

BCL6\_KO\_VS\_WT\_TREG\_UP, GSE40493\_BCL6\_KO\_VS\_WT\_TREG\_UP

GSE2770\_IL12\_VS\_IL4\_TREATED\_ACT\_CD4\_TCELL\_2H\_UP, GSE2770\_IL12\_VS\_IL4\_TREATED\_ACT\_CD4\_TCELL\_2H\_UP  
HP\_NASOGASTRIC\_TUBE\_FEEDING, HP\_NASOGASTRIC\_TUBE\_FEEDING  
MIR449B\_3P, MIR449B\_3P  
HP\_ASTHMA, HP\_ASTHMA  
HP\_LARGE\_BASAL\_GANGLIA, HP\_LARGE\_BASAL\_GANGLIA  
GOBP\_POSITIVE\_REGULATION\_OF\_CELL\_GROWTH, GOBP\_POSITIVE\_REGULATION\_OF\_CELL\_GROWTH  
REACTOME\_RHOV\_GTPASE\_CYCLE, REACTOME\_RHOV\_GTPASE\_CYCLE  
MIR5571\_3P, MIR5571\_3P  
MIR582\_3P, MIR582\_3P  
GOBP\_NEGATIVE\_REGULATION\_OF\_DEPHOSPHORYLATION, GOBP\_NEGATIVE\_REGULATION\_OF\_DEPHOSPHORYLATION  
GOBP\_PROTEIN\_QUALITY\_CONTROL\_FOR\_MISFOLDED\_OR\_INCOMPLETELY\_SYNTHESIZED\_PROTEINS, GOBP\_PROTEIN\_QUALITY\_CONTROL\_FOR\_MISFOLDED\_OR\_INCOMPLETELY\_SYNTHESIZED\_PROTEINS  
MIR12127, MIR12127  
REACTOME\_ACTIVATION\_OF\_RAC1, REACTOME\_ACTIVATION\_OF\_RAC1  
GOCC\_CUL4A\_RING\_E3\_UBIQUITIN\_LIGASE\_COMPLEX, GOCC\_CUL4A\_RING\_E3\_UBIQUITIN\_LIGASE\_COMPLEX  
HP\_ABNORMAL\_ANTERIOR\_HORN\_CELL\_MORPHOLOGY, HP\_ABNORMAL\_ANTERIOR\_HORN\_CELL\_MORPHOLOGY  
GOMF\_METALLOENDOPEPTIDASE\_ACTIVITY, GOMF\_METALLOENDOPEPTIDASE\_ACTIVITY  
chr6q16, chr6q16  
GOMF\_NEUROTRANSMITTER\_RECEPTOR\_ACTIVITY, GOMF\_NEUROTRANSMITTER\_RECEPTOR\_ACTIVITY  
GOBP\_REGULATION\_OF\_CGMP\_MEDIATED\_SIGNALING, GOBP\_REGULATION\_OF\_CGMP\_MEDIATED\_SIGNALING  
YIH\_RESPONSE\_TO\_ARSENITE\_C4, YIH\_RESPONSE\_TO\_ARSENITE\_C4  
GOBP\_SECONDARY\_METABOLITE\_BIOSYNTHETIC\_PROCESS, GOBP\_SECONDARY\_METABOLITE\_BIOSYNTHETIC\_PROCESS  
WP\_HEDGEHOG\_SIGNALING\_PATHWAY\_NETPATH, WP\_HEDGEHOG\_SIGNALING\_PATHWAY\_NETPATH  
GOBP\_SMOOTHENED\_SIGNALING\_PATHWAY\_INVOLVED\_IN\_VENTRAL\_SPINAL\_CORD\_PATTERNING, GOBP\_SMOOTHENED\_SIGNALING\_PATHWAY\_INVOLVED\_IN\_VENTRAL\_SPINAL\_CORD\_PATTERNING  
GOBP\_CELL\_FATE\_COMMITMENT\_INVOLVED\_IN\_PATTERN\_SPECIFICATION, GOBP\_CELL\_FATE\_COMMITMENT\_INVOLVED\_IN\_PATTERN\_SPECIFICATION