

G_TREATED_ENDOTHELIAL_CELL_DN,GSE3920_UNTREATED_VS_IFNG_TREATED_ENDOTHELIAL_CELL_DN

GSE17721_POLYIC_VS_CPG_1H_BMDC_UP,GSE17721_POLYIC_VS_CPG_1H_BMDC_UP
GSE27434_WT_VS_DNMT1_KO_TREG_UP,GSE27434_WT_VS_DNMT1_KO_TREG_UP
GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4MID_BCELL_DN,GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4MID_BCELL_DN
GSE27859_MACROPHAGE_VS_CD11C_INT_F480_HI_MACROPHAGE_DN,GSE27859_MACROPHAGE_VS_CD11C_INT_F480_HI_MACROPHAGE_DN
GSE6092_CTRL_VS_BORRELIA_BIRGDOFERI_INF_ENDOTHELIAL_CELL_UP,GSE6092_CTRL_VS_BORRELIA_BIRGDOFERI_INF_ENDOTHELIAL_CELL_UP
GSE20715_WT_VS_TLR4_KO_6H_OZONE_LUNG_UP,GSE20715_WT_VS_TLR4_KO_6H_OZONE_LUNG_UP
GSE8685_IL15_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP,GSE8685_IL15_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP
GSE22611_UNSTIM_VS_6H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP,GSE22611_UNSTIM_VS_6H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP
GSE40274_CTRL_VS_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP,GSE40274_CTRL_VS_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
REACTOME_AUTOPHAGY,REACTOME_AUTOPHAGY
GSE18281_PERIMEDULLARY_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_UP,GSE18281_PERIMEDULLARY_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_UP
GSE23321_CENTRAL_VS_EFFECTOR_MEMORY_CD8_TCELL_UP,GSE23321_CENTRAL_VS_EFFECTOR_MEMORY_CD8_TCELL_UP
MIR939_3P,MIR939_3P
GSE10240_IL17_VS_IL17_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN,GSE10240_IL17_VS_IL17_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN
GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_IRF8_KO_UP,GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_IRF8_KO_UP
REACTOME_GLYCEROPHOSPHOLIPID_BIOSYNTHESIS,REACTOME_GLYCEROPHOSPHOLIPID_BIOSYNTHESIS
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_12,YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_12
MIR6765_3P,MIR6765_3P
GSE6259_33D1_POS_DC_VS_BCELL_UP,GSE6259_33D1_POS_DC_VS_BCELL_UP
KMCATNNWGGA_UNKNOWN,KMCATNNWGGA_UNKNOWN
GOBP_MULTIVESICULAR_BODY_SORTING_PATHWAY,GOBP_MULTIVESICULAR_BODY_SORTING_PATHWAY
HALLMARK_BILE_ACID_METABOLISM,HALLMARK_BILE_ACID_METABOLISM
GOBP_PHOSPHATIDYLCHOLINE_METABOLIC_PROCESS,GOBP_PHOSPHATIDYLCHOLINE_METABOLIC_PROCESS
GOBP_ENDOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY,GOBP_ENDOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY
MIR4685_5P,MIR4685_5P
GOBP_BRANCHED_CHAIN_AMINO_ACID_METABOLIC_PROCESS,GOBP_BRANCHED_CHAIN_AMINO_ACID_METABOLIC_PROCESS
ADNP_TARGET_GENES,ADNP_TARGET_GENES
MIR6754_5P,MIR6754_5P
GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4_KO_BCELL_UP,GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4_KO_BCELL_UP
MIR6748_5P,MIR6748_5P
GOBP_PROTEIN_LOCALIZATION_TO_CELL_SURFACE,GOBP_PROTEIN_LOCALIZATION_TO_CELL_SURFACE
MIR6834_3P,MIR6834_3P
GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY,GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY
HP_NASOGASTRIC_TUBE_FEEDING,HP_NASOGASTRIC_TUBE_FEEDING
STARK_HYPOCAMPUS_22Q11_DELETION_UP,STARK_HYPOCAMPUS_22Q11_DELETION_UP
MIR331_5P,MIR331_5P
MIR1225_3P,MIR1225_3P
MIR3675_5P,MIR3675_5P
REACTOME_TRANSCRIPTIONAL_REGULATION_BY_MECP2,REACTOME_TRANSCRIPTIONAL_REGULATION_BY_MECP2
GOBP_GENITALIA_DEVELOPMENT,GOBP_GENITALIA_DEVELOPMENT
MIR7106_3P,MIR7106_3P
BIOCARTA_IL6_PATHWAY,BIOCARTA_IL6_PATHWAY
HP_ABNORMAL_EATING_BEHAVIOR,HP_ABNORMAL_EATING_BEHAVIOR
HP_BROAD_PHILTRUM,HP_BROAD_PHILTRUM
GOBP_NUCLEOSIDE_TRIPHOSPHATE_CATABOLIC_PROCESS,GOBP_NUCLEOSIDE_TRIPHOSPHATE_CATABOLIC_PROCESS
GOMF_NUCLEOSIDE_TRIPHOSPHATE_DIPHOSPHATASE_ACTIVITY,GOMF_NUCLEOSIDE_TRIPHOSPHATE_DIPHOSPHATASE_ACTIVITY
GOCC_ENDOPLASMIC_RETICULUM_PALMITOYLTRANSFERASE_COMPLEX,GOCC_ENDOPLASMIC_RETICULUM_PALMITOYLTRANSFERASE_COMPLEX
GOBP_PHOSPHATIDIC_ACID_METABOLIC_PROCESS,GOBP_PHOSPHATIDIC_ACID_METABOLIC_PROCESS
MIR5739,MIR5739
GOBP_HORMONE_BIOSYNTHETIC_PROCESS,GOBP_HORMONE_BIOSYNTHETIC_PROCESS
HP_GINGIVAL_FIBROMATOSIS,HP_GINGIVAL_FIBROMATOSIS
GOCC_PALMITOYLTRANSFERASE_COMPLEX,GOCC_PALMITOYLTRANSFERASE_COMPLEX
TACGGGT_MIR99A_MIR100_MIR99B,TACGGGT_MIR99A_MIR100_MIR99B
HP_EEG_WITH_BURST_SUPPRESSION,HP_EEG_WITH_BURST_SUPPRESSION
ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_6HR_UP,ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_6HR_UP
REACTOME_RETINOID_CYCLE_DISEASE_EVENTS,REACTOME_RETINOID_CYCLE_DISEASE_EVENTS
HP_PULMONARY_HEMORRHAGE,HP_PULMONARY_HEMORRHAGE
GOBP_PURINE_DEOXYRIBONUCLEOTIDE_METABOLIC_PROCESS,GOBP_PURINE_DEOXYRIBONUCLEOTIDE_METABOLIC_PROCESS
GSE13887_ACT_CD4_VS_NO_TREATED_CD4_TCELL_DN,GSE13887_ACT_CD4_VS_NO_TREATED_CD4_TCELL_DN
WP_NANOPARTICLE_TRIGGERED_AUTOPHAGIC_CELL_DEATH,WP_NANOPARTICLE_TRIGGERED_AUTOPHAGIC_CELL_DEATH
GOBP_PURINE_DEOXYRIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS,GOBP_PURINE_DEOXYRIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
HP_AGENESIS_OF_PERMANENT_TEETH,HP_AGENESIS_OF_PERMANENT_TEETH
GOBP_ANDROGEN_BIOSYNTHETIC_PROCESS,GOBP_ANDROGEN_BIOSYNTHETIC_PROCESS
GOBP_SYNAPTIC_VESICLE_DOCKING,GOBP_SYNAPTIC_VESICLE_DOCKING
KAZMIN_PBMCP_FALCIPARUM_RTSS_AS01_UNKN_AGE_IMM_WITH_ARR_VS_IMM_BY_RRR_PRIMARY_IMMUNIZ_WITH_RECOMB_ADENOVIRUS_35_1DY_UP,KAZMIN_PBMCP_FALCIPARUM_RTSS_AS01_UNKN_AGE_IMM_WITH_ARR_VS_IMM_BY_RRR_PRIMARY_IMMUNIZ_WITH_RECOMB_ADENOVIRUS_35_1DY_UP
GOMF_SPHINGOSINE_1_PHOSPHATE_PHOSPHATASE_ACTIVITY,GOMF_SPHINGOSINE_1_PHOSPHATE_PHOSPHATASE_ACTIVITY
HP_ABNORMALITY_OF_THE_ANTITRAGUS,HP_ABNORMALITY_OF_THE_ANTITRAGUS
GOMF_DOPAMINE_RECEPTOR_BINDING,GOMF_DOPAMINE_RECEPTOR_BINDING
GOBP_NEGATIVE_REGULATION_OF_INFLAMMATORY_RESPONSE,GOBP_NEGATIVE_REGULATION_OF_INFLAMMATORY_RESPONSE
GOBP_VAGINA_DEVELOPMENT,GOBP_VAGINA_DEVELOPMENT
HP_VERTICAL_NYSTAGMUS,HP_VERTICAL_NYSTAGMUS
GOBP_PURINE_DEOXYRIBONUCLEOTIDE_CATABOLIC_PROCESS,GOBP_PURINE_DEOXYRIBONUCLEOTIDE_CATABOLIC_PROCESS
WP_IRINOTECAN_PATHWAY,WP_IRINOTECAN_PATHWAY
GOBP_REGULATION_OF_EATING_BEHAVIOR,GOBP_REGULATION_OF_EATING_BEHAVIOR
BIOCARTA_ARAP_PATHWAY,BIOCARTA_ARAP_PATHWAY
GOBP_SECRETION_OF_LYSOSOMAL_ENZYMES,GOBP_SECRETION_OF_LYSOSOMAL_ENZYMES
GOBP_NEGATIVE_REGULATION_OF_RELAXATION_OF_MUSCLE,GOBP_NEGATIVE_REGULATION_OF_RELAXATION_OF_MUSCLE
GOCC_PHOTORECEPTOR_OUTER_SEGMENT_MEMBRANE,GOCC_PHOTORECEPTOR_OUTER_SEGMENT_MEMBRANE
GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_SYNAPSE,GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_SYNAPSE
GOBP_NEGATIVE_REGULATION_OF_STRIATED_MUSCLE_CONTRACTION,GOBP_NEGATIVE_REGULATION_OF_STRIATED_MUSCLE_CONTRACTION
GOCC_INTEGRAL_COMPONENT_OF_LYSOSOMAL_MEMBRANE,GOCC_INTEGRAL_COMPONENT_OF_LYSOSOMAL_MEMBRANE
REACTOME_THE_CANONICAL_RETINOID_CYCLE_IN_RODS_TWILIGHT_VISION,REACTOME_THE_CANONICAL_RETINOID_CYCLE_IN_RODS_TWILIGHT_VISION
GOBP_ISOPRENOID_CATABOLIC_PROCESS,GOBP_ISOPRENOID_CATABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_NODAL_SIGNALING_PATHWAY,GOBP_NEGATIVE_REGULATION_OF_NODAL_SIGNALING_PATHWAY
HP_ABNORMAL_RAPID_EYE_MOVEMENT_SLEEP,HP_ABNORMAL_RAPID_EYE_MOVEMENT_SLEEP
GOBP_PROTEIN_LOCALIZATION_TO_NON_MOTILE_CILIUM,GOBP_PROTEIN_LOCALIZATION_TO_NON_MOTILE_CILIUM
HP_ABNORMAL_ENERGY_EXPENDITURE,HP_ABNORMAL_ENERGY_EXPENDITURE
HP_APNEIC_EPISODES_IN_INFANCY,HP_APNEIC_EPISODES_IN_INFANCY
GOBP_PROTEIN_LOCALIZATION_TO_PHOTORECEPTOR_OUTER_SEGMENT,GOBP_PROTEIN_LOCALIZATION_TO_PHOTORECEPTOR_OUTER_SEGMENT
GOMF_PROTEIN_GLUTAMINE_GAMMA_GLUTAMYLTRANSFERASE_ACTIVITY,GOMF_PROTEIN_GLUTAMINE_GAMMA_GLUTAMYLTRANSFERASE_ACTIVITY
GOBP_NEGATIVE_REGULATION_OF_FEEDING_BEHAVIOR,GOBP_NEGATIVE_REGULATION_OF_FEEDING_BEHAVIOR
HP_ABNORMAL_CELLULAR_COMPOSITION_OF_BRONCHOALVEOLAR_FLUID,HP_ABNORMAL_CELLULAR_COMPOSITION_OF_BRONCHOALVEOLAR_FLUID
GOMF_ENONE_REDUCTASE_ACTIVITY,GOMF_ENONE_REDUCTASE_ACTIVITY
HP_DUPLICATION_OF_RENAL_PELVIS,HP_DUPLICATION_OF_RENAL_PELVIS
GOMF_ALPHA_2A_ADRENERGIC_RECEPTOR_BINDING,GOMF_ALPHA_2A_ADRENERGIC_RECEPTOR_BINDING