```
H_MACROPHAGE_VS_CD11C_ING_F480_INT_DC_UP, GSE27859_CD11C_INT_F480_HI_MACROPHAGE_VS_CD11C_ING_F480_INT_DC_UP {
                                                                                                                MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_29, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_29
                                                                                                                GOBP_REGULATION_OF_VASCULOGENESIS, GOBP_REGULATION_OF_VASCULOGENESIS
```

GSE21360\_TERTIARY\_VS\_QUATERNARY\_MEMORY\_CD8\_TCELL\_UP, GSE21360\_TERTIARY\_VS\_QUATERNARY\_MEMORY\_CD8\_TCELL\_UP MIR1913, MIR1913 GSE42088\_UNINF\_VS\_LEISHMANIA\_INF\_DC\_8H\_DN, GSE42088\_UNINF\_VS\_LEISHMANIA\_INF\_DC\_8H\_DN HP\_APLASIA\_HYPOPLASIA\_OF\_THE\_LUNGS, HP\_APLASIA\_HYPOPLASIA\_OF\_THE\_LUNGS GTCNYYATGR\_UNKNOWN, GTCNYYATGR\_UNKNOWN HSF4\_TARGET\_GENES, HSF4\_TARGET\_GENES GOBP\_POSITIVE\_REGULATION\_OF\_PEPTIDYL\_LYSINE\_ACETYLATION, GOBP\_POSITIVE\_REGULATION\_OF\_PEPTIDYL\_LYSINE\_ACETYLATION MODULE 146, MODULE 146 GOBP\_AMINO\_SUGAR\_METABOLIC\_PROCESS, GOBP\_AMINO\_SUGAR\_METABOLIC\_PROCESS GOBP\_RESPONSE\_TO\_TUMOR\_CELL, GOBP\_RESPONSE\_TO\_TUMOR\_CELL GSE19512\_NAUTRAL\_VS\_INDUCED\_TREG\_UP, GSE19512\_NAUTRAL\_VS\_INDUCED\_TREG\_UP GOBP\_CELL\_COMMUNICATION\_BY\_ELECTRICAL\_COUPLING\_INVOLVED\_IN\_CARDIAC\_CONDUCTION, GOBP\_CELL\_COMMUNICATION\_BY\_ELECTRICAL\_COUPLING GOBP\_POTASSIUM\_ION\_HOMEOSTASIS, GOBP\_POTASSIUM\_ION\_HOMEOSTASIS GOBP\_CELL\_COMMUNICATION\_BY\_ELECTRICAL\_COUPLING, GOBP\_CELL\_COMMUNICATION\_BY\_ELECTRICAL\_COUPLING GOBP\_SURFACTANT\_HOMEOSTASIS, GOBP\_SURFACTANT\_HOMEOSTASIS WP\_NUCLEOTIDE\_GPCRS, WP\_NUCLEOTIDE\_GPCRS