```
ANOBACTERIUM LPSLIKE STIM DC 3H UP, GSE4748 CTRL VS LPS AND CYANOBACTERIUM LPSLIKE STIM DC 3H UP 🤘
                                                                                                    GOBP_POSITIVE_REGULATION_OF_PROTEIN_KINASE_A_SIGNALING, GOBP_POSITIVE_REGULATION_OF_PROT
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GSE5589\_UNSTIM\_VS\_45MIN\_LPS\_STIM\_MACROPHAGE\_UP, GSE5589\_UNSTIM\_VS\_45MIN\_LPS\_STIM\_MACROPH CORRE MULTIPLE MYELOMA UP, CORRE MULTIPLE MYELOMA UP REACTOME\_PEPTIDE\_LIGAND\_BINDING\_RECEPTORS, REACTOME\_PEPTIDE\_LIGAND\_BINDING\_RECEPTORS MEISSNER\_NPC\_HCP\_WITH\_H3K4ME3\_AND\_H3K27ME3, MEISSNER\_NPC\_HCP\_WITH\_H3K4ME3\_AND\_H3K27ME

GOBP REGULATION OF SYNAPSE ASSEMBLY, GOBP REGULATION OF SYNAPSE ASSEMBLY

KRAS.LUNG\_UP.V1\_UP, KRAS.LUNG\_UP.V1\_UP

GOBP KIDNEY MORPHOGENESIS, GOBP KIDNEY MORPHOGENESIS DESCARTES FETAL EYE PDE11A FAM19A2 POSITIVE CELLS, DESCARTES FETAL EYE PDE11A FAM19A2 POSI

KRAS.50 UP.V1 UP, KRAS.50 UP.V1 UP

DESCARTES FETAL LUNG VASCULAR ENDOTHELIAL CELLS, DESCARTES FETAL LUNG VASCULAR ENDOT PIEPOLI\_LGI1\_TARGETS\_DN, PIEPOLI\_LGI1\_TARGETS\_DN

GOBP\_RETINAL\_GANGLION\_CELL\_AXON\_GUIDANCE, GOBP\_RETINAL\_GANGLION\_CELL\_AXON\_GUIDANCE

PID ERBB NETWORK PATHWAY, PID ERBB NETWORK PATHWAY