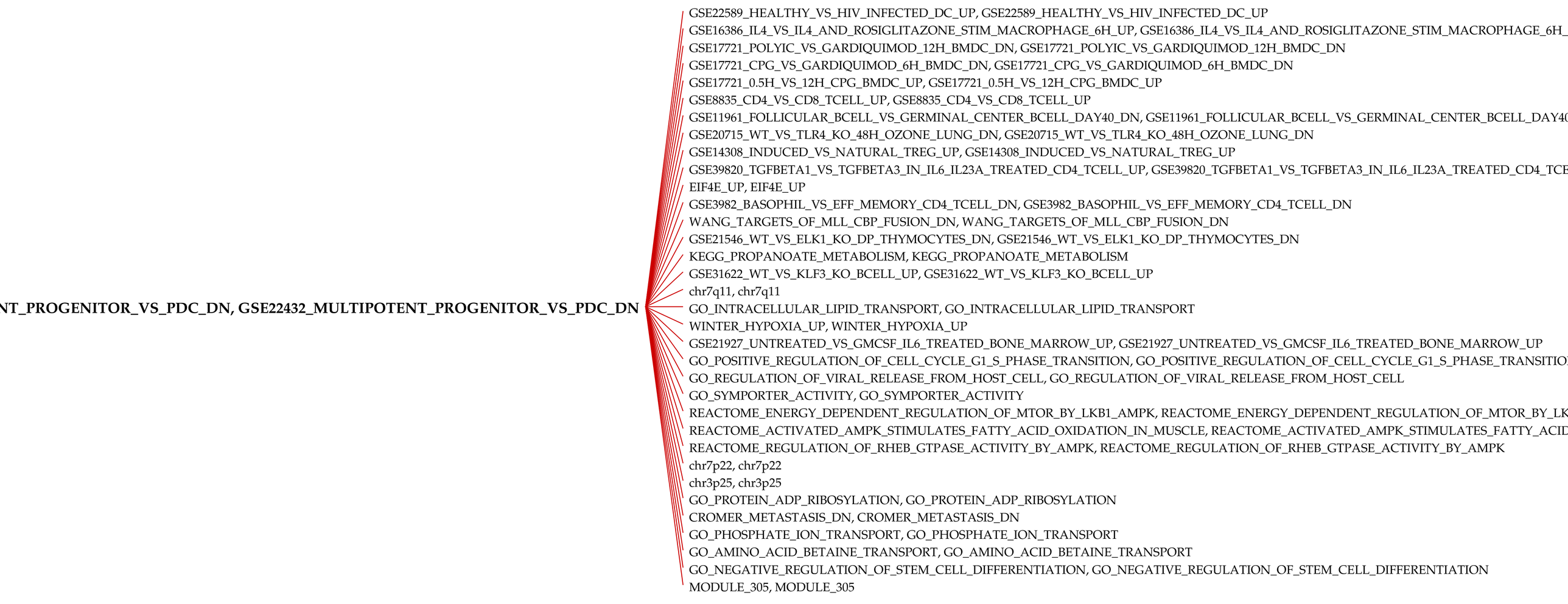
NT_PROGENITOR_VS_PDC_DN, GSE22432_MULTIPOTENT_PROGENITOR_VS_PDC_DN
GSE22589_HEALTHY_VS_HIV_INFECTED_DC_UP, GSE22589_HEALTHY_VS_HIV_INFECTED_DC_UP
GSE16386_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_6H_UP, GSE16386_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_6H_
GSE17721_POLYIC_VS_GARDIQUIMOD_12H_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_12H_BMDC_DN
GSE17721_CPG_VS_GARDIQUIMOD_6H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_6H_BMDC_DN
GSE17721_0.5H_VS_12H_CPG_BMDC_UP, GSE17721_0.5H_VS_12H_CPG_BMDC_UP
GSE8835_CD4_VS_CD8_TCELL_UP, GSE8835_CD4_VS_CD8_TCELL_UP
GSE11961_FOLLICULAR_BCELL_VS_GERMINAL_CENTER_BCELL_DAY40_DN, GSE11961_FOLLICULAR_BCELL_VS_GERMINAL_CENTER_BCELL_DAY40
GSE20715_WT_VS_TLR4_KO_48H_OZONE_LUNG_DN, GSE20715_WT_VS_TLR4_KO_48H_OZONE_LUNG_DN
GSE14308_INDUCED_VS_NATURAL_TREG_UP, GSE14308_INDUCED_VS_NATURAL_TREG_UP
GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCE
EIF4E_UP, EIF4E_UP
GSE3982_BASOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN, GSE3982_BASOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN
WANG_TARGETS_OF_MLL_CBP_FUSION_DN, WANG_TARGETS_OF_MLL_CBP_FUSION_DN
GSE21546_WT_VS_ELK1_KO_DP_THYMOCYTES_DN, GSE21546_WT_VS_ELK1_KO_DP_THYMOCYTES_DN
KEGG_PROPANOATE_METABOLISM, KEGG_PROPANOATE_METABOLISM
GSE31622_WT_VS_KLF3_KO_BCELL_UP, GSE31622_WT_VS_KLF3_KO_BCELL_UP
chr7q11, chr7q11
GO_INTRACELLULAR_LIPID_TRANSPORT, GO_INTRACELLULAR_LIPID_TRANSPORT
WINTER_HYPOXIA_UP, WINTER_HYPOXIA_UP
GSE21927_UNTREATED_VS_GMCSF_IL6_TREATED_BONE_MARROW_UP, GSE21927_UNTREATED_VS_GMCSF_IL6_TREATED_BONE_MARROW_UP
GO_POSITIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION, GO_POSITIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITIO
GO_REGULATION_OF_VIRAL_RELEASE_FROM_HOST_CELL, GO_REGULATION_OF_VIRAL_RELEASE_FROM_HOST_CELL
GO_SYMPORTER_ACTIVITY, GO_SYMPORTER_ACTIVITY
REACTOME_ENERGY_DEPENDENT_REGULATION_OF_MTOR_BY_LKB1_AMPK, REACTOME_ENERGY_DEPENDENT_REGULATION_OF_MTOR_BY_LK
REACTOME_ACTIVATED_AMPK_STIMULATES_FATTY_ACID_OXIDATION_IN_MUSCLE, REACTOME_ACTIVATED_AMPK_STIMULATES_FATTY_ACID
REACTOME_REGULATION_OF_RHEB_GTPASE_ACTIVITY_BY_AMPK, REACTOME_REGULATION_OF_RHEB_GTPASE_ACTIVITY_BY_AMPK
chr7p22, chr7p22
chr3p25, chr3p25
GO_PROTEIN_ADP_RIBOSYLATION, GO_PROTEIN_ADP_RIBOSYLATION
CROMER_METASTASIS_DN, CROMER_METASTASIS_DN
GO_PHOSPHATE_ION_TRANSPORT, GO_PHOSPHATE_ION_TRANSPORT
GO_AMINO_ACID_BETAINE_TRANSPORT, GO_AMINO_ACID_BETAINE_TRANSPORT
GO_NEGATIVE_REGULATION_OF_STEM_CELL_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_STEM_CELL_DIFFERENTIATION
MODULE_305, MODULE_305