

**1C\_INT\_F480\_HI\_MACROPHAGE\_DN, GSE27859\_DC\_VS\_CD11C\_INT\_F480\_HI\_MACROPHAGE\_DN**

GSE29618\_LAIV\_VS\_TIV\_FLU\_VACCINE\_DAY7\_MONOCYTE\_UP, GSE29618\_LAIV\_VS\_TIV\_FLU\_VACCINE\_DAY7\_MONOCYTE\_UP  
GSE26488\_WT\_VS\_HDAC7\_KO\_DOUBLE\_POSITIVE\_THYMOCYTE\_UP, GSE26488\_WT\_VS\_HDAC7\_KO\_DOUBLE\_POSITIVE\_THYMOCYTE\_UP  
PURBEY\_TARGETS\_OF\_CTBP1\_AND\_SATB1\_DN, PURBEY\_TARGETS\_OF\_CTBP1\_AND\_SATB1\_DN  
GSE5589\_LPS\_VS\_LPS\_AND\_IL10\_STIM\_MACROPHAGE\_45MIN\_DN, GSE5589\_LPS\_VS\_LPS\_AND\_IL10\_STIM\_MACROPHAGE\_45MIN\_DN  
GSE17721\_CPG\_VS\_GARDIQUIMOD\_8H\_BMDC\_UP, GSE17721\_CPG\_VS\_GARDIQUIMOD\_8H\_BMDC\_UP  
GSE13522\_CTRL\_VS\_T\_CRUZI\_G\_STRAIN\_INF\_SKIN\_UP, GSE13522\_CTRL\_VS\_T\_CRUZI\_G\_STRAIN\_INF\_SKIN\_UP  
GO\_REGULATION\_OF\_GLUCOSE\_METABOLIC\_PROCESS, GO\_REGULATION\_OF\_GLUCOSE\_METABOLIC\_PROCESS  
GO\_REGULATION\_OF\_CELLULAR\_SENESCENCE, GO\_REGULATION\_OF\_CELLULAR\_SENESCENCE  
REACTOME\_RAS\_ACTIVATION\_UOPN\_CA2\_INFUX\_THROUGH\_NMDA\_RECEPTOR, REACTOME\_RAS\_ACTIVATION\_UOPN\_CA2\_INFUX\_THROUGH\_NMDA\_  
EPPERT\_CE\_HSC\_LSC, EPPERT\_CE\_HSC\_LSC  
GO\_REGULATION\_OF\_RYANODINE\_SENSITIVE\_CALCIUM\_RELEASE\_CHANNEL\_ACTIVITY, GO\_REGULATION\_OF\_RYANODINE\_SENSITIVE\_CALCIUM\_REL  
GO\_MELANOCYTE\_DIFFERENTIATION, GO\_MELANOCYTE\_DIFFERENTIATION  
GO\_NEUROTRANSMITTER\_METABOLIC\_PROCESS, GO\_NEUROTRANSMITTER\_METABOLIC\_PROCESS