

	GSE6674_UNSTIM_VS_ANTL_IJGM_AND_CPG_STIM_BCELL_UP, GSE6674_UNSTIM_VS_ANTL_IJGM_AND_CPG_STIM_BCELL_UP
	GSE9650_EFFECTOR_VS_MEMORY_CD8_TCELL_UP, GSE9650_EFFECTOR_VS_MEMORY_CD8_TCELL_UP
	GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_DN, GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_DN
	GSE9878_CTRL_VS_EBF_TRANSDUCED_PAX5_KO_PRO_BCELL_DN, GSE9878_CTRL_VS_EBF_TRANSDUCED_PAX5_KO_PRO_BCELL_DN
	GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_DN, GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_DN
	GSE17812_WT_VS_THPOK_KO_MEMORY_CD8_TCELL_DN, GSE17812_WT_VS_THPOK_KO_MEMORY_CD8_TCELL_DN
	GSE7509_UNSTIM_VS_IFNA_STIM_IMMATURE_DC_UP, GSE7509_UNSTIM_VS_IFNA_STIM_IMMATURE_DC_UP
	GSE4984_UNTREATED_VS_LPS_TREATED_DC_DN, GSE4984_UNTREATED_VS_LPS_TREATED_DC_DN
	GSE8685_IL2_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP, GSE8685_IL2_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP
	HP_ABNORMALITY_OF_THE_UVULA, HP_ABNORMALITY_OF_THE_UVULA
	REACTOME_G2_M_DNA_DAMAGE_CHECKPOINT, REACTOME_G2_M_DNA_DAMAGE_CHECKPOINT
	GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP
	HP_OLIGOHYDRAMNIOS, HP_OLIGOHYDRAMNIOS
	GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP, GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP
	GOBP_MITOTIC_DNA_INTEGRITY_CHECKPOINT, GOBP_MITOTIC_DNA_INTEGRITY_CHECKPOINT
	GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_IRF8_KO_DN, GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_IRF8_KO_DN
	GSE32034_LY6C_HIGH_VS_LOW_ROSIGLIZATONE_TREATED_MONOCYTE_DN, GSE32034_LY6C_HIGH_VS_LOW_ROSIGLIZATONE_TREATED_MONOCYTE_DN
	GOBP_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION, GOBP_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION
	GSE40441_NRP1_POS_INDUCED_TREG_VS_NRP1_NEG_NATURAL_TREG_DN, GSE40441_NRP1_POS_INDUCED_TREG_VS_NRP1_NEG_NATURAL_TREG_DN
	GOBP_REGULATION_OF_CELL_CYCLE_ARREST, GOBP_REGULATION_OF_CELL_CYCLE_ARREST
	MIR1236_3P, MIR1236_3P
	HP_ABNORMAL_TRACHEA_MORPHOLOGY, HP_ABNORMAL_TRACHEA_MORPHOLOGY
	SRC_UP.V1_DN, SRC_UP.V1_DN
	GSE20366_EX_VIVO_VS_HOMEOSTATIC_CONVERSION_NAIVE_CD4_TCELL_DN, GSE20366_EX_VIVO_VS_HOMEOSTATIC_CONVERSION_NAIVE_CD4_TCELL_DN
	HP_OSTEOLYSIS, HP_OSTEOLYSIS
	GSE27786_LSK_VS_NEUTROPHIL_DN, GSE27786_LSK_VS_NEUTROPHIL_DN
	HP_APLASIA_HYPOPLASIA_INVOLVING_BONES_OF_THE_THORAX, HP_APLASIA_HYPOPLASIA_INVOLVING_BONES_OF_THE_THORAX
	HP_SHORT_PALPEBRAL_FISSURE, HP_SHORT_PALPEBRAL_FISSURE
	MIR629_5P, MIR629_5P
	WP_ATM_SIGNALING_PATHWAY, WP_ATM_SIGNALING_PATHWAY
	GSE24574_BCL6_LOW_TFH_VS_TCONV_CD4_TCELL_UP, GSE24574_BCL6_LOW_TFH_VS_TCONV_CD4_TCELL_UP
	MIR6729_3P, MIR6729_3P
	MIR320E, MIR320E
	MIR2682_5P, MIR2682_5P
	REACTOME_ENDOSOMAL_SORTING_COMPLEX_REQUIRED_FOR_TRANSPORT_ESCRT, REACTOME_ENDOSOMAL_SORTING_COMPLEX_REQUIRED_FOR_TRANSPORT_ESCRT
	HP_TRACHEOSOPHAGEAL_FISTULA, HP_TRACHEOSOPHAGEAL_FISTULA
	HP_ABNORMALITY_OF_THE_HUMERORADIAL_JOINT, HP_ABNORMALITY_OF_THE_HUMERORADIAL_JOINT
	GSE9006_1MONTH_VS_4MONTH_POST_TYPE_1_DIABETES_DX_PBMCDN, GSE9006_1MONTH_VS_4MONTH_POST_TYPE_1_DIABETES_DX_PBMCDN
	NADLER_HYPERGLYCEMIA_AT_OBESITY, NADLER_HYPERGLYCEMIA_AT_OBESITY
	LEE_LIVER_CANCER_MYC_UP, LEE_LIVER_CANCER_MYC_UP
	GOBP_MULTIVESICULAR_BODY_ORGANIZATION, GOBP_MULTIVESICULAR_BODY_ORGANIZATION
	HP_TRACHEOMALACIA, HP_TRACHEOMALACIA
	HP_ABNORMALITY_OF_THE_RADIAL_HEAD, HP_ABNORMALITY_OF_THE_RADIAL_HEAD
	HP_FATIGABLE_WEAKNESS_OF_BULBAR_MUSCLES, HP_FATIGABLE_WEAKNESS_OF_BULBAR_MUSCLES
	JIANG_TIP30_TARGETS_UP, JIANG_TIP30_TARGETS_UP
	HP_PIERRE_ROBIN_SEQUENCE, HP_PIERRE_ROBIN_SEQUENCE
	RYCACNNRNNRNCAG_UNKNOWN, RYCACNNRNNRNCAG_UNKNOWN
	MIR4710, MIR4710
	HP_ABNORMAL_EPIPHYSEAL_OSSIFICATION, HP_ABNORMAL_EPIPHYSEAL_OSSIFICATION
	GSE4590_SMALL_VS_VPREB_POS_LARGE_PRE_BCELL_DN, GSE4590_SMALL_VS_VPREB_POS_LARGE_PRE_BCELL_DN
	LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_LARGE_VS_TINY_DN, LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_LARGE_VS_TINY_DN
	GOCC_MULTIVESICULAR_BODY, GOCC_MULTIVESICULAR_BODY
	GOMF_MEMBRANE_INSERTASE_ACTIVITY, GOMF_MEMBRANE_INSERTASE_ACTIVITY
	GOBP_RESPONSE_TO_VITAMIN_E, GOBP_RESPONSE_TO_VITAMIN_E
	BIOCARTA_MPR_PATHWAY, BIOCARTA_MPR_PATHWAY
	GOBP_MIDBODY_ABSCISSION, GOBP_MIDBODY_ABSCISSION
	GOBP_TAIL_ANCHORED_MEMBRANE_PROTEIN_INSERTION_INTO_ER_MEMBRANE, GOBP_TAIL_ANCHORED_MEMBRANE_PROTEIN_INSERTION_INTO_ER_MEMBRANE
	DESCARTES_FETAL_THYMUS_THYMOCYTES, DESCARTES_FETAL_THYMUS_THYMOCYTES
	HP_PUSTULE, HP_PUSTULE
	GOBP_RESPONSE_TO_VITAMIN, GOBP_RESPONSE_TO_VITAMIN
	GOBP_PROTEIN_INSERTION_INTO_ER_MEMBRANE_BY_STOP_TRANSFER_MEMBRANE_ANCHOR_SEQUENCE, GOBP_PROTEIN_INSERTION_INTO_ER_MEMBRANE_BY_STOP_TRANSFER_MEMBRANE_ANCHOR_SEQUENCE
	GOCC EMC_COMPLEX, GOCC EMC_COMPLEX
	HP_OVARIAN_CARCINOMA, HP_OVARIAN_CARCINOMA
	HP_ANAL_STENOSIS, HP_ANAL_STENOSIS
	GOBP_NEGATIVE_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS
	MIR4535, MIR4535
	GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY
	HP_ANHIDROSIS, HP_ANHIDROSIS
	MIR6813_3P, MIR6813_3P
S_KLRG1INT_EFF_CD8_TCELL_DN, GSE10239_MEMORY_VS_KLRG1INT_EFF_CD8_TCELL_DN	REACTOME_DEPOLYMERISATION_OF_THE_NUCLEAR_LAMINA, REACTOME_DEPOLYMERISATION_OF_THE_NUCLEAR_LAMINA
	GOBP_LUNG_ALVEOLUS_DEVELOPMENT, GOBP_LUNG_ALVEOLUS_DEVELOPMENT
	GOBP_NEGATIVE_REGULATION_OF_CENTROSOME_DUPLICATION, GOBP_NEGATIVE_REGULATION_OF_CENTROSOME_DUPLICATION
	ZNF26_TARGET_GENES, ZNF26_TARGET_GENES
	GSE16386_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_6H_DN, GSE16386_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_6H_DN
	GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_SALMON_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_SALMON_DN
	HP_URETERAL_OBSTRUCTION, HP_URETERAL_OBSTRUCTION
	GOBP_PEPTIDYL_ARGININE_METHYLATION, GOBP_PEPTIDYL_ARGININE_METHYLATION
	PID_ARF6_DOWNSTREAM_PATHWAY, PID_ARF6_DOWNSTREAM_PATHWAY
	PARK_TRETINOLIN_RESPONSE, PARK_TRETINOLIN_RESPONSE
	HP_SCLEROCORNEA, HP_SCLEROCORNEA
	HOEK_T_CELL_2011_2012_TIV_ADULT_1DY_UP, HOEK_T_CELL_2011_2012_TIV_ADULT_1DY_UP
	GOBP_SPONTANEOUS_NEUROTRANSMITTER_SECRETION, GOBP_SPONTANEOUS_NEUROTRANSMITTER_SECRETION
	GOBP_XENOBIOTIC_CATABOLIC_PROCESS, GOBP_XENOBIOTIC_CATABOLIC_PROCESS
	GOBP_BILE_ACID_BIOSYNTHETIC_PROCESS, GOBP_BILE_ACID_BIOSYNTHETIC_PROCESS
	HP_ULNAR_DEVIATION_OF_THE_HAND_OR_OF_FINGERS_OF_THE_HAND, HP_ULNAR_DEVIATION_OF_THE_HAND_OR_OF_FINGERS_OF_THE_HAND
	HP_BROAD_ALVEOLAR_RIDGES, HP_BROAD_ALVEOLAR_RIDGES
	GOBP_LYMPHANGIOGENESIS, GOBP_LYMPHANGIOGENESIS
	GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_CYAN_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_CYAN_UP
	GOBP_LYMPH_VESSEL_DEVELOPMENT, GOBP_LYMPH_VESSEL_DEVELOPMENT
	GOBP_ENDOTHELIAL_CELL_FATE_COMMITMENT, GOBP_ENDOTHELIAL_CELL_FATE_COMMITMENT
	GOBP_SKELETAL_MUSCLE_THIN_FILAMENT_ASSEMBLY, GOBP_SKELETAL_MUSCLE_THIN_FILAMENT_ASSEMBLY
	HOWLIN_CITED1_TARGETS_2_UP, HOWLIN_CITED1_TARGETS_2_UP
	GOBP_POSITIVE_REGULATION_OF_CELL_DIVISION, GOBP_POSITIVE_REGULATION_OF_CELL_DIVISION
	GOBP_LYMPH_VESSEL_MORPHOGENESIS, GOBP_LYMPH_VESSEL_MORPHOGENESIS
	GOBP_RESPONSE_TO_PHENYLPROPANOID, GOBP_RESPONSE_TO_PHENYLPROPANOID
	GOBP_NEGATIVE_REGULATION_OF_CELLULAR_EXTRAVASATION, GOBP_NEGATIVE_REGULATION_OF_CELLULAR_EXTRAVASATION
	GOBP_SKELETAL_MYOFIBRIL_ASSEMBLY, GOBP_SKELETAL_MYOFIBRIL_ASSEMBLY
	HP_ABNORMAL_MORPHOLOGY_OF_MYOCARDIAL TRABECULAE, HP_ABNORMAL_MORPHOLOGY_OF_MYOCARDIAL TRABECULAE
	GOCC_ER_MEMBRANE_INSERTION_COMPLEX, GOCC_ER_MEMBRANE_INSERTION_COMPLEX
	KIM_PTEN_TARGETS_UP, KIM_PTEN_TARGETS_UP
	REACTOME_PROSTANOID_LIGAND_RECEPTORS, REACTOME_PROSTANOID_LIGAND_RECEPTORS
	HP_ABNORMAL_ENERGY_EXPENDITURE, HP_ABNORMAL_ENERGY_EXPENDITURE
	GOBP_ENDOCARDIUM_MORPHOGENESIS, GOBP_ENDOCARDIUM_MORPHOGENESIS
	GOBP_VENOUS_BLOOD_VESSEL_DEVELOPMENT, GOBP_VENOUS_BLOOD_VESSEL_DEVELOPMENT
	GOMF_PROSTANOID_RECEPTOR_ACTIVITY, GOMF_PROSTANOID_RECEPTOR_ACTIVITY
	REACTOME_GLUCURONIDATION, REACTOME_GLUCURONIDATION
	GOBP_G_PROTEIN_COUPLED_PURINERGIC_RECEPTOR_SIGNALING_PATHWAY, GOBP_G_PROTEIN_COUPLED_PURINERGIC_RECEPTOR_SIGNALING_PATHWAY
	ZHOU_PANCREATIC_EXOCRINE_PROGENITOR, ZHOU_PANCREATIC_EXOCRINE_PROGENITOR
	GOBP_ENDOCARDIUM_DEVELOPMENT, GOBP_ENDOCARDIUM_DEVELOPMENT
	HP_PALMAR_HYPERKERATOSIS, HP_PALMAR_HYPERKERATOSIS
	GOBP_ANTEROGRADE_DENDRITIC_TRANSPORT, GOBP_ANTEROGRADE_DENDRITIC_TRANSPORT
	GOBP_REGULATION_OF_LYMPHANGIOGENESIS, GOBP_REGULATION_OF_LYMPHANGIOGENESIS
	HP_DECREASED_CSF_HOMOVANILLIC_ACID, HP_DECREASED_CSF_HOMOVANILLIC_ACID
	HP_ATROPHY_OF_ALVEOLAR_RIDGES, HP_ATROPHY_OF_ALVEOLAR_RIDGES
	GOCC_SUPER_ELONGATION_COMPLEX, GOCC_SUPER_ELONGATION_COMPLEX
	GOMF_SULFATE_BINDING, GOMF_SULFATE_BINDING
	GOBP_ATRIAL_CARDIAC_MUSCLE_TISSUE_DEVELOPMENT, GOBP_ATRIAL_CARDIAC_MUSCLE_TISSUE_DEVELOPMENT
	GOBP_OTIC_VESICLE_FORMATION, GOBP_OTIC_VESICLE_FORMATION
	GOBP_VENOUS_BLOOD_VESSEL_MORPHOGENESIS, GOBP_VENOUS_BLOOD_VESSEL_MORPHOGENESIS
	GOBP_POSITIVE_REGULATION_OF_MUSCLE_CONTRACTION, GOBP_POSITIVE_REGULATION_OF_MUSCLE_CONTRACTION
	MODULE_381, MODULE_381
	GOBP_REGULATION_OF_BILE_ACID_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_BILE_ACID_BIOSYNTHETIC_PROCESS
	HP_OPEN_ANGLE_GLAUCOMA, HP_OPEN_ANGLE_GLAUCOMA
	GOBP_ATRIAL_CARDIAC_MUSCLE_TISSUE_MORPHOGENESIS, GOBP_ATRIAL_CARDIAC_MUSCLE_TISSUE_MORPHOGENESIS
	HP_CIGARETTE_PAPER_SCARS, HP_CIGARETTE_PAPER_SCARS
	HP_ABNORMALITY_OF_THE_DISTAL_PHALANX_OF_THE_THUMB, HP_ABNORMALITY_OF_THE_DISTAL_PHALANX_OF_THE_THUMB
	GOBP_REGULATION_OF_MUCUS_SECRETION, GOBP_REGULATION_OF_MUCUS_SECRETION
	GOBP_ADENYLATE_CYCLASE_INHIBITING_G_PROTEIN_COUPLED_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY, GOBP_ADENYLATE_CYCLASE_INHIBITING_G_PROTEIN_COUPLED_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY
	GOBP_REGULATION_OF_BILE_ACID_METABOLIC_PROCESS, GOBP_REGULATION_OF_BILE_ACID_METABOLIC_PROCESS
	HP_PREMATURE_LOSS_OF_PERMANENT_TEETH, HP_PREMATURE_LOSS_OF_PERMANENT_TEETH
	GOBP_LENS_FIBER_CELL_MORPHOGENESIS, GOBP_LENS_FIBER_CELL_MORPHOGENESIS
	GOBP_REGULATION_OF_SKELETAL_MUSCLE_CONTRACTION_BY_CALCIIUM_ION_SIGNALING, GOBP_REGULATION_OF_SKELETAL_MUSCLE_CONTRACTION_BY_CALCIIUM_ION_SIGNALING
	GOBP_NEGATIVE_REGULATION_OF_CIRCADIAN_SLEEP_WAKE_CYCLE_SLEEP, GOBP_NEGATIVE_REGULATION_OF_CIRCADIAN_SLEEP_WAKE_CYCLE_SLEEP
	GOBP_POSITIVE_REGULATION_OF_SARCOMERE_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_SARCOMERE_ORGANIZATION
	GOBP_REGULATION_OF_SKELETAL_MUSCLE_CONTRACTION, GOBP_REGULATION_OF_SKELETAL_MUSCLE_CONTRACTION
	GOMF_G_PROTEIN_COUPLED_GLUTAMATE_RECEPTOR_ACTIVITY, GOMF_G_PROTEIN_COUPLED_GLUTAMATE_RECEPTOR_ACTIVITY
	GOBP_REGULATION_OF_SARCOMERE_ORGANIZATION, GOBP_REGULATION_OF_SARCOMERE_ORGANIZATION
	WP_GPCRS_CLASS_C_METABOTROPIC_GLUTAMATE_PHEROMONE, WP_GPCRS_CLASS_C_METABOTROPIC_GLUTAMATE_PHEROMONE
	MODULE_153, MODULE_153