

	GSE34156_NOD2_LIGAND_VS_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP
	GSE34156_NOD2_LIGAND_VS_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_DN
	GSE34156_NOD2_LIGAND_VS_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_DN
	GSE34156_UNTREATED_VS_24H_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_DN
	GSE34156_UNTREATED_VS_24H_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_DN
	GSE25088_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_UP
	GSE25088_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_UP
	GSE25123_CTRL_VS_IL4_STIM_MACROPHAGE_DN
	GSE25123_CTRL_VS_IL4_STIM_MACROPHAGE_DN
	GSE11057_NAIVE_VS_EFF_MEMORY_CD4_TCELL_UP
	GSE11057_NAIVE_VS_EFF_MEMORY_CD4_TCELL_UP
	GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_2H_ACT_CD4_TCELL_UP
	GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_2H_ACT_CD4_TCELL_UP
	GSE28237_FOLLICULAR_VS_EARLY_GC_BCELL_UP
	GSE28237_FOLLICULAR_VS_EARLY_GC_BCELL_UP
	GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_UP
	GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_UP
	GSE21670_IL6_VS_TGFB_AND_IL6_TREATED_CD4_TCELL_UP
	GSE21670_IL6_VS_TGFB_AND_IL6_TREATED_CD4_TCELL_UP
	GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_DN
	GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_DN
	GSE5099_DAY3_VS_DAY7_MCSF_TREATED_MACROPHAGE_UP
	GSE5099_DAY3_VS_DAY7_MCSF_TREATED_MACROPHAGE_UP
	GSE26495_NAIVE_VS_PD1LOW_CD8_TCELL_UP
	GSE26495_NAIVE_VS_PD1LOW_CD8_TCELL_UP
	GSE11057_NAIVE_VS_MEMORY_CD4_TCELL_UP
	GSE11057_NAIVE_VS_MEMORY_CD4_TCELL_UP
	GSE11057_EFF_MEM_VS_CENT_MEM_CD4_TCELL_DN
	GSE11057_EFF_MEM_VS_CENT_MEM_CD4_TCELL_DN
	GSE26495_NAIVE_VS_PD1HIGH_CD8_TCELL_UP
	GSE26495_NAIVE_VS_PD1HIGH_CD8_TCELL_UP
	GSE5542_UNTREATED_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP
	GSE5542_UNTREATED_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP
	GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_10H_DN
	GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_10H_DN
	HP_CLEFT_SOFT_PALATE
	HP_CLEFT_SOFT_PALATE
	GOBP_LIPOPROTEIN_METABOLIC_PROCESS
	GOBP_LIPOPROTEIN_METABOLIC_PROCESS
	GSE22045_TREG_VS_TCONV_DN
	GSE22045_TREG_VS_TCONV_DN
	GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_UP
	GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_UP
	GSE4984_LPS_VS_VEHICLE_CTRL_TREATED_DC_UP
	GSE4984_LPS_VS_VEHICLE_CTRL_TREATED_DC_UP
	HP_NEOPLASM_OF_THE_COLON
	HP_NEOPLASM_OF_THE_COLON
	GOBP_LIPOPROTEIN_BIOSYNTHETIC_PROCESS
	GOBP_LIPOPROTEIN_BIOSYNTHETIC_PROCESS
	GSE11057_NAIVE_VS_CENT_MEMORY_CD4_TCELL_UP
	GSE11057_NAIVE_VS_CENT_MEMORY_CD4_TCELL_UP
	GSE25087_TREG_VS_TCONV_ADULT_DN
	GSE25087_TREG_VS_TCONV_ADULT_DN
	MIR4424
	MIR4424
	HP_WEBBED_NECK
	HP_WEBBED_NECK
	GSE22886_NAIVE_CD4_TCELL_VS_NKCELL_UP
	GSE22886_NAIVE_CD4_TCELL_VS_NKCELL_UP
	HP_APLASIA_HYPOPLASIA_OF_THE_THUMB
	HP_APLASIA_HYPOPLASIA_OF_THE_THUMB
	GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS
	GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS
	GSE24081_CONTROLLER_VS_PROGRESSOR_HIV_SPECIFIC_CD8_TCELL_UP
	GSE24081_CONTROLLER_VS_PROGRESSOR_HIV_SPECIFIC_CD8_TCELL_UP
	GSE22886_CD8_VS_CD4_NAIVE_TCELL_DN
	GSE22886_CD8_VS_CD4_NAIVE_TCELL_DN
	MIR1286
	MIR1286
	MIR1302
	MIR1302
	HP_DUPLICATION_OF_THUMB_PHALANX
	HP_DUPLICATION_OF_THUMB_PHALANX
	MIR4722_5P
	MIR4722_5P
	BCAT_BILD_ET_AL_DN
	BCAT_BILD_ET_AL_DN
	HP_TRIPHALANGEAL_THUMB
	HP_TRIPHALANGEAL_THUMB
	MIR668_3P
	MIR668_3P
	REACTOME_INTRA_GOLGI_TRAFFIC
	REACTOME_INTRA_GOLGI_TRAFFIC
	HP_APLASIA_OF_THE_FINGERS
	HP_APLASIA_OF_THE_FINGERS
	REACTOME_TNF_SIGNALING
	REACTOME_TNF_SIGNALING
	GAZDA_DIAMOND_BLACKFAN_ANEMIA_MYELOID_UP
	GAZDA_DIAMOND_BLACKFAN_ANEMIA_MYELOID_UP
	LEE_EARLY_T_LYMPHOCYTE_DN
	LEE_EARLY_T_LYMPHOCYTE_DN
	HP_TAPERED_FINGER
	HP_TAPERED_FINGER
	DESCARTES_MAIN_FETAL_HEMATOPOIETIC_STEM_CELLS
	DESCARTES_MAIN_FETAL_HEMATOPOIETIC_STEM_CELLS
	HP_TONIC_SEIZURE
	HP_TONIC_SEIZURE
	FARMER_BREAST_CANCER_CLUSTER_5
	FARMER_BREAST_CANCER_CLUSTER_5
	GSE2706_R848_VS_LPS_8H_STIM_DC_UP
	GSE2706_R848_VS_LPS_8H_STIM_DC_UP
	GOBP_PROTEIN_DEGLYCOSYLATION
	GOBP_PROTEIN_DEGLYCOSYLATION
	TRIP13_TARGET_GENES
	TRIP13_TARGET_GENES
	HP_NORMOCHROMIC_ANEMIA
	HP_NORMOCHROMIC_ANEMIA
	GOBP_P_BODY_ASSEMBLY
	GOBP_P_BODY_ASSEMBLY
	REACTOME_TNFR1_INDUCED_NFKAPPAB_SIGNALING_PATHWAY
	REACTOME_TNFR1_INDUCED_NFKAPPAB_SIGNALING_PATHWAY
	GOBP_MRNA_5_SPLICE_SITE_RECOGNITION
	GOBP_MRNA_5_SPLICE_SITE_RECOGNITION
	HP_PARTIAL_DUPLICATION_OF_THE_PHALANX_OF_HAND
	HP_PARTIAL_DUPLICATION_OF_THE_PHALANX_OF_HAND
	HP_ADENOCARCINOMA_OF_THE_INTESTINES
	HP_ADENOCARCINOMA_OF_THE_INTESTINES
	HP_CUPPED_EAR
	HP_CUPPED_EAR
	KYNG_DNA_DAMAGE_BY_GAMMA_RADIATION
	KYNG_DNA_DAMAGE_BY_GAMMA_RADIATION
	GOBP_POSITIVE_REGULATION_OF_AXON_EXTENSION
	GOBP_POSITIVE_REGULATION_OF_AXON_EXTENSION
	REACTOME_TAK1_ACTIVATES_NFKB_BY_PHOSPHORYLATION_AND_ACTIVATION_OF_IKKS_COMPLEX
	REACTOME_TAK1_ACTIVATES_NFKB_BY_PHOSPHORYLATION_AND_ACTIVATION_OF_IKKS_COMPLEX
	ABE_INNER_EAR
	ABE_INNER_EAR
	PID_IL8_CXCR2_PATHWAY
	PID_IL8_CXCR2_PATHWAY
	HP_AGYRIA
	HP_AGYRIA
	GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
	GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
	HP_CONTRACTURES_INVOLVING_THE_JOINTS_OF_THE_FEET
	HP_CONTRACTURES_INVOLVING_THE_JOINTS_OF_THE_FEET
	HP_EEG_WITH_POLYSPIKE_WAVE_COMPLEXES
	HP_EEG_WITH_POLYSPIKE_WAVE_COMPLEXES
	GRATIAS_RETINOBLASTOMA_16Q24
	GRATIAS_RETINOBLASTOMA_16Q24
	HP_MASK_LIKE_FACIES
	HP_MASK_LIKE_FACIES
	GOBP_REGULATION_OF_CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY
	GOBP_REGULATION_OF_CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY
	TRAVAGLINI_LUNG_CD8_MEMORY_EFFECTOR_T_CELL
	TRAVAGLINI_LUNG_CD8_MEMORY_EFFECTOR_T_CELL
	REACTOME_STRIATED_MUSCLE_CONTRACTION
	REACTOME_STRIATED_MUSCLE_CONTRACTION
	RASHI_RESPONSE_TO_IONIZING_RADIATION_3
	RASHI_RESPONSE_TO_IONIZING_RADIATION_3
	HP_LIPOATROPHY
	HP_LIPOATROPHY
	GOBP_MALE_MEIOSIS_I
	GOBP_MALE_MEIOSIS_I
	REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATION_MEDIATED_BY_ACTIVATED_HUMAN_TAK1
	REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATION_MEDIATED_BY_ACTIVATED_HUMAN_TAK1
	HP_ABNORMAL_SARCOMERE_MORPHOLOGY
	HP_ABNORMAL_SARCOMERE_MORPHOLOGY
	GOBP_COBALAMIN_METABOLIC_PROCESS
	GOBP_COBALAMIN_METABOLIC_PROCESS
	REACTOME_ACTIVATED_TAK1_MEDIATES_P38_MAPK_ACTIVATION
	REACTOME_ACTIVATED_TAK1_MEDIATES_P38_MAPK_ACTIVATION
	HALLMARK_NOTCH_SIGNALING
	HALLMARK_NOTCH_SIGNALING
	REACTOME_SUPPRESSION_OF_PHAGOSOMAL_MATURATION
	REACTOME_SUPPRESSION_OF_PHAGOSOMAL_MATURATION
	GOCC_TRANSCRIPTION_EXPORT_COMPLEX
	GOCC_TRANSCRIPTION_EXPORT_COMPLEX
	GOBP_LEUKOTRIENE_TRANSPORT
	GOBP_LEUKOTRIENE_TRANSPORT
	BUSSLINGER_DUODENAL_EC_CELLS
	BUSSLINGER_DUODENAL_EC_CELLS
	GOBP_MUSCLE_FILAMENT_SLIDING
	GOBP_MUSCLE_FILAMENT_SLIDING
	GOBP_SARCOMERE_ORGANIZATION
	GOBP_SARCOMERE_ORGANIZATION
	GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H3_K4_SPECIFIC
	GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H3_K4_SPECIFIC
	chr3q13
	chr3q13
	HEDENFALK_BREAST_CANCER_BRACX_UP
	HEDENFALK_BREAST_CANCER_BRACX_UP
	GOBP_POSITIVE_REGULATION_OF_EMBRYONIC_DEVELOPMENT
	GOBP_POSITIVE_REGULATION_OF_EMBRYONIC_DEVELOPMENT
	GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
	GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
	PGBD5_TARGET_GENES
	PGBD5_TARGET_GENES
	GOBP_CELLULAR_RESPONSE_TO_ACIDIC_PH
	GOBP_CELLULAR_RESPONSE_TO_ACIDIC_PH
	GOMF_ABC_TYPE GLUTATHIONE_S_CONJUGATE_TRANSPORTER_ACTIVITY
	GOMF_ABC_TYPE GLUTATHIONE_S_CONJUGATE_TRANSPORTER_ACTIVITY
	GOCC_SET1C_COMPASS_COMPLEX
	GOCC_SET1C_COMPASS_COMPLEX
	HP_INTERNALLY_NUCLEATED_SKELETAL_MUSCLE_FIBERS
	HP_INTERNALLY_NUCLEATED_SKELETAL_MUSCLE_FIBERS