

	GSE9988_LPS_VS_CTRL_TREATED_MONOCYTE_DN
	GSE9988_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN
	GSE9988_ANTL_TREM1_VS_CTRL_TREATED_MONOCYTES_DN
	GSE9988_LOW_LPS_VS_VEHICLE_TREATED_MONOCYTE_DN
	GSE9988_LOW_LPS_VS_CTRL_TREATED_MONOCYTE_DN
	GSE9988_ANTL_TREM1_AND_LPS_VS_VEHICLE_TREATED_MONOCYTES_DN
	GSE9988_ANTL_TREM1_AND_LPS_VS_CTRL_TREATED_MONOCYTES_DN
	GSE2706_UNSTIM_VS_8H_LPS_AND_R848_DC_UP
	GSE14000_UNSTIM_VS_16H_LPS_DC_TRANSLATED_RNA_UP
	GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_DN
	GSE9988_ANTL_TREM1_VS_VEHICLE_TREATED_MONOCYTES_DN
	GSE7509_DC_VS_MONOCYTE_WITH_FCGRIIB_STIM_UP
	GSE14000_UNSTIM_VS_16H_LPS_DC_UP
	GSE36826_NORMAL_VS_STAPH_AUREUS_INF_IL1R_KO_SKIN_DN
	GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_4H_DN
	GSE13484_3H_UNSTIM_VS_YF17D_VACCINE_STIM_PBMC_UP
	GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_KO_2H_DN
	GSE9960_HEALTHY_VS_GRAM_NEG_SEPSIS_PBMC_UP
	GSE16755_CTRL_VS_IFNA_TREATED_MAC_UP
	GSE41176_UNSTIM_VS_ANTL_IGM_STIM_BCELL_3H_DN
	GSE42088_UNINF_VS_LEISHMANIA_INF_DC_24H_UP
	GSE17721_CTRL_VS_LPS_4H_BMDC_UP
	GSE1432_CTRL_VS_IFNG_6H_MICROGLIA_UP
	GSE2706_UNSTIM_VS_8H_LPS_DC_UP
	GSE13484_UNSTIM_VS_12H_YF17D_VACCINE_STIM_PBMC_UP
	GSE13484_UNSTIM_VS_3H_YF17D_VACCINE_STIM_PBMC_UP
	GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CYS_IN_MAC_UP
	GSE3982_EOSINOPHIL_VS_CENT_MEMORY_CD4_TCELL_UP
	GSE13484_12H_UNSTIM_VS_YF17D_VACCINE_STIM_PBMC_UP
	GO_TRANSCRIPTION_COUPLED_NUCLEOTIDE_EXCISION_REPAIR
	GSE360_CTRL_VS_M_TUBERCULOSIS_DC_UP
	GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_DN
	GSE41867_NAIVE_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN
	GSE6681_DELETED_FOXP3_VS_WT_TREG_DN
	GSE1432_1H_VS_6H_IFNG_MICROGLIA_UP
	GSE15930_STIM_VS_STIM_AND_IL12_72H_CD8_T_CELL_UP
	GSE36476_CTRL_VS_TSST_ACT_40H_MEMORY_CD4_TCELL_YOUNG_UP
	GCM_SUFU
	GSE29618_PRE_VS_DAY7_FLU_VACCINE_MDC_DN
	GSE42021_CD24HI_VS_CD24INT_TREG_THYMUS_UP
	GSE1432_CTRL_VS_IFNG_24H_MICROGLIA_UP
	GSE42021_TREG_VS_TCONV_PLN_DN
	LINDGREN_BLADDER_CANCER_CLUSTER_1_UP
	MODULE_277
	GSE41867_DAY15_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN
	GSE3982_CTRL_VS_LPS_1H_NEUTROPHIL_UP
	GSE360_LOW_DOSE_B_MALAYI_VS_M_TUBERCULOSIS_DC_UP
	GSE3982_MAST_CELL_VS_TH2_UP
	GSE18281_SUBCAPSULAR_VS_PERIMEDULLARY_CORTICAL_REGION_OF_THYMUS_UP
	MORF_PPP5C
	GSE21360_NAIVE_VS_TERTIARY_MEMORY_CD8_TCELL_DN
	GCM_BAG5
	GSE13522_CTRL_VS_T_CRUZY_STRAIN_INF_SKIN_IFNG_KO_UP
	GSE37301_MULTIPOTENT_PROGENITOR_VS_RAG2_KO_NK_CELL_UP
	GSE3203_HEALTHY_VS_INFLUENZA_INFECTED_LN_BCELL_UP
	GSE42021_TREG_PLN_VS_CD24LO_TREG_THYMUS_UP
	MORF_REV3L
	GAGCTGG_MIR337
	GSE10325_CD4_TCELL_VS_MYELOID_UP
	GSE19401_PLN_VS_Peyers_PATCH_FOLLICULAR_DC_UP
	GSE27786_LSK_VS_NKTCCELL_DN
	GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_DN
	GSE14308_TH2_VS_NAIVE_CD4_TCELL_DN
	GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_DN
	GNF2_BNIP2
	GSE35685_CD34POS_CD38NEG_VS_CD34POS_CD10POS_BONE_MARROW_UP
	GSE20715_0H_VS_48H_OZONE_LUNG_UP
	GNF2_TYK2
	GSE36476_CTRL_VS_TSST_ACT_16H_MEMORY_CD4_TCELL_OLD_UP
	GSE7831_UNSTIM_VS_INFLUENZA_STIM_PDC_4H_UP
	GNF2_CD53
	GSE360_T_GONDII_VS_B_MALAYI_HIGH_DOSE_DC_DN
	GSE21360_NAIVE_VS_TERTIARY_MEMORY_CD8_TCELL_UP
	GSE1460_DP_THYMOCYTE_VS_THYMIC_STROMAL_CELL_UP
	GSE11924_TFH_VS_TH2_CD4_TCELL_DN
	YAGL_AML_WITH_T_9_11_TRANSLOCATION
	GSE27859_CD11C_INT_F480_HI_MACROPHAGE_VS_CD11C_ING_F480_INT_DC_DN
	GSE22611_UNSTIM_VS_6H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_DN
	GCM_CSNK1D
	PECE_MAMMARY_STEM_CELL_DN
	GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_UP
	GSE1448_CTRL_VS_ANTL_VBETA5_DP_THYMOCYTE_DN
	GO_POSITIVE_REGULATION_OF_PROTEIN_COMPLEX_ASSEMBLY
	GSE42021_CD24INT_VS_CD24LOW_TREG_THYMUS_UP
	MORF_PSMF1
	GSE41978_ID2_KO_AND_BIM_KO_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN
	REACTOME_DIABETES_PATHWAYS
	GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_WITH_EGR2_DELETED_DN
	GSE37605_C57BL6_VS_NOD_FOXP3_FUSION_GFP_TCONV_UP
	PENG_LEUCINE_DEPRIVATION_UP
	GSE1448_CTRL_VS_ANTL_VALPHA2_DP_THYMOCYTE_DN
	GSE360_DC_VS_MAC_T_GONDII_DN
	chr22q13
	PID_CDC42_PATHWAY
	ROSS_AML_WITH_CBFb_MYH11_FUSION
	CHNG_MULTIPLE_MYELOMA_HYPERPLOID_DN
	GCM_RAP2A
	GO_NEGATIVE_REGULATION_OF_CHROMATIN_MODIFICATION
	GSE15930_STIM_VS_STIM_AND_IFNAB_24H_CD8_T_CELL_UP
	GO_CYTOPLASMIC_MICROTUBULE_ORGANIZATION
	GNF2_VAV1
	BOYAUULT_LIVER_CANCER_SUBCLASS_G12_UP
	FAELT_B_CLL_WITH_VH_REARRANGEMENTS_UP
	GO_NUCLEOTIDE_EXCISION_REPAIR_PREINCISION_COMPLEX_STABILIZATION
	MODULE_243
	PID_INSULIN_PATHWAY
	MORF_CDC2L5
	GCM_DEAF1
	GSE14308_TH1_VS_NAIVE_CD4_TCELL_DN
	GSE20366_EX_VIVO_VS_HOMEOSTATIC_CONVERSION_NAIVE_CD4_TCELL_DN
	GSE37532_VISCERAL_ADIPOSE_TISSUE_VS_LN_DERIVED_PPARG_KO_TREG_CD4_TCELL_DN
	LTE2_UP.V1_UP
	SASSON_RESPONSE_TO_FORSKOLIN_DN
	GO_REGULATION_OF_HISTONE_H3_K9_METHYLATION
	TBK1.DN.48HRS_UP
	GSE8515_CTRL_VS_IL1_4H_STIM_MAC_UP
	GO_ACTIN_NUCLEATION
	GSE27786_BCELL_VS_CD8_TCELL_DN
	GO_PROTEIN_TYROSINE_PHOSPHATASE_ACTIVITY
	MARIADASON_RESPONSE_TO_CURCUMIN_SULINDAC_5
	GO_RNA_POLYMERASE_II_CORE_BINDING
	GO_HISTONE_MRNA_CATABOLIC_PROCESS
	GENTILE_UV_RESPONSE_CLUSTER_D8
	GSE24210_IL35_TREATED_VS_UNTREATED_TCONV_CD4_TCELL_DN
	GO_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
	GO_PEPTIDYL_CYSSTEINE_MODIFICATION
	JAZAG_TGFB1_SIGNALING_UP
	GO_POSITIVE_REGULATION_OF_CYTOSKELETON_ORGANIZATION
	SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5
	GO_CELL_CORTEX_PART
	BASSO_HAIRY_CELL_LEUKEMIA_UP
	GO_DSRNA_FRAGMENTATION
	SIG_REGULATION_OF_THE_ACTIN_CYTOSKELETON_BY_RHO_GTPASES
	GO_PROTEIN_DEALKYLATION
	PID_RAC1_PATHWAY
	AGTCAGC_MIR345
	GO_HISTONE_DEMETHYLASE_ACTIVITY
	GSE18203_CTRL_VS_INTRATUMORAL_CPG_INJ_MC38_TUMOR_UP
	GO_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY_INTERCONVERTING_ALDOSES_AND_KETOSES
	GO_DNA_REPLICATION_FACTOR_A_COMPLEX
	REACTOME_SYNTHESIS_SECRETION_AND_DEACYLATION_OF_GHRELIN
	GO_POSITIVE_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY
	PID_IL1_PATHWAY
	ASTON_MAJOR_DEPRESSIVE_DISORDER_UP
	WIKMAN_ASBESTOS_LUNG_CANCER_UP
	GGTAACC_MIR4095P
	GO_FAT_CELL_DIFFERENTIATION
	GO_NEGATIVE_REGULATION_OF_HISTONE_METHYLATION
	GO_METHIONINE_METABOLIC_PROCESS
	GO_METHIONINE_BIOSYNTHETIC_PROCESS
	GSE16450_IMMATURE_VS_MATURE_NEURON_CELL_LINE_6H_IFNA_STIM_UP
	PARK_TRETINOIN_RESPONSE_AND_RARA_PLZF_FUSION
	GO_RNAI_EFFECTOR_COMPLEX
	PID_AR_TF_PATHWAY
	GO_SNRNA_METABOLIC_PROCESS
	GSE360_T_GONDII_VS_M_TUBERCULOSIS_DC_DN
	GSE19401_PAM2CSK4_VS_RETINOIC_ACID_STIM_FOLLICULAR_DC_DN
	GO_PRE_MIRNA_PROCESSING
	LUI_THYROID_CANCER_PAX8_PPARG_UP
	PHESSE_TARGETS_OF_APC_AND_MBD2_UP
	GO_SULFUR_AMINO_ACID_BIOSYNTHETIC_PROCESS
	RUAN_RESPONSE_TO_TROGLITAZONE_DN
	GO_CORTICAL_CYTOSKELETON