROTEIN_EXPORT_FROM_NUCLEUS, GO_PROTEIN_EXPORT_FROM_NUCLEUS
GSE34006_WT_VS_A2AR_KO_TREG_UP, GSE34006_WT_VS_A2AR_KO_TREG_UP
AGCTCCT_MIR28, AGCTCCT_MIR28
GO_PURINE_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS, GO_PURINE_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS
KOBAYASHI_EGFR_SIGNALING_24HR_UP, KOBAYASHI_EGFR_SIGNALING_24HR_UP
SRF_01, SRF_01
GSE5099_MONOCYTE_VS_CLASSICAL_M1_MACROPHAGE_UP, GSE5099_MONOCYTE_VS_CLASSICAL_M1_MACROPHAGE_UP
GO_PRE_MRNA_BINDING, GO_PRE_MRNA_BINDING
GSE17721_CTRL_VS_POLYIC_0.5H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_0.5H_BMDC_DN
GSE37301_HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_UP, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_UP
PID_RAC1_PATHWAY, PID_RAC1_PATHWAY
HONMA_DOCETAXEL_RESISTANCE, HONMA_DOCETAXEL_RESISTANCE
SASSON_RESPONSE_TO_GONADOTROPHINS_DN, SASSON_RESPONSE_TO_GONADOTROPHINS_DN
GSE30971_2H_VS_4H_LPS_STIM_MACROPHAGE_WBP7_KO_DN, GSE30971_2H_VS_4H_LPS_STIM_MACROPHAGE_WBP7_KO_DN
LIU_PROSTATE_CANCER_UP, LIU_PROSTATE_CANCER_UP
GSE17721_CTRL_VS_POLYIC_1H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_1H_BMDC_DN
GO_RESPONSE_TO_NUTRIENT, GO_RESPONSE_TO_NUTRIENT
GO_ATPASE_REGULATOR_ACTIVITY, GO_ATPASE_REGULATOR_ACTIVITY
GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN, GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN
GO_POSITIVE_REGULATION_OF_ATPASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_ATPASE_ACTIVITY
KHETCHOUMIAN_TRIM24_TARGETS_UP, KHETCHOUMIAN_TRIM24_TARGETS_UP
GO_CELLULAR_RESPONSE_TO_INORGANIC_SUBSTANCE, GO_CELLULAR_RESPONSE_TO_INORGANIC_SUBSTANCE
MODULE_116, MODULE_116
GSE21360_PRIMARY_VS_SECONDARY_MEMORY_CD8_TCELL_DN, GSE21360_PRIMARY_VS_SECONDARY_MEMORY_CD8_TCELL_DN
BAE_BRCA1_TARGETS_DN, BAE_BRCA1_TARGETS_DN
SATO_SILENCED_BY_DEACETYLATION_IN_PANCREATIC_CANCER, SATO_SILENCED_BY_DEACETYLATION_IN_PANCREATIC_CANCER
GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4_KO_BCELL_UP, GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4_KO_BCELL_UP
GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_UP, GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_UP
GO_NEGATIVE_REGULATION_OF_CELL_ACTIVATION, GO_NEGATIVE_REGULATION_OF_CELL_ACTIVATION
GSE28783_ANTL_MIR33_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_UP, GSE28783_ANTL_MIR33_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_UP
LABBE_TARGETS_OF_TGFB1_AND_WNT3A_UP, LABBE_TARGETS_OF_TGFB1_AND_WNT3A_UP
GO_TRANSCRIPTION_FACTOR_TFIID_COMPLEX, GO_TRANSCRIPTION_FACTOR_TFIID_COMPLEX
GO_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY, GO_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY
GO_ATPASE_ACTIVATOR_ACTIVITY, GO_ATPASE_ACTIVATOR_ACTIVITY
PID_RHOA_PATHWAY, PID_RHOA_PATHWAY
GO_REGULATION_OF_DNA_BINDING, GO_REGULATION_OF_DNA_BINDING
MODULE_419, MODULE_419
MODULE_524, MODULE_524
GCM_VAV1, GCM_VAV1
GO_RESPONSE_TO_CADMIUM_ION, GO_RESPONSE_TO_CADMIUM_ION
GO_REGULATION_OF_B_CELL_ACTIVATION, GO_REGULATION_OF_B_CELL_ACTIVATION
ACCTGTG_UNKNOWN, ACCTGTG_UNKNOWN
BIOCARTA_RHO_PATHWAY, BIOCARTA_RHO_PATHWAY
PDGF_ERK_DN.V1_UP, PDGF_ERK_DN.V1_UP
GO_POSITIVE_REGULATION_OF_CYTOSKELETON_ORGANIZATION, GO_POSITIVE_REGULATION_OF_CYTOSKELETON_ORGANIZATION
PRAMOONJAGO_SOX4_TARGETS_DN, PRAMOONJAGO_SOX4_TARGETS_DN
WU_HBX_TARGETS_1_DN, WU_HBX_TARGETS_1_DN
GSE17721_CTRL_VS_CPG_0.5H_BMDC_DN, GSE17721_CTRL_VS_CPG_0.5H_BMDC_DN
GSE3982_DC_VS_NEUTROPHIL_LPS_STIM_DN, GSE3982_DC_VS_NEUTROPHIL_LPS_STIM_DN
GO_ATP_HYDROLYSIS_COUPLED_TRANSMEMBRANE_TRANSPORT, GO_ATP_HYDROLYSIS_COUPLED_TRANSMEMBRANE_TRANSPORT
BIOCARTA_CTCF_PATHWAY, BIOCARTA_CTCF_PATHWAY
GHO_ATF5_TARGETS_DN, GHO_ATF5_TARGETS_DN
GO_POSITIVE_REGULATION_OF_PROTEIN_DEACETYLATION, GO_POSITIVE_REGULATION_OF_PROTEIN_DEACETYLATION
GO_PROTEIN_DEPOLYMERIZATION, GO_PROTEIN_DEPOLYMERIZATION
GO_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY_TRANSPOSING_S_S_BONDS, GO_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY_TRANSPOSING_S_S_BONDS
GO_POSITIVE_REGULATION_OF_PEPTIDYL_TYROSINE_PHOSPHORYLATION, GO_POSITIVE_REGULATION_OF_PEPTIDYL_TYROSINE_PHOSPHORYLATION
BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_48HR_DN, BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_48HR_DN
WANG_PROSTATE_CANCER_ANDROGEN_INDEPENDENT, WANG_PROSTATE_CANCER_ANDROGEN_INDEPENDENT
GO_POSITIVE_REGULATION_OF_LIPID_KINASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_LIPID_KINASE_ACTIVITY
BONOME_OVARIAN_CANCER_POOR_SURVIVAL_DN, BONOME_OVARIAN_CANCER_POOR_SURVIVAL_DN
GO_ENTRAINMENT_OF_CIRCADIAN_CLOCK_BY_PHOTOPERIOD, GO_ENTRAINMENT_OF_CIRCADIAN_CLOCK_BY_PHOTOPERIOD
GO_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_PROTON_TRANSPORTING_DOMAIN, GO_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_PROTON_TRANSPORTING_DOMAIN
ZHENG_RESPONSE_TO_ARSENITE_UP, ZHENG_RESPONSE_TO_ARSENITE_UP
PAX5_02, PAX5_02
GO_PRP19_COMPLEX, GO_PRP19_COMPLEX
RIZKI_TUMOR_INVASIVENESS_2D_UP, RIZKI_TUMOR_INVASIVENESS_2D_UP
GO_REGULATION_OF_EXOSOMAL_SECRETION, GO_REGULATION_OF_EXOSOMAL_SECRETION
GO_CELLULAR_GLUCOSE_HOMEOSTASIS, GO_CELLULAR_GLUCOSE_HOMEOSTASIS
CHEOK_RESPONSE_TO_MERCAPTOPURINE_AND_HD_MTX_DN, CHEOK_RESPONSE_TO_MERCAPTOPURINE_AND_HD_MTX_DN
GO_ENDOCYTIC_RECYCLING, GO_ENDOCYTIC_RECYCLING
MALONEY_RESPONSE_TO_17AAG_UP, MALONEY_RESPONSE_TO_17AAG_UP
VERRECCHIA_RESPONSE_TO_TGFB1_C4, VERRECCHIA_RESPONSE_TO_TGFB1_C4
REACTOME_SEMA3A_PAK_DEPENDENT_AXON_REPULSION, REACTOME_SEMA3A_PAK_DEPENDENT_AXON_REPULSION
GO_NEGATIVE_REGULATION_OF_HOMEOSTATIC_PROCESS, GO_NEGATIVE_REGULATION_OF_HOMEOSTATIC_PROCESS
GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_PROLIFERATION, GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_PROLIFERATION
GO_PROTEIN_SERINE_THREONINE_KINASE_ACTIVATOR_ACTIVITY, GO_PROTEIN_SERINE_THREONINE_KINASE_ACTIVATOR_ACTIVITY
GO_BLOOD_MICROPARTICLE, GO_BLOOD_MICROPARTICLE
GO_PURINE_CONTAINING_COMPOUND_SALVAGE, GO_PURINE_CONTAINING_COMPOUND_SALVAGE
GO_MITOCHONDRIAL_CALCIIUM_ION_HOMEOSTASIS, GO_MITOCHONDRIAL_CALCIIUM_ION_HOMEOSTASIS
GO_CALCIIUM_DEPENDENT_PROTEIN_BINDING, GO_CALCIIUM_DEPENDENT_PROTEIN_BINDING
GO_CHONDROITIN_SULFATE_CATABOLIC_PROCESS, GO_CHONDROITIN_SULFATE_CATABOLIC_PROCESS
SPELMAN_LYMPHOBLAST_EUROPEAN_VS_ASIAN_2FC_UP, SPELMAN_LYMPHOBLAST_EUROPEAN_VS_ASIAN_2FC_UP
JIANG_TIP30_TARGETS_DN, JIANG_TIP30_TARGETS_DN
BIOCARTA_RAC1_PATHWAY, BIOCARTA_RAC1_PATHWAY
GO_GOLGI_LOCALIZATION, GO_GOLGI_LOCALIZATION
GALIE_TUMOR_ANGIOGENESIS, GALIE_TUMOR_ANGIOGENESIS
GO_NEGATIVE_REGULATION_OF_B_CELL_ACTIVATION, GO_NEGATIVE_REGULATION_OF_B_CELL_ACTIVATION
HASLINGER_B_CLL_WITH_CHROMOSOME_12_TRISOMY, HASLINGER_B_CLL_WITH_CHROMOSOME_12_TRISOMY
MEDINA_SMARCA4_TARGETS, MEDINA_SMARCA4_TARGETS
BANDRES_RESPONSE_TO_CARMUSTIN_MGMT_48HR_DN, BANDRES_RESPONSE_TO_CARMUSTIN_MGMT_48HR_DN
GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_UP, GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_UP
KYN, WERNER_SYNDROM_DN, KYN, WERNER_SYNDROM_DN
REACTOME_CS_DS_DEGRADATION, REACTOME_CS_DS_DEGRADATION
GO_ALTERNATIVE_MRNA_SPLICING_VIA_SPLICEOSOME, GO_ALTERNATIVE_MRNA_SPLICING_VIA_SPLICEOSOME
SASAKI_TARGETS_OF_TP73_AND_TP63, SASAKI_TARGETS_OF_TP73_AND_TP63
GSE3994_WT_VS_PAC1_KO_ACTIVATED_MAST_CELL_UP, GSE3994_WT_VS_PAC1_KO_ACTIVATED_MAST_CELL_UP
DASU_IL6_SIGNALING_UP, DASU_IL6_SIGNALING_UP
WU_HBX_TARGETS_2_UP, WU_HBX_TARGETS_2_UP
GO_POSITIVE_REGULATION_OF_HISTONE_DEACETYLATION, GO_POSITIVE_REGULATION_OF_HISTONE_DEACETYLATION
REACTOME_GENERATION_OF_SECOND_MESSENGER_MOLECULES, REACTOME_GENERATION_OF_SECOND_MESSENGER_MOLECULES
GO_SULFUR_COMPOUND_CATABOLIC_PROCESS, GO_SULFUR_COMPOUND_CATABOLIC_PROCESS
CHEOK_RESPONSE_TO_MERCAPTOPURINE_UP, CHEOK_RESPONSE_TO_MERCAPTOPURINE_UP
GO_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX_COUPLING_FACTOR_F_O, GO_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX_COUPLING_FACTOR_F_O
GO_CELL_TRAILING_EDGE, GO_CELL_TRAILING_EDGE
REACTOME_HS_GAG_DEGRADATION, REACTOME_HS_GAG_DEGRADATION
GO_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_BODY_SORTING_PATHWAY, GO_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_BODY_SORTING_PATHWAY

_TYPE_2_DIABETES_PBM_AT_DX_UP, GSE9006_HEALTHY_VS_TYPE_2_DIABETES_PBM_AT_DX_UP