

	GSE37532_CTRL_VS_TCONV_PPARG_KO_CD4_TCELL_FROM_LN_DN, GSE37532_TREG_VS_TCONV_PPARG_KO_CD4_TCELL_FROM_LN_DN GSE17974_05H_VS_72H_IL4_AND_ANTIL_IL12_ACT_CD4_TCELL_DN, GSE17974_05H_VS_72H_IL4_AND_ANTIL_IL12_ACT_CD4_TCELL_DN GSE17974_CTRL_VS_ACT_IL4_AND_ANTIL_IL12_72H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTIL_IL12_72H_CD4_TCELL_DN GSE17974_CTRL_VS_ACT_IL4_AND_ANTIL_IL12_6H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTIL_IL12_6H_CD4_TCELL_DN GSE14415_INDUCED_TREG_VS_FAILED_INDUCED_TREG_DN, GSE14415_INDUCED_TREG_VS_FAILED_INDUCED_TREG_DN GSE6092_CTRL_VS_BORRELIA_BIRGDOFERI_INF_ENDOTHELIAL_CELL_UP, GSE6092_CTRL_VS_BORRELIA_BIRGDOFERI_INF_ENDOTHELIAL_CELL_UP GSE21678_WT_VS_FOXO1_FOXO3_KO_TREG_DN, GSE21678_WT_VS_FOXO1_FOXO3_KO_TREG_DN GSE11864_CSFL_VS_CSFL_IFNG_IN_MAC_DN, GSE11864_CSFL_VS_CSFL_IFNG_IN_MAC_DN GSE3565_CTRL_VS_LPS_INJECTED_DUSP1_KO_SPLENOCYTES_UP, GSE3565_CTRL_VS_LPS_INJECTED_DUSP1_KO_SPLENOCYTES_UP GSE15930_STIM_VS_STIM_AND_IFNAB_48H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_IFNAB_48H_CD8_T_CELL_DN HALLMARK_ESTROGEN_RESPONSE_LATE, HALLMARK_ESTROGEN_RESPONSE_LATE DARWICHE_PAPILLOMA_RISK_LOW_DN, DARWICHE_PAPILLOMA_RISK_LOW_DN HP_CARIOUS_TEETH, HP_CARIOUS_TEETH ZHAN_MULTIPLE_MYELOMA_UP, ZHAN_MULTIPLE_MYELOMA_UP GSE360_L_MAJOR_VS_T_GONDIL_DC_DN, GSE360_L_MAJOR_VS_T_GONDIL_DC_DN HP_LACRIMATION_ABNORMALITY, HP_LACRIMATION_ABNORMALITY KIM_GASTRIC_CANCER_CHEMOSENSITIVITY, KIM_GASTRIC_CANCER_CHEMOSENSITIVITY HP_PULMONARY_ARTERY_STENOSIS, HP_PULMONARY_ARTERY_STENOSIS GSE15930_STIM_VS_STIM_AND_IL12_24H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_IL12_24H_CD8_T_CELL_DN HP_PETECHIAE, HP_PETECHIAE CHEN_HOXA5_TARGETS_9HR_DN, CHEN_HOXA5_TARGETS_9HR_DN HP_PERSONALITY_CHANGES, HP_PERSONALITY_CHANGES HP_MICRODONTIA, HP_MICRODONTIA HP_CHILDHOOD_ONSET, HP_CHILDHOOD_ONSET LEE_TARGETS_OF_PTCH1_AND_SUFU_UP, LEE_TARGETS_OF_PTCH1_AND_SUFU_UP HP_HYPOPLASIA_OF_THE_ZYGOMATIC_BONE, HP_HYPOPLASIA_OF_THE_ZYGOMATIC_BONE HP_CNS_DEMYELINATION, HP_CNS_DEMYELINATION HP_RECTAL_PROLAPSE, HP_RECTAL_PROLAPSE ZHAN_MULTIPLE_MYELOMA_CD2_DN, ZHAN_MULTIPLE_MYELOMA_CD2_DN BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_DN, BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_DN REACTOME_GAMMA_CARBOXYLATION_HYPUSINE_FORMATION_AND_ARYLSULFATASE_ACTIVATION, REACTOME_GAMMA_CARBOXYLATION_HYPUSINE_FORMATION_AND_ARYLSULFATASE_ACTIVATION MIR4772_5P, MIR4772_5P MODULE_138, MODULE_138 GOBP_T_CELL_MEDIATED_CYTOTOXICITY, GOBP_T_CELL_MEDIATED_CYTOTOXICITY MIR6759_5P, MIR6759_5P HP_HYPOPLASTIC_FACIAL_BONES, HP_HYPOPLASTIC_FACIAL_BONES GOBP_MANNOSYLATION, GOBP_MANNOSYLATION GOBP_PURINE_NUCLEOBASE_BIOSYNTHETIC_PROCESS, GOBP_PURINE_NUCLEOBASE_BIOSYNTHETIC_PROCESS GOMF_MANNOSYLTRANSFERASE_ACTIVITY, GOMF_MANNOSYLTRANSFERASE_ACTIVITY GOBP_RESPONSE_TO_ISCHEMIA, GOBP_RESPONSE_TO_ISCHEMIA PID_P38_ALPHA_BETA_DOWNSTREAM_PATHWAY, PID_P38_ALPHA_BETA_DOWNSTREAM_PATHWAY GOBP_PURINE_NUCLEOBASE_METABOLIC_PROCESS, GOBP_PURINE_NUCLEOBASE_METABOLIC_PROCESS HP_ATROPHY_DEGENERATION_INVOLVING_THE_CORTICOSPINAL_TRACTS, HP_ATROPHY_DEGENERATION_INVOLVING_THE_CORTICOSPINAL_TRACTS GOMF_CARBON_CARBON_LYASE_ACTIVITY, GOMF_CARBON_CARBON_LYASE_ACTIVITY GOBP_GAMMA_DELTA_T_CELL_ACTIVATION, GOBP_GAMMA_DELTA_T_CELL_ACTIVATION MIR652_5P, MIR652_5P HP_ABNORMALITY_OF_BLADDER_MORPHOLOGY, HP_ABNORMALITY_OF_BLADDER_MORPHOLOGY GOMF_TRANSFORMING_GROWTH_FACTOR_BETA_BINDING, GOMF_TRANSFORMING_GROWTH_FACTOR_BETA_BINDING SCHERER_PBM_C_APSV_WETVAX_AGE_18_40YO_5_TO_7DY_UP, SCHERER_PBM_C_APSV_WETVAX_AGE_18_40YO_5_TO_7DY_UP HP_INVERTED_NIPPLES, HP_INVERTED_NIPPLES GOBP_PROTEIN_HOMOTETRAMERIZATION, GOBP_PROTEIN_HOMOTETRAMERIZATION BONOME_OVARIAN_CANCER_POOR_SURVIVAL_DN, BONOME_OVARIAN_CANCER_POOR_SURVIVAL_DN SOBOLEV_PBM_C_PANDEMRIX_AGE_18_64YO_CORRELATED_WITH_AGE_1DY_POSITIVE, SOBOLEV_PBM_C_PANDEMRIX_AGE_18_64YO_CORRELATED_WITH_AGE_1DY_POSITIVE SOBOLEV_PBM_C_PANDEMRIX_AGE_18_64YO_RESPONDERS_VS_NONRESPONDERS_7DY_UP, SOBOLEV_PBM_C_PANDEMRIX_AGE_18_64YO_RESPONDERS_VS_NONRESPONDERS_7DY_UP MIR219B_3P, MIR219B_3P GOBP_PROTEIN_MATURATION_BY_IRON_SULFUR_CLUSTER_TRANSFER, GOBP_PROTEIN_MATURATION_BY_IRON_SULFUR_CLUSTER_TRANSFER GOMF_FATTY_ACID_LIGASE_ACTIVITY, GOMF_FATTY_ACID_LIGASE_ACTIVITY LEIN_CEREBELLUM_MARKERS, LEIN_CEREBELLUM_MARKERS HP_PERIPHERAL_PULMONARY_ARTERY_STENOSIS, HP_PERIPHERAL_PULMONARY_ARTERY_STENOSIS WANG_RECURRENT_LIVER_CANCER_DN, WANG_RECURRENT_LIVER_CANCER_DN HP_PELVIC_KIDNEY, HP_PELVIC_KIDNEY GOMF_EPHRIN_RECEPTOR_BINDING, GOMF_EPHRIN_RECEPTOR_BINDING MIR9903, MIR9903 HP_VISUAL_HALLUCINATIONS, HP_VISUAL_HALLUCINATIONS GOBP_PROTEIN_MANNOSYLATION, GOBP_PROTEIN_MANNOSYLATION HP_SUPRAVALVULAR_AORTIC_STENOSIS, HP_SUPRAVALVULAR_AORTIC_STENOSIS GOBP_PROTEIN_O_LINKED_MANNOSYLATION, GOBP_PROTEIN_O_LINKED_MANNOSYLATION DESERT_EXTRACELLULAR_MATRIX_HEPATOCELLULAR_CARCINOMA_SUBCLASS_UP, DESERT_EXTRACELLULAR_MATRIX_HEPATOCELLULAR_CARCINOMA_SUBCLASS_UP GOBP_RESPONSE_TO_INTERLEUKIN_15, GOBP_RESPONSE_TO_INTERLEUKIN_15 GOBP_PROSTATE_GLANDULAR_ACINUS_DEVELOPMENT, GOBP_PROSTATE_GLANDULAR_ACINUS_DEVELOPMENT WP_DEVELOPMENT_AND_HETEROGENEITY_OF_THE_ILC_FAMILY, WP_DEVELOPMENT_AND_HETEROGENEITY_OF_THE_ILC_FAMILY ZWANG_EGF_INTERVAL_UP, ZWANG_EGF_INTERVAL_UP WP_OMEGA3OMEGA6_FA_SYNTHESIS, WP_OMEGA3OMEGA6_FA_SYNTHESIS HP_APLASTIC_ANEMIA, HP_APLASTIC_ANEMIA REACTOME_BETA_OXIDATION_OF_VERY_LONG_CHAIN_FATTY_ACIDS, REACTOME_BETA_OXIDATION_OF_VERY_LONG_CHAIN_FATTY_ACIDS WEST_ADRENOCORTICAL_CARCINOMA_VS_ADENOMA_UP, WEST_ADRENOCORTICAL_CARCINOMA_VS_ADENOMA_UP HP_ABNORMAL_FULL_FIELD_ELECTRORETINOGRAM, HP_ABNORMAL_FULL_FIELD_ELECTRORETINOGRAM HP_SECONDARY_AMENORRHEA, HP_SECONDARY_AMENORRHEA PIONTEK_PKD1_TARGETS_UP, PIONTEK_PKD1_TARGETS_UP STREICHER_LSM1_TARGETS_DN, STREICHER_LSM1_TARGETS_DN HP_ACNE_INVERSA, HP_ACNE_INVERSA CHIANG_LIVER_CANCER_SUBCLASS_POLYSOMY7_UP, CHIANG_LIVER_CANCER_SUBCLASS_POLYSOMY7_UP HP_PHONOPHOBIA, HP_PHONOPHOBIA PALOMERO_GSL_SENSITIVITY_UP, PALOMERO_GSL_SENSITIVITY_UP HP_RETINAL_ARTERIOLAR_TORTUOSITY, HP_RETINAL_ARTERIOLAR_TORTUOSITY EGFR_TARGET_GENES, EGFR_TARGET_GENES HP_SEVERELY_REDUCED_VISUAL_ACUITY, HP_SEVERELY_REDUCED_VISUAL_ACUITY GOMF_GALACTOSYLTRANSFERASE_ACTIVITY, GOMF_GALACTOSYLTRANSFERASE_ACTIVITY MIR6854_3P, MIR6854_3P HP_ABNORMAL_POSTURING, HP_ABNORMAL_POSTURING HP_PILOMATRIXOMA, HP_PILOMATRIXOMA CARDOSO_RESPONSE_TO_GAMMA_RADIATION_AND_3AB, CARDOSO_RESPONSE_TO_GAMMA_RADIATION_AND_3AB GOBP_SEGMENTATION, GOBP_SEGMENTATION GOBP_VERY_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS, GOBP_VERY_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS GOMF_LONG_CHAIN_FATTY_ACID_COA_LIGASE_ACTIVITY, GOMF_LONG_CHAIN_FATTY_ACID_COA_LIGASE_ACTIVITY GOBP_EMBRYONIC_HEART_TUBE_MORPHOGENESIS, GOBP_EMBRYONIC_HEART_TUBE_MORPHOGENESIS TSENG_ADIPOGENIC_POTENTIAL_UP, TSENG_ADIPOGENIC_POTENTIAL_UP GOMF_PROTEIN_BINDING_INVOLVED_IN_HETEROTYPIC_CELL_CELL_ADHESION, GOMF_PROTEIN_BINDING_INVOLVED_IN_HETEROTYPIC_CELL_CELL_ADHESION GOMF_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY GOMF_PYRIMIDINE_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_PYRIMIDINE_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY HP_GENERALIZED_DYSTONIA, HP_GENERALIZED_DYSTONIA ZWANG_CLASS_2_TRANSIENTLY_INDUCED_BY_EGF, ZWANG_CLASS_2_TRANSIENTLY_INDUCED_BY_EGF GOBP_LOCOMOTION_INVOLVED_IN_LOCOMOTORY_BEHAVIOR, GOBP_LOCOMOTION_INVOLVED_IN_LOCOMOTORY_BEHAVIOR GOBP_EPITHELIAL_CELL_PROLIFERATION_INVOLVED_IN_PROSTATE_GLAND_DEVELOPMENT, GOBP_EPITHELIAL_CELL_PROLIFERATION_INVOLVED_IN_PROSTATE_GLAND_DEVELOPMENT GOMF_METHYL_CPG_BINDING, GOMF_METHYL_CPG_BINDING GOBP_RESPONSE_TO_LEAD_ION, GOBP_RESPONSE_TO_LEAD_ION HP_PROMINENT_STERNUM, HP_PROMINENT_STERNUM GOBP_POSITIVE_REGULATION_OF_HUMORAL_IMMUNE_RESPONSE_MEDIATED_BY_CIRCULATING_IMMUNOGLOBULIN, GOBP_POSITIVE_REGULATION_OF_HUMORAL_IMMUNE_RESPONSE_MEDIATED_BY_CIRCULATING_IMMUNOGLOBULIN GOBP_LONG_CHAIN_FATTY_ACID_IMPORT_INTO_CELL, GOBP_LONG_CHAIN_FATTY_ACID_IMPORT_INTO_CELL GOBP_REGULATION_OF_HEMOGLOBIN_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_HEMOGLOBIN_BIOSYNTHETIC_PROCESS HP_AORTIC_ARCH_ANEURYSM, HP_AORTIC_ARCH_ANEURYSM BIOCARTA_BTG2_PATHWAY, BIOCARTA_BTG2_PATHWAY GOBP_EMBRYONIC_HEART_TUBE_DEVELOPMENT, GOBP_EMBRYONIC_HEART_TUBE_DEVELOPMENT GOMF_NATURAL_KILLER_CELL_LECTIN_LIKE_RECEPTOR_BINDING, GOMF_NATURAL_KILLER_CELL_LECTIN_LIKE_RECEPTOR_BINDING GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_METABOLIC_PROCESS MIR6506_3P, MIR6506_3P GOBP_NEGATIVE_REGULATION_OF_T_CELL_MIGRATION, GOBP_NEGATIVE_REGULATION_OF_T_CELL_MIGRATION REACTOME_PRE_NOTCH_PROCESSING_IN_THE_ENDOPLASMIC_RETICULUM, REACTOME_PRE_NOTCH_PROCESSING_IN_THE_ENDOPLASMIC_RETICULUM GOBP_PROGRAMMED_CELL_DEATH_IN_RESPONSE_TO_REACTIVE_OXYGEN_SPECIES, GOBP_PROGRAMMED_CELL_DEATH_IN_RESPONSE_TO_REACTIVE_OXYGEN_SPECIES HP_ABNORMAL_LIGHT_AND_DARK_ADAPTED_ELECTRORETINOGRAM, HP_ABNORMAL_LIGHT_AND_DARK_ADAPTED_ELECTRORETINOGRAM GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_BIOSYNTHETIC_PROCESS GOBP_SPHINGOMYELIN_METABOLIC_PROCESS, GOBP_SPHINGOMYELIN_METABOLIC_PROCESS GOBP_MANGANESE_ION_TRANSPORT, GOBP_MANGANESE_ION_TRANSPORT GOMF_VERY_LONG_CHAIN_FATTY_ACID_COA_LIGASE_ACTIVITY, GOMF_VERY_LONG_CHAIN_FATTY_ACID_COA_LIGASE_ACTIVITY THILLAINADESAN_ZNF217_TARGETS_DN, THILLAINADESAN_ZNF217_TARGETS_DN HP_ABNORMALITY_OF_THE_GLENOID_FOSSA, HP_ABNORMALITY_OF_THE_GLENOID_FOSSA HP_FOOD_INTOLERANCE, HP_FOOD_INTOLERANCE GOBP_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORT, GOBP_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORT HP_PATELLAR_HYPOPLASIA, HP_PATELLAR_HYPOPLASIA MODULE_148, MODULE_148 WP_GAMMAGLUTAMYL_CYCLE_FOR_THE_BIOSYNTHESIS_AND_DEGRADATION_OF_GLUTATHIONE_INCLUDING_DISEASES, WP_GAMMAGLUTAMYL_CYCLE_FOR_THE_BIOSYNTHESIS_AND_DEGRADATION_OF_GLUTATHIONE_INCLUDING_DISEASES GOCC_EXTRINSIC_COMPONENT_OF_ENDOSOME_MEMBRANE, GOCC_EXTRINSIC_COMPONENT_OF_ENDOSOME_MEMBRANE HP_CAUDATE_ATROPHY, HP_CAUDATE_ATROPHY BARRIER_CANCER_RELAPSE_NORMAL_SAMPLE_UP, BARRIER_CANCER_RELAPSE_NORMAL_SAMPLE_UP GOBP_C_TERMINAL_PROTEIN_AMINO_ACID_MODIFICATION, GOBP_C_TERMINAL_PROTEIN_AMINO_ACID_MODIFICATION HP_PLANTAR_HYPERKERATOSIS, HP_PLANTAR_HYPERKERATOSIS GOBP_MACROPHAGE_INFLAMMATORY_PROTEIN_1_ALPHA_PRODUCTION, GOBP_MACROPHAGE_INFLAMMATORY_PROTEIN_1_ALPHA_PRODUCTION HP_BICORONAL_SYNOSTOSIS, HP_BICORONAL_SYNOSTOSIS GOBP_PYRIMIDINE_NUCLEOSIDE_TRANSPORT, GOBP_PYRIMIDINE_NUCLEOSIDE_TRANSPORT GOMF_URIDINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_URIDINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY HP_WIDE_PUBIC_SYMPHYSIS, HP_WIDE_PUBIC_SYMPHYSIS GOBP_POSITIVE_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS GOBP_EMBRYONIC_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY, GOBP_EMBRYONIC_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY GOMF_NAADP_SENSITIVE_CALCIUM_RELEASE_CHANNEL_ACTIVITY, GOMF_NAADP_SENSITIVE_CALCIUM_RELEASE_CHANNEL_ACTIVITY HP_PERINEAL_FISTULA, HP_PERINEAL_FISTULA GOBP_FLOOR_PLATE_DEVELOPMENT, GOBP_FLOOR_PLATE_DEVELOPMENT GOMF_LYSOPHOSPHATIDIC_ACID_BINDING, GOMF_LYSOPHOSPHATIDIC_ACID_BINDING HP_BRACHYTURRICEPHALY, HP_BRACHYTURRICEPHALY HP_HYPERGRANULOSIS, HP_HYPERGRANULOSIS ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_HIGH_ICM_RESPONDERS_3DY_7D_DN, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_HIGH_ICM_RESPONDERS_3DY_7D_DN GOBP_AMINO_ACID_HOMEOSTASIS, GOBP_AMINO_ACID_HOMEOSTASIS REACTOME_SCAVENGING_BY_CLASS_F_RECEPTORS, REACTOME_SCAVENGING_BY_CLASS_F_RECEPTORS GOBP_CORNEA_DEVELOPMENT_IN_CAMERA_TYPE_EYE, GOBP_CORNEA_DEVELOPMENT_IN_CAMERA_TYPE_EYE GOBP_RESPONSE_TO_PHEROMONE, GOBP_RESPONSE_TO_PHEROMONE GOBP_SPHINGOMYELIN_BIOSYNTHETIC_PROCESS, GOBP_SPHINGOMYELIN_BIOSYNTHETIC_PROCESS HP_VEGETATIVE_STATE, HP_VEGETATIVE_STATE HP_FLARING_OF_RIB_CAGE, HP_FLARING_OF_RIB_CAGE BIOCARTA_HSP27_PATHWAY, BIOCARTA_HSP27_PATHWAY GOMF_MAP_KINASE_PHOSPHATASE_ACTIVITY, GOMF_MAP_KINASE_PHOSPHATASE_ACTIVITY HP_REDUCED_DELAYED_HYPERSENSITIVITY, HP_REDUCED_DELAYED_HYPERSENSITIVITY GOBP_GLUTAMATE_HOMEOSTASIS, GOBP_GLUTAMATE_HOMEOSTASIS GOMF_SULFATE_BINDING, GOMF_SULFATE_BINDING HP_ALVEOLAR_RHABDOMYOSARCOMA, HP_ALVEOLAR_RHABDOMYOSARCOMA HP_NEOPLASM_OF_THE_TRACHEOBRONCHIAL_SYSTEM, HP_NEOPLASM_OF_THE_TRACHEOBRONCHIAL_SYSTEM GOBP_NEURON_REMODELING, GOBP_NEURON_REMODELING REACTOME_POU5F1_OCT4_SOX2_NANOG_REPRESS_GENES_RELATED_TO_DIFFERENTIATION, REACTOME_POU5F1_OCT4_SOX2_NANOG_REPRESS_GENES_RELATED_TO_DIFFERENTIATION HP_ABSENT_BRAINSTEM_AUDITORY_RESPONSES, HP_ABSENT_BRAINSTEM_AUDITORY_RESPONSES GOBP_VERY_LONG_CHAIN_FATTY_ACID_CATABOLIC_PROCESS, GOBP_VERY_LONG_CHAIN_FATTY_ACID_CATABOLIC_PROCESS GOBP_ENDODERMAL_CELL_FATE_SPECIFICATION, GOBP_ENDODERMAL_CELL_FATE_SPECIFICATION	
2H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_12H_IN_VITRO_ACT_CD4_TCELL_DN		