

A2AR_AGONIST_TREATED_TREG_UP, GSE34006_UNTREATED_VS_A2AR_AGONIST_TREATED_TREG_UP

chr15q21, chr15q21

GO_GDP_BINDING, GO_GDP_BINDING

GSE19941_UNSTIM_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_DN, GSE19941_UNSTIM_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_DN

MARSON_FOXP3_TARGETS_UP, MARSON_FOXP3_TARGETS_UP

GSE32423_CTRL_VS_IL4_MEMORY_CD8_TCELL_UP, GSE32423_CTRL_VS_IL4_MEMORY_CD8_TCELL_UP

GSE4748_CTRL_VS_LPS_STIM_DC_3H_DN, GSE4748_CTRL_VS_LPS_STIM_DC_3H_DN

GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_DC_UP, GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_DC_UP

GSE37301_HEMATOPOIETIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_UP, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_UP

SMIRNOV_RESPONSE_TO_IR_2HR_UP, SMIRNOV_RESPONSE_TO_IR_2HR_UP

GO_NEGATIVE_REGULATION_OF_CYTOPLASMIC_TRANSPORT, GO_NEGATIVE_REGULATION_OF_CYTOPLASMIC_TRANSPORT

GSE17721_CTRL_VS_LPS_0.5H_BMDC_DN, GSE17721_CTRL_VS_LPS_0.5H_BMDC_DN

ADDYA_ERYTHROID_DIFFERENTIATION_BY_HEMIN, ADDYA_ERYTHROID_DIFFERENTIATION_BY_HEMIN

PID_ECADHERIN_NASCENT_AJ_PATHWAY, PID_ECADHERIN_NASCENT_AJ_PATHWAY

KEGG_AXON_GUIDANCE, KEGG_AXON_GUIDANCE

KEGG_ERBB_SIGNALING_PATHWAY, KEGG_ERBB_SIGNALING_PATHWAY

GO_ACTIN_BASED_CELL_PROJECTION, GO_ACTIN_BASED_CELL_PROJECTION

SESTO_RESPONSE_TO_UV_C7, SESTO_RESPONSE_TO_UV_C7

PID_MYC_REPRESS_PATHWAY, PID_MYC_REPRESS_PATHWAY

GO_NEGATIVE_REGULATION_OF_INTRACELLULAR_TRANSPORT, GO_NEGATIVE_REGULATION_OF_INTRACELLULAR_TRANSPORT

GSE2706_R848_VS_R848_AND_LPS_8H_STIM_DC_DN, GSE2706_R848_VS_R848_AND_LPS_8H_STIM_DC_DN

GO_GLIAL_CELL_DEVELOPMENT, GO_GLIAL_CELL_DEVELOPMENT

IKEDA_MIR133_TARGETS_UP, IKEDA_MIR133_TARGETS_UP

ELVIDGE_HYPOXIA_BY_DMOG_UP, ELVIDGE_HYPOXIA_BY_DMOG_UP

GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_UP, GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_UP

GO_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS, GO_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS

GO_SNARE_BINDING, GO_SNARE_BINDING

GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_DN, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_DN

GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_UP, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_UP

GSE9650_NAIVE_VS_EXHAUSTED_CD8_TCELL_DN, GSE9650_NAIVE_VS_EXHAUSTED_CD8_TCELL_DN

LIU_PROSTATE_CANCER_UP, LIU_PROSTATE_CANCER_UP

WANG_RESPONSE_TO_ANDROGEN_UP, WANG_RESPONSE_TO_ANDROGEN_UP

GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_UP, GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_UP

GO_NUCLEAR_TRANSCRIPTION_FACTOR_COMPLEX, GO_NUCLEAR_TRANSCRIPTION_FACTOR_COMPLEX

GROSS_HYPOXIA_VIA_ELK3_AND_HIF1A_UP, GROSS_HYPOXIA_VIA_ELK3_AND_HIF1A_UP

MODULE_312, MODULE_312

FARDIN_HYPOXIA_11, FARDIN_HYPOXIA_11

GSE9960_GRAM_POS_VS_GRAM_NEG_AND_POS_SEPSIS_PBMCDN, GSE9960_GRAM_POS_VS_GRAM_NEG_AND_POS_SEPSIS_PBMCDN

GSE5589_LPS_AND_IL10_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_UP, GSE5589_LPS_AND_IL10_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_UP

GO_REGULATION_OF_STEM_CELL_DIFFERENTIATION, GO_REGULATION_OF_STEM_CELL_DIFFERENTIATION

LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_SMALL_VS_HUGE_UP, LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_SMALL_VS_HUGE_UP

GO_ORGANELLE_MEMBRANE_FUSION, GO_ORGANELLE_MEMBRANE_FUSION

BIOCARTA_P53HYPOXIA_PATHWAY, BIOCARTA_P53HYPOXIA_PATHWAY

GSE17721_CPG_VS_GARDIQUIMOD_2H_BMDC_UP, GSE17721_CPG_VS_GARDIQUIMOD_2H_BMDC_UP

GO_NCRNA_3_END_PROCESSING, GO_NCRNA_3_END_PROCESSING

GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TREATED_CD4_TCELL_DN, GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TREATED_CD4_TCELL_DN

GSE27786_CD4_TCELL_VS_NEUTROPHIL_DN, GSE27786_CD4_TCELL_VS_NEUTROPHIL_DN

CHEN_ETV5_TARGETS_TESTIS, CHEN_ETV5_TARGETS_TESTIS

NAKAYAMA_FRA2_TARGETS, NAKAYAMA_FRA2_TARGETS

GO_INTRAMOLECULAR_TRANSFERASE_ACTIVITY, GO_INTRAMOLECULAR_TRANSFERASE_ACTIVITY

GO_CIRCADIAN_RHYTHM, GO_CIRCADIAN_RHYTHM

GO_VESICLE_DOCKING, GO_VESICLE_DOCKING

GCM_DDX11, GCM_DDX11

PID_P13KCI_PATHWAY, PID_P13KCI_PATHWAY

MCCLUNG_CREB1_TARGETS_UP, MCCLUNG_CREB1_TARGETS_UP

GSE26488_WT_VS_HDAC7_DELTAP_TG_OT2_THYMOCYTE_WITH_PEPTIDE_INJECTION_UP, GSE26488_WT_VS_HDAC7_DELTAP_TG_OT2_THYMOCYTE_WITH_PEPTIDE_INJECTION_UP

YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_6, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_6

GO_VESICLE_DOCKING_INVOLVED_IN_EXOCYTOSIS, GO_VESICLE_DOCKING_INVOLVED_IN_EXOCYTOSIS

GO_PROTEIN_DEPOLYMERIZATION, GO_PROTEIN_DEPOLYMERIZATION

GSE33292_DN3_THYMOCYTE_VS_TCELL_LYMPHOMA_FROM_TCF1_KO_DN, GSE33292_DN3_THYMOCYTE_VS_TCELL_LYMPHOMA_FROM_TCF1_KO_DN

ROVERSI_GLIOMA_COPY_NUMBER_DN, ROVERSI_GLIOMA_COPY_NUMBER_DN

KEGG_RENAL_CELL_CARCINOMA, KEGG_RENAL_CELL_CARCINOMA

GO_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION, GO_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION

BIOCARTA_HCMV_PATHWAY, BIOCARTA_HCMV_PATHWAY

RAY_TARGETS_OF_P210_BCR_ABL_FUSION_DN, RAY_TARGETS_OF_P210_BCR_ABL_FUSION_DN

GO_INTERACTION_WITH_SYMBIONT, GO_INTERACTION_WITH_SYMBIONT

GO_PHOSPHATIDYLETHANOLAMINE_METABOLIC_PROCESS, GO_PHOSPHATIDYLETHANOLAMINE_METABOLIC_PROCESS

PID_RHOA_PATHWAY, PID_RHOA_PATHWAY

BIOCARTA_EIF2_PATHWAY, BIOCARTA_EIF2_PATHWAY

LEONARD_HYPOXIA, LEONARD_HYPOXIA

GO_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS, GO_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS

GO_RECYCLING_ENDOSOME_MEMBRANE, GO_RECYCLING_ENDOSOME_MEMBRANE

GO_POSITIVE_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS, GO_POSITIVE_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS

REACTOME_RORA_ACTIVATES_CIRCADIAN_EXPRESSION, REACTOME_RORA_ACTIVATES_CIRCADIAN_EXPRESSION

AGTCTAG_MIR151, AGTCTAG_MIR151

GO_TAU_PROTEIN_KINASE_ACTIVITY, GO_TAU_PROTEIN_KINASE_ACTIVITY

REACTOME_G_ALPHA_Z_SIGNALLING_EVENTS, REACTOME_G_ALPHA_Z_SIGNALLING_EVENTS

ZHENG_FOXP3_TARGETS_UP, ZHENG_FOXP3_TARGETS_UP

WANG_RESPONSE_TO_FORSKOLIN_UP, WANG_RESPONSE_TO_FORSKOLIN_UP

GSE15750_WT_VS_TRAF6KO_DAY6_EFF_CD8_TCELL_DN, GSE15750_WT_VS_TRAF6KO_DAY6_EFF_CD8_TCELL_DN

GO_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION, GO_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION

GO_PERINUCLEAR_ENDOPLASMIC_RETICULUM, GO_PERINUCLEAR_ENDOPLASMIC_RETICULUM

chr14q21, chr14q21

GO_MYTUBE_DIFFERENTIATION, GO_MYTUBE_DIFFERENTIATION

BIOCARTA_MET_PATHWAY, BIOCARTA_MET_PATHWAY

KIM_HYPOXIA, KIM_HYPOXIA

WANG_METASTASIS_OF_BREAST_CANCER_ESR1_DN, WANG_METASTASIS_OF_BREAST_CANCER_ESR1_DN

GSE17721_CTRL_VS_LPS_4H_BMDC_DN, GSE17721_CTRL_VS_LPS_4H_BMDC_DN

GO_POSITIVE_REGULATION_OF_AXON_EXTENSION, GO_POSITIVE_REGULATION_OF_AXON_EXTENSION

GERHOLD_RESPONSE_TO_TZD_DN, GERHOLD_RESPONSE_TO_TZD_DN

KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_UV_IN_OLD, KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_UV_IN_OLD

GO_MODIFICATION_OF_MORPHOLOGY_OR_PHYSIOLOGY_OF_OTHER_ORGANISM, GO_MODIFICATION_OF_MORPHOLOGY_OR_PHYSIOLOGY_OF_OTHER_ORGANISM

GO_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GO_EPITHELIAL_TO_MESENCHYMAL_TRANSITION

GSE2585_THYMIC_DC_VS_THYMIC_MACROPHAGE_UP, GSE2585_THYMIC_DC_VS_THYMIC_MACROPHAGE_UP

GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR

TORCHIA_TARGETS_OF_EWSR1_FLI1_FUSION_TOP20_DN, TORCHIA_TARGETS_OF_EWSR1_FLI1_FUSION_TOP20_DN

LANDIS_BREAST_CANCER_PROGRESSION_UP, LANDIS_BREAST_CANCER_PROGRESSION_UP

GO_RNA_POLYADENYLATION, GO_RNA_POLYADENYLATION

ABBUD_LIF_SIGNALING_1_DN, ABBUD_LIF_SIGNALING_1_DN

GO_SYNCYTIIUM_FORMATION, GO_SYNCYTIIUM_FORMATION