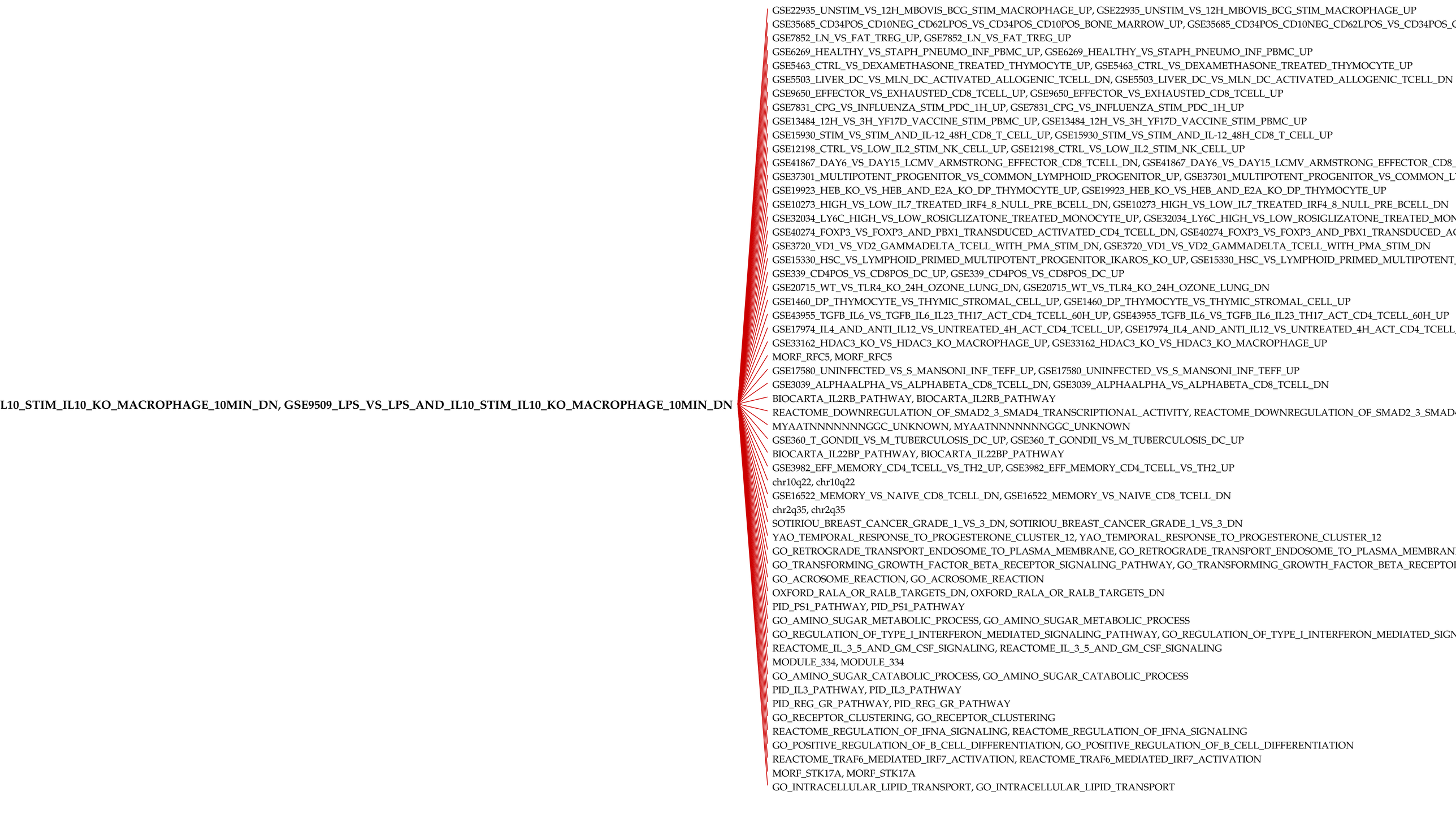
L10_STIM_IL10_KO_MACROPHAGE_10MIN_DN, GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_10MIN_DN
GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MACROPHAGE_UP, GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MACROPHAGE_UP
GSE35685_CD34POS_CD10NEG_CD62LPOS_VS_CD34POS_CD10POS_BONE_MARROW_UP, GSE35685_CD34POS_CD10NEG_CD62LPOS_VS_CD34POS_C
GSE7852_LN_VS_FAT_TREG_UP, GSE7852_LN_VS_FAT_TREG_UP
GSE6269_HEALTHY_VS_STAPH_PNEUMO_INF_PBMC_UP, GSE6269_HEALTHY_VS_STAPH_PNEUMO_INF_PBMC_UP
GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_UP, GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_UP
GSE5503_LIVER_DC_VS_MLN_DC_ACTIVATED_ALLOGENIC_TCELL_DN, GSE5503_LIVER_DC_VS_MLN_DC_ACTIVATED_ALLOGENIC_TCELL_DN
GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_UP, GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_UP
GSE7831_CPG_VS_INFLUENZA_STIM_PDC_1H_UP, GSE7831_CPG_VS_INFLUENZA_STIM_PDC_1H_UP
GSE13484_12H_VS_3H_YF17D_VACCINE_STIM_PBMC_UP, GSE13484_12H_VS_3H_YF17D_VACCINE_STIM_PBMC_UP
GSE15930_STIM_VS_STIM_AND_IL-12_48H_CD8_T_CELL_UP, GSE15930_STIM_VS_STIM_AND_IL-12_48H_CD8_T_CELL_UP
GSE12198_CTRL_VS_LOW_IL2_STIM_NK_CELL_UP, GSE12198_CTRL_VS_LOW_IL2_STIM_NK_CELL_UP
GSE41867_DAY6_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN, GSE41867_DAY6_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_
GSE37301_MULTIPOTENT_PROGENITOR_VS_COMMON_LYMPHOID_PROGENITOR_UP, GSE37301_MULTIPOTENT_PROGENITOR_VS_COMMON_L
GSE19923_HEB_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP, GSE19923_HEB_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP
GSE10273_HIGH_VS_LOW_IL7_TREATED_IRF4_8_NULL_PRE_BCELL_DN, GSE10273_HIGH_VS_LOW_IL7_TREATED_IRF4_8_NULL_PRE_BCELL_DN
GSE32034_LY6C_HIGH_VS_LOW_ROSIGLIZATONE_TREATED_MONOCYTE_UP, GSE32034_LY6C_HIGH_VS_LOW_ROSIGLIZATONE_TREATED_MON
GSE40274_FOXP3_VS_FOXP3_AND_PBX1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_PBX1_TRANSDUCED_AC
GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_PMA_STIM_DN, GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_PMA_STIM_DN
GSE15330_HSC_VS_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_IKAROS_KO_UP, GSE15330_HSC_VS_LYMPHOID_PRIMED_MULTIPOTENT_
GSE339_CD4POS_VS_CD8POS_DC_UP, GSE339_CD4POS_VS_CD8POS_DC_UP
GSE20715_WT_VS_TLR4_KO_24H_OZONE_LUNG_DN, GSE20715_WT_VS_TLR4_KO_24H_OZONE_LUNG_DN
GSE1460_DP_THYMOCYTE_VS_THYMIC_STROMAL_CELL_UP, GSE1460_DP_THYMOCYTE_VS_THYMIC_STROMAL_CELL_UP
GSE43955_TGFB_IL6_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP, GSE43955_TGFB_IL6_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP
GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_4H_ACT_CD4_TCELL_UP, GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_4H_ACT_CD4_TCELL_
GSE33162_HDAC3_KO_VS_HDAC3_KO_MACROPHAGE_UP, GSE33162_HDAC3_KO_VS_HDAC3_KO_MACROPHAGE_UP
MORF_RFC5, MORF_RFC5
GSE17580_UNINFECTED_VS_S_MANSONI_INF_TEFF_UP, GSE17580_UNINFECTED_VS_S_MANSONI_INF_TEFF_UP
GSE3039_ALPHAALPHA_VS_ALPHABETA_CD8_TCELL_DN, GSE3039_ALPHAALPHA_VS_ALPHABETA_CD8_TCELL_DN
BIOCARTA_IL2RB_PATHWAY, BIOCARTA_IL2RB_PATHWAY
REACTOME_DOWNREGULATION_OF_SMAD2_3_SMAD4_TRANSCRIPTIONAL_ACTIVITY, REACTOME_DOWNREGULATION_OF_SMAD2_3_SMAD4
MYAATNNNNNNNGGC_UNKNOWN, MYAATNNNNNNNGGC_UNKNOWN
GSE360_T_GONDII_VS_M_TUBERCULOSIS_DC_UP, GSE360_T_GONDII_VS_M_TUBERCULOSIS_DC_UP
BIOCARTA_IL22BP_PATHWAY, BIOCARTA_IL22BP_PATHWAY
GSE3982_EFF_MEMORY_CD4_TCELL_VS_TH2_UP, GSE3982_EFF_MEMORY_CD4_TCELL_VS_TH2_UP
chr10q22, chr10q22
GSE16522_MEMORY_VS_NAIVE_CD8_TCELL_DN, GSE16522_MEMORY_VS_NAIVE_CD8_TCELL_DN
chr2q35, chr2q35
SOTIRIOU_BREAST_CANCER_GRADE_1_VS_3_DN, SOTIRIOU_BREAST_CANCER_GRADE_1_VS_3_DN
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_12, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_12
GO_RETROGRADE_TRANSPORT_ENDOSOME_TO_PLASMA_MEMBRANE, GO_RETROGRADE_TRANSPORT_ENDOSOME_TO_PLASMA_MEMBRAN
GO_TRANSFORMING_GROWTH_FACTOR_BETA_RECEPTOR_SIGNALING_PATHWAY, GO_TRANSFORMING_GROWTH_FACTOR_BETA_RECEPTO
GO_ACROSOME_REACTION, GO_ACROSOME_REACTION
OXFORD_RALA_OR_RALB_TARGETS_DN, OXFORD_RALA_OR_RALB_TARGETS_DN
PID_PS1_PATHWAY, PID_PS1_PATHWAY
GO_AMINO_SUGAR_METABOLIC_PROCESS, GO_AMINO_SUGAR_METABOLIC_PROCESS
GO_REGULATION_OF_TYPE_I_INTERFERON_MEDIATED_SIGNALING_PATHWAY, GO_REGULATION_OF_TYPE_I_INTERFERON_MEDIATED_SIGN
REACTOME_IL_3_5_AND_GM_CSF_SIGNALING, REACTOME_IL_3_5_AND_GM_CSF_SIGNALING
MODULE_334, MODULE_334
GO_AMINO_SUGAR_CATABOLIC_PROCESS, GO_AMINO_SUGAR_CATABOLIC_PROCESS
PID_IL3_PATHWAY, PID_IL3_PATHWAY
PID_REG_GR_PATHWAY, PID_REG_GR_PATHWAY
GO_RECEPTOR_CLUSTERING, GO_RECEPTOR_CLUSTERING
REACTOME_REGULATION_OF_IFNA_SIGNALING, REACTOME_REGULATION_OF_IFNA_SIGNALING
GO_POSITIVE_REGULATION_OF_B_CELL_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_B_CELL_DIFFERENTIATION
REACTOME_TRAF6_MEDIATED_IRF7_ACTIVATION, REACTOME_TRAF6_MEDIATED_IRF7_ACTIVATION
MORF_STK17A, MORF_STK17A
GO_INTRACELLULAR_LIPID_TRANSPORT, GO_INTRACELLULAR_LIPID_TRANSPORT