

VS_IKAROS_KO_HSC_UP, GSE15330_WT_VS_IKAROS_KO_HSC_UP

- GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN, GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN
- GSE12845_IGD_POS_BLOOD_VS_PRE_GC_TONSIL_BCELL_UP, GSE12845_IGD_POS_BLOOD_VS_PRE_GC_TONSIL_BCELL_UP
- GSE7460_WT_VS_FOXP3_HET_ACT_WITH_TGFB_TCONV_UP, GSE7460_WT_VS_FOXP3_HET_ACT_WITH_TGFB_TCONV_UP
- GO_PDZ_DOMAIN_BINDING, GO_PDZ_DOMAIN_BINDING
- GO_PROTEIN_DEGLYCOSYLATION, GO_PROTEIN_DEGLYCOSYLATION
- GO_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_MIGRATION, GO_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_MIGRATION
- MCCABE_HOXC6_TARGETS_DN, MCCABE_HOXC6_TARGETS_DN
- GO_LATERAL_PLASMA_MEMBRANE, GO_LATERAL_PLASMA_MEMBRANE
- GO_POSITIVE_REGULATION_OF_HORMONE_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_HORMONE_METABOLIC_PROCESS