

ANGIOGENIC_MARKERS_UP, VART_KSHV_INFECTION_ANGIOGENIC_MARKERS_UP

GO_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_PROLIFERATION, GO_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_PROLIFERATION
GSE9988_LPS_VS_LOW_LPS_MONOCYTE_DN, GSE9988_LPS_VS_LOW_LPS_MONOCYTE_DN
GSE22443_NAIVE_VS_ACT_AND_IL2_TREATED_CD8_TCELL_DN, GSE22443_NAIVE_VS_ACT_AND_IL2_TREATED_CD8_TCELL_DN
MIKKELSEN_ES_LCP_WITH_H3K4ME3, MIKKELSEN_ES_LCP_WITH_H3K4ME3
GSE22611_NOD2_VS_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_DN, GSE22611_NOD2_VS_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_DN
GO_PEPTIDASE_INHIBITOR_ACTIVITY, GO_PEPTIDASE_INHIBITOR_ACTIVITY
GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_TYROSINE_KINASE_ACTIVITY, GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_TYROSINE_KINASE_ACTIVITY
GO_MESONEPHRIC_TUBULE_MORPHOGENESIS, GO_MESONEPHRIC_TUBULE_MORPHOGENESIS
KRAS.BREAST_UP.V1_DN, KRAS.BREAST_UP.V1_DN
GSE11961_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_DAY7_DN, GSE11961_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_DAY7_DN
MORF_RAB3A, MORF_RAB3A
GSE19825_NAIVE_VS_DAY3_EFF_CD8_TCELL_UP, GSE19825_NAIVE_VS_DAY3_EFF_CD8_TCELL_UP
GSE38697_LIGHT_ZONE_VS_DARK_ZONE_BCELL_DN, GSE38697_LIGHT_ZONE_VS_DARK_ZONE_BCELL_DN
GSE23502_BM_VS_COLON_TUMOR_HDC_KO_MYELOID_DERIVED_SUPPRESSOR_CELL_UP, GSE23502_BM_VS_COLON_TUMOR_HDC_KO_MYELOID_DERIVED_SUPPRESSOR_CELL_UP
GO_ACTIVATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_TYROSINE_KINASE_ACTIVITY, GO_ACTIVATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_TYROSINE_KINASE_ACTIVITY
GO_REGULATION_OF_HAIR_FOLLICLE_DEVELOPMENT, GO_REGULATION_OF_HAIR_FOLLICLE_DEVELOPMENT
HAEGERSTRAND_RESPONSE_TO_IMATINIB, HAEGERSTRAND_RESPONSE_TO_IMATINIB
GO_POSITIVE_REGULATION_OF_OSTEOBLAST_PROLIFERATION, GO_POSITIVE_REGULATION_OF_OSTEOBLAST_PROLIFERATION
GSE13522_WT_VS_IFNG_KO_SKIN_UP, GSE13522_WT_VS_IFNG_KO_SKIN_UP
GO_ORGAN_INDUCTION, GO_ORGAN_INDUCTION
GSE37336_LY6C_POS_VS_NEG_NAIVE_CD4_TCELL_DN, GSE37336_LY6C_POS_VS_NEG_NAIVE_CD4_TCELL_DN