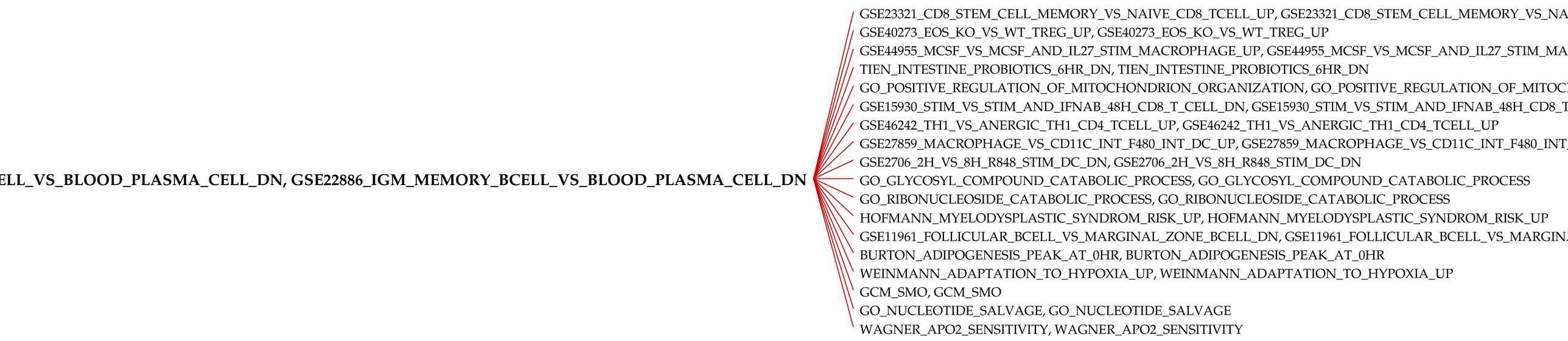
ELL_VS_BLOOD_PLASMA_CELL_DN, GSE22886_IGM_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_DN
GSE23321_CD8_STEM_CELL_MEMORY_VS_NAIVE_CD8_TCELL_UP, GSE23321_CD8_STEM_CELL_MEMORY_VS_NA
GSE40273_EOS_KO_VS_WT_TREG_UP, GSE40273_EOS_KO_VS_WT_TREG_UP
GSE44955_MCSF_VS_MCSF_AND_IL27_STIM_MACROPHAGE_UP, GSE44955_MCSF_VS_MCSF_AND_IL27_STIM_MA
TIEN_INTESTINE_PROBIOTICS_6HR_DN, TIEN_INTESTINE_PROBIOTICS_6HR_DN
GO_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GO_POSITIVE_REGULATION_OF_MITOC
GSE15930_STIM_VS_STIM_AND_IFNAB_48H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_IFNAB_48H_CD8_T
GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_UP, GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_UP
GSE27859_MACROPHAGE_VS_CD11C_INT_F480_INT_DC_UP, GSE27859_MACROPHAGE_VS_CD11C_INT_F480_INT
GSE2706_2H_VS_8H_R848_STIM_DC_DN, GSE2706_2H_VS_8H_R848_STIM_DC_DN
GO_GLYCOSYL_COMPOUND_CATABOLIC_PROCESS, GO_GLYCOSYL_COMPOUND_CATABOLIC_PROCESS
GO_RIBONUCLEOSIDE_CATABOLIC_PROCESS, GO_RIBONUCLEOSIDE_CATABOLIC_PROCESS
HOFMANN_MYELODYSPLASTIC_SYNDROM_RISK_UP, HOFMANN_MYELODYSPLASTIC_SYNDROM_RISK_UP
GSE11961_FOLLICULAR_BCELL_VS_MARGINAL_ZONE_BCELL_DN, GSE11961_FOLLICULAR_BCELL_VS_MARGIN
BURTON_ADIPOGENESIS_PEAK_AT_0HR, BURTON_ADIPOGENESIS_PEAK_AT_0HR
WEINMANN_ADAPTATION_TO_HYPOXIA_UP, WEINMANN_ADAPTATION_TO_HYPOXIA_UP
GCM_SMO, GCM_SMO
GO_NUCLEOTIDE_SALVAGE, GO_NUCLEOTIDE_SALVAGE
WAGNER_APO2_SENSITIVITY, WAGNER_APO2_SENSITIVITY