

ND_IFNG_TREATED_EPITHELIAL_CELLS_24H_UP, GSE5542_IFNG_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_24H_UP

GSE5542_UNTREATED_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_24H_UP, GSE5542_UNTREATED_VS_IFNA_AND_IFNG_TREATED_E
GSE411_UNSTIM_VS_400MIN_IL6_STIM_SOCS3_KO_MACROPHAGE_DN, GSE411_UNSTIM_VS_400MIN_IL6_STIM_SOCS3_KO_MACROPHAGE_DN
GSE45881_CXCR6HI_VS_CXCR1LO_COLONIC_LAMINA_PROPRIA_DN, GSE45881_CXCR6HI_VS_CXCR1LO_COLONIC_LAMINA_PROPRIA_DN
GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_MEMORY_BCELL_DAY40_UP, GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_MEMORY_BO
GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_DN, GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_DN
GSE5542_IFNA_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP, GSE5542_IFNA_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP
GSE27786_LSK_VS_NEUTROPHIL_DN, GSE27786_LSK_VS_NEUTROPHIL_DN
GSE21670_UNTREATED_VS_TGFB_TREATED_CD4_TCELL_DN, GSE21670_UNTREATED_VS_TGFB_TREATED_CD4_TCELL_DN
ACTGTAG_MIR139, ACTGTAG_MIR139
GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_DN, GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_DN
MIR8057, MIR8057
WP_INTEGRINMEDIATED_CELL_ADHESION, WP_INTEGRINMEDIATED_CELL_ADHESION
WP_THE_HUMAN_IMMUNE_RESPONSE_TO_TUBERCULOSIS, WP_THE_HUMAN_IMMUNE_RESPONSE_TO_TUBERCULOSIS
GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_UP, GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_UP
TAGGTCA_MIR192_MIR215, TAGGTCA_MIR192_MIR215
HP_LIPODYSTROPHY, HP_LIPODYSTROPHY
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREY_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREY_DN
GOBP_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR, GOBP_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR
MIR6888_3P, MIR6888_3P
BOWIE_RESPONSE_TO_EXTRACELLULAR_MATRIX, BOWIE_RESPONSE_TO_EXTRACELLULAR_MATRIX
GOBP_XENOBIOTIC_TRANSPORT, GOBP_XENOBIOTIC_TRANSPORT
GOBP_CELLULAR_GLUCAN_METABOLIC_PROCESS, GOBP_CELLULAR_GLUCAN_METABOLIC_PROCESS
HP_HIGH_MYOPIA, HP_HIGH_MYOPIA
HP_FACIAL_HYPOTONIA, HP_FACIAL_HYPOTONIA
REACTOME_GLYCOGEN_BREAKDOWN_GLYCOGENOLYSIS, REACTOME_GLYCOGEN_BREAKDOWN_GLYCOGENOLYSIS
HP_CARCINOMA, HP_CARCINOMA
PID_SHP2_PATHWAY, PID_SHP2_PATHWAY
MIR6747_5P, MIR6747_5P
HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT, HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT
MIR12114, MIR12114
GU_PDEF_TARGETS_UP, GU_PDEF_TARGETS_UP
MIR3690, MIR3690
MIKKELSEN_ES_LCP_WITH_H3K4ME3, MIKKELSEN_ES_LCP_WITH_H3K4ME3
GOBP_INTERLEUKIN_27_MEDIATED_SIGNALING_PATHWAY, GOBP_INTERLEUKIN_27_MEDIATED_SIGNALING_PATHWAY
REACTOME_INTERLEUKIN_27_SIGNALING, REACTOME_INTERLEUKIN_27_SIGNALING
TSAI_DNAJB4_TARGETS_UP, TSAI_DNAJB4_TARGETS_UP
REACTOME_INTERLEUKIN_35_SIGNALLING, REACTOME_INTERLEUKIN_35_SIGNALLING
GOBP_LEFT_RIGHT_AXIS_SPECIFICATION, GOBP_LEFT_RIGHT_AXIS_SPECIFICATION
REACTOME_ORGANIC_CATION_TRANSPORT, REACTOME_ORGANIC_CATION_TRANSPORT