

S\_WT\_UNSTIM\_CD4\_TCELL\_DN, GSE45739\_NRAS\_KO\_VS\_WT\_UNSTIM\_CD4\_TCELL\_DN

GSE45739\_NRAS\_KO\_VS\_WT\_ACD3\_ACD28\_STIM\_CD4\_TCELL\_DN, GSE45739\_NRAS\_KO\_VS\_WT\_ACD3\_ACD28\_STIM\_CD4\_TCELL\_DN  
GSE25123\_WT\_VS\_PPARG\_KO\_MACROPHAGE\_IL4\_AND\_ROSIGLITAZONE\_STIM\_DN, GSE25123\_WT\_VS\_PPARG\_KO\_MACROPHAGE\_IL4\_AND\_ROSIGLITAZONE\_STIM\_DN  
LAKE\_ADULT\_KIDNEY\_C27\_VASCULAR\_SMOOTH\_MUSCLE\_CELLS\_AND\_PERICYTES, LAKE\_ADULT\_KIDNEY\_C27\_VASCULAR\_SMOOTH\_MUSCLE\_CELLS\_AND\_PERICYTES  
MIR199A\_3P\_MIR199B\_3P, MIR199A\_3P\_MIR199B\_3P  
MIR3129\_5P, MIR3129\_5P  
MIR450A\_1\_3P, MIR450A\_1\_3P  
HP\_CIRRHOSIS, HP\_CIRRHOSIS  
GSE21927\_UNTREATED\_VS\_GMCSF\_IL6\_TREATED\_BONE\_MARROW\_UP, GSE21927\_UNTREATED\_VS\_GMCSF\_IL6\_TREATED\_BONE\_MARROW\_UP  
HP\_EXODEVIATION, HP\_EXODEVIATION  
TTNNANAGCYR\_UNKNOWN, TTTNNANAGCYR\_UNKNOWN  
GSE16385\_IFNG\_TNF\_VS\_IL4\_STIM\_MACROPHAGE\_ROSIGLITAZONE\_TREATED\_UP, GSE16385\_IFNG\_TNF\_VS\_IL4\_STIM\_MACROPHAGE\_ROSIGLITAZONE\_TREATED\_UP  
MED25\_TARGET\_GENES, MED25\_TARGET\_GENES  
MIR6748\_3P, MIR6748\_3P  
HP\_CRANIAL\_HYPEROSTOSIS, HP\_CRANIAL\_HYPEROSTOSIS  
MIR549A\_5P, MIR549A\_5P  
GOMF\_METALLOPEPTIDASE\_ACTIVITY, GOMF\_METALLOPEPTIDASE\_ACTIVITY  
LAKE\_ADULT\_KIDNEY\_C1\_EPITHELIAL\_CELLS\_UNASSIGNED, LAKE\_ADULT\_KIDNEY\_C1\_EPITHELIAL\_CELLS\_UNASSIGNED  
GOBP\_HEAT\_GENERATION, GOBP\_HEAT\_GENERATION  
DESCARTES\_FETAL\_HEART\_VISCERAL\_NEURONS, DESCARTES\_FETAL\_HEART\_VISCERAL\_NEURONS  
GOBP\_REGULATION\_OF\_NON\_CANONICAL\_WNT\_SIGNALING\_PATHWAY, GOBP\_REGULATION\_OF\_NON\_CANONICAL\_WNT\_SIGNALING\_PATHWAY  
GOBP\_POSITIVE\_REGULATION\_OF\_VOLTAGE\_GATED\_CALCIUM\_CHANNEL\_ACTIVITY, GOBP\_POSITIVE\_REGULATION\_OF\_VOLTAGE\_GATED\_CALCIUM\_CHANNEL\_ACTIVITY  
GOBP\_REGULATION\_OF\_HEAT\_GENERATION, GOBP\_REGULATION\_OF\_HEAT\_GENERATION