

DIATE_FILAMENT, GO_INTERMEDIATE_FILAMENT

GO_KERATIN_FILAMENT, GO_KERATIN_FILAMENT
GO_KERATINIZATION, GO_KERATINIZATION
MODULE_298, MODULE_298
HUPER_BREAST_BASAL_VS_LUMINAL_DN, HUPER_BREAST_BASAL_VS_LUMINAL_DN
MUELLER_METHYLATED_IN_GLIOBLASTOMA, MUELLER_METHYLATED_IN_GLIOBLASTOMA
BOSCO_EPITHELIAL_DIFFERENTIATION_MODULE, BOSCO_EPITHELIAL_DIFFERENTIATION_MODULE
GO_CELL_CELL_ADHERENS_JUNCTION, GO_CELL_CELL_ADHERENS_JUNCTION
MODULE_68, MODULE_68
TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_DUCTAL_NORMAL_DN, TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_DUCTAL_NORMAL_DN
chr21q22, chr21q22
GSE10325_MYELOID_VS_LUPUS_MYELOID_UP, GSE10325_MYELOID_VS_LUPUS_MYELOID_UP
HALLMARK_KRAS_SIGNALING_DN, HALLMARK_KRAS_SIGNALING_DN
GO_CELL_JUNCTION_ORGANIZATION, GO_CELL_JUNCTION_ORGANIZATION
HUPER_BREAST_BASAL_VS_LUMINAL_UP, HUPER_BREAST_BASAL_VS_LUMINAL_UP
SNF5_DN.V1_DN, SNF5_DN.V1_DN
SENESE_HDAC2_TARGETS_DN, SENESE_HDAC2_TARGETS_DN
GO_MICROTUBULE_END, GO_MICROTUBULE_END
KRAS.300_UP.V1_DN, KRAS.300_UP.V1_DN
PID_CASPASE_PATHWAY, PID_CASPASE_PATHWAY
GSE29949_MICROGLIA_BRAIN_VS_CD8_NEG_DC_SPLEEN_DN, GSE29949_MICROGLIA_BRAIN_VS_CD8_NEG_DC_SPLEEN_DN
JECHLINGER_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_DN, JECHLINGER_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_DN
GSE41867_DAY6_VS_DAY15_LCMV_CLONE13_EFFECTOR_CD8_TCELL_UP, GSE41867_DAY6_VS_DAY15_LCMV_CLONE13_EFFECTOR_CD8_TCELL_UP
GO_ACTIN_FILAMENT_BASED_MOVEMENT, GO_ACTIN_FILAMENT_BASED_MOVEMENT
GSE27670_CTRL_VS_BLIMP1_TRANSDUCED_GC_BCELL_UP, GSE27670_CTRL_VS_BLIMP1_TRANSDUCED_GC_BCELL_UP
GO_CELL_CELL_JUNCTION_ASSEMBLY, GO_CELL_CELL_JUNCTION_ASSEMBLY
GO_MICROTUBULE_PLUS_END, GO_MICROTUBULE_PLUS_END
BIOCARTA_FAS_PATHWAY, BIOCARTA_FAS_PATHWAY
GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMOCYTE_UP, GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMOCYTE_UP
LIN_SILENCED_BY_TUMOR_MICROENVIRONMENT, LIN_SILENCED_BY_TUMOR_MICROENVIRONMENT
GO_CELL_JUNCTION_ASSEMBLY, GO_CELL_JUNCTION_ASSEMBLY
GO_KERATINOCYTE_DIFFERENTIATION, GO_KERATINOCYTE_DIFFERENTIATION
BIOCARTA_TNFR1_PATHWAY, BIOCARTA_TNFR1_PATHWAY
GNF2_SPRR1B, GNF2_SPRR1B
KEGG_ARRHYTHMOGENIC_RIGHT_VENTRICULAR_CARDIOMYOPATHY_ARVC, KEGG_ARRHYTHMOGENIC_RIGHT_VENTRICULAR_CARDIOMYOPATHY_ARVC
GSE27092_WT_VS_HDAC7_PHOSPHO_DEFICIENT_CD8_TCELL_UP, GSE27092_WT_VS_HDAC7_PHOSPHO_DEFICIENT_CD8_TCELL_UP
GSE41867_DAY8_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_DAY8_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP
GO_HETEROTYPIC_CELL_CELL_ADHESION, GO_HETEROTYPIC_CELL_CELL_ADHESION
GSE37532_WT_VS_PPARG_KO_LN_TREG_DN, GSE37532_WT_VS_PPARG_KO_LN_TREG_DN
GO_ACTIN_MEDIATED_CELL_CONTRACTION, GO_ACTIN_MEDIATED_CELL_CONTRACTION
GSE40666_STAT1_KO_VS_STAT4_KO_CD8_TCELL_DN, GSE40666_STAT1_KO_VS_STAT4_KO_CD8_TCELL_DN
KRAS.LUNG_UP.V1_DN, KRAS.LUNG_UP.V1_DN
GO_REGULATION_OF_ACTION_POTENTIAL, GO_REGULATION_OF_ACTION_POTENTIAL
GO_MULTICELLULAR_ORGANISMAL_SIGNALING, GO_MULTICELLULAR_ORGANISMAL_SIGNALING
GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN, GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN
BROWNE_HCMV_INFECTION_16HR_DN, BROWNE_HCMV_INFECTION_16HR_DN
ONDER_CDH1_TARGETS_3_DN, ONDER_CDH1_TARGETS_3_DN
MISHRA_CARCINOMA_ASSOCIATED_FIBROBLAST_DN, MISHRA_CARCINOMA_ASSOCIATED_FIBROBLAST_DN
LIANG_SILENCED_BY_METHYLATION_UP, LIANG_SILENCED_BY_METHYLATION_UP
MODULE_297, MODULE_297
MODULE_357, MODULE_357
GSE40666_UNTREATED_VS_IFNA_STIM_CD8_TCELL_90MIN_UP, GSE40666_UNTREATED_VS_IFNA_STIM_CD8_TCELL_90MIN_UP
GO_PROTEIN_BINDING_INVOLVED_IN_CELL_ADHESION, GO_PROTEIN_BINDING_INVOLVED_IN_CELL_ADHESION
JI_CARCINOGENESIS_BY_KRAS_AND_STK11_UP, JI_CARCINOGENESIS_BY_KRAS_AND_STK11_UP
KEGG_AMYOTROPHIC_LATERAL_SCLEROSIS_ALS, KEGG_AMYOTROPHIC_LATERAL_SCLEROSIS_ALS
GCM_ATM, GCM_ATM
MODULE_154, MODULE_154
GSE37605_TREG_VS_TCONV_C57BL6_FOXP3_IRES_GFP_UP, GSE37605_TREG_VS_TCONV_C57BL6_FOXP3_IRES_GFP_UP
TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_LOBULAR_NORMAL_DN, TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_LOBULAR_NORMAL_DN
chr17q12, chr17q12
GU_PDEF_TARGETS_DN, GU_PDEF_TARGETS_DN
GO_I_BAND, GO_I_BAND
MEISSNER_NPC_ICP_WITH_H3_UNMETHYLATED, MEISSNER_NPC_ICP_WITH_H3_UNMETHYLATED
KRAS.50_UP.V1_DN, KRAS.50_UP.V1_DN
GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_DN, GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_DN
HINATA_NFKB_TARGETS_KERATINOCYTE_UP, HINATA_NFKB_TARGETS_KERATINOCYTE_UP
GSE27786_LIN_NEG_VS_ERYTHROBLAST_DN, GSE27786_LIN_NEG_VS_ERYTHROBLAST_DN
GO_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEMBLY, GO_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEMBLY
GO_INTERMEDIATE_FILAMENT_BASED_PROCESS, GO_INTERMEDIATE_FILAMENT_BASED_PROCESS
MCCLUNG_DELTA_FOSB_TARGETS_2WK, MCCLUNG_DELTA_FOSB_TARGETS_2WK
CERIBELLI_PROMOTERS_INACTIVE_AND_BOUND_BY_NFY, CERIBELLI_PROMOTERS_INACTIVE_AND_BOUND_BY_NFY
GSE17721_CTRL_VS_CPG_1H_BMDC_DN, GSE17721_CTRL_VS_CPG_1H_BMDC_DN
TANAKA_METHYLATED_IN_ESOPHAGEAL_CARCINOMA, TANAKA_METHYLATED_IN_ESOPHAGEAL_CARCINOMA
GO_RESPONSE_TO_SALT, GO_RESPONSE_TO_SALT
POU3F2_01, POU3F2_01
AP2REP_01, AP2REP_01
GO_REGULATION_OF_VENTRICULAR_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL, GO_REGULATION_OF_VENTRICULAR_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL
REACTOME_APOPTOTIC_EXECUTION_PHASE, REACTOME_APOPTOTIC_EXECUTION_PHASE
KORKOLA_EMBRYONIC_CARCINOMA_VS_SEMINOMA_UP, KORKOLA_EMBRYONIC_CARCINOMA_VS_SEMINOMA_UP
GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_UP, GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_UP
POOLA_INVASIVE_BREAST_CANCER_DN, POOLA_INVASIVE_BREAST_CANCER_DN
GO_RECEPTOR_SIGNALING_COMPLEX_SCAFFOLD_ACTIVITY, GO_RECEPTOR_SIGNALING_COMPLEX_SCAFFOLD_ACTIVITY
GO_MULTICELLULAR_ORGANISMAL_WATER_HOMEOSTASIS, GO_MULTICELLULAR_ORGANISMAL_WATER_HOMEOSTASIS
DOANE_BREAST_CANCER_ESR1_DN, DOANE_BREAST_CANCER_ESR1_DN
SCHUETZ_BREAST_CANCER_DUCTAL_INVASIVE_DN, SCHUETZ_BREAST_CANCER_DUCTAL_INVASIVE_DN
CROMER_TUMORIGENESIS_DN, CROMER_TUMORIGENESIS_DN
TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_DN, TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_DN
BIOCARTA_CASPASE_PATHWAY, BIOCARTA_CASPASE_PATHWAY