
GO_KERATIN_FILAMENT, GO_KERATIN_FILAMENT
KIM_RESPONSE_TO_TSA_AND_DECITABINE_UP, KIM_RESPONSE_TO_TSA_AND_DECITABINE_UP
GO_KERATINIZATION, GO_KERATINIZATION
LEF1_UP.V1_UP, LEF1_UP.V1_UP
BOSCO_EPITHELIAL_DIFFERENTIATION_MODULE, BOSCO_EPITHELIAL_DIFFERENTIATION_MODULE
MODULE_298, MODULE_298
MODULE_68, MODULE_68
HUPER_BREAST_BASAL_VS_LUMINAL_DN, HUPER_BREAST_BASAL_VS_LUMINAL_DN
GSE40666_STAT1_KO_VS_STAT4_KO_CD8_TCELL_DN, GSE40666_STAT1_KO_VS_STAT4_KO_CD8_TCELL_DN
RB_P130_DN.V1_DN, RB_P130_DN.V1_DN
GO_ACTIN_FILAMENT_BASED_MOVEMENT, GO_ACTIN_FILAMENT_BASED_MOVEMENT
CHIBA_RESPONSE_TO_TSA_UP, CHIBA_RESPONSE_TO_TSA_UP
GO_CELL_JUNCTION_ORGANIZATION, GO_CELL_JUNCTION_ORGANIZATION
GO_ACTIN_MEDIATED_CELL_CONTRACTION, GO_ACTIN_MEDIATED_CELL_CONTRACTION
LIANG_SILENCED_BY_METHYLATION_UP, LIANG_SILENCED_BY_METHYLATION_UP
GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMOCYTE_UP, GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMO
KEGG_ARRHYTHMOGENIC_RIGHT_VENTRICULAR_CARDIOMYOPATHY_ARVC, KEGG_ARRHYTHMOGENIC_RIGHT_VENTRICULAR_CARDIOMYOPATHY_ARVC
MODULE_111, MODULE_111
GSE41867_DAY6_VS_DAY15_LCMV_CLONE13_EFFECTOR_CD8_TCELL_UP, GSE41867_DAY6_VS_DAY15_LCMV_CLONE13_EFFECTOR_CD8_TCELL_UP
BIOCARTA_CASPASE_PATHWAY, BIOCARTA_CASPASE_PATHWAY
CROMER_TUMORIGENESIS_DN, CROMER_TUMORIGENESIS_DN
TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_LOBULAR_NORMAL_UP, TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_LOBULAR_NORMAL_UP
GSE10325_MYELOID_VS_LUPUS_MYELOID_UP, GSE10325_MYELOID_VS_LUPUS_MYELOID_UP
GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_8H_DN, GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_8H_DN
WANG_BARRETTS_ESOPHAGUS_AND_ESOPHAGUS_CANCER_DN, WANG_BARRETTS_ESOPHAGUS_AND_ESOPHAGUS_CANCER_DN
RICKMAN_HEAD_AND_NECK_CANCER_C, RICKMAN_HEAD_AND_NECK_CANCER_C
chr17q12, chr17q12
MODULE_157, MODULE_157
GSE27670_CTRL_VS_BLIMP1_TRANSDUCED_GC_BCELL_UP, GSE27670_CTRL_VS_BLIMP1_TRANSDUCED_GC_BCELL_UP
TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_DN, TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_DN
GO_REGULATION_OF_ACTIN_FILAMENT_BASED_MOVEMENT, GO_REGULATION_OF_ACTIN_FILAMENT_BASED_MOVEMENT
GO_SCAFFOLD_PROTEIN_BINDING, GO_SCAFFOLD_PROTEIN_BINDING
GSE29949_MICROGLIA_BRAIN_VS_CD8_NEG_DC_SPLEEN_DN, GSE29949_MICROGLIA_BRAIN_VS_CD8_NEG_DC_SPLEEN_DN
TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_DUCTAL_NORMAL_DN, TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_DUCTAL_NORMAL_DN
GSE17721_CTRL_VS_CPG_1H_BMDC_DN, GSE17721_CTRL_VS_CPG_1H_BMDC_DN
LIN_SILENCED_BY_TUMOR_MICROENVIRONMENT, LIN_SILENCED_BY_TUMOR_MICROENVIRONMENT
GO_I_BAND, GO_I_BAND
REACTOME_APOPTOTIC_CLEAVAGE_OF_CELLULAR_PROTEINS, REACTOME_APOPTOTIC_CLEAVAGE_OF_CELLULAR_PROTEINS
GNF2_SPRR1B, GNF2_SPRR1B
AP2REP_01, AP2REP_01
chr21q22, chr21q22
HENDRICKS_SMARCA4_TARGETS_UP, HENDRICKS_SMARCA4_TARGETS_UP
HINATA_NFKB_TARGETS_KERATINOCYTE_UP, HINATA_NFKB_TARGETS_KERATINOCYTE_UP
GSE37532_WT_VS_PPARG_KO_LN_TREG_DN, GSE37532_WT_VS_PPARG_KO_LN_TREG_DN
POOLA_INVASIVE_BREAST_CANCER_DN, POOLA_INVASIVE_BREAST_CANCER_DN
GO_KERATINOCYTE_DIFFERENTIATION, GO_KERATINOCYTE_DIFFERENTIATION
GO_STRUCTURAL_CONSTITUENT_OF_EYE_LENS, GO_STRUCTURAL_CONSTITUENT_OF_EYE_LENS
GO_CELL_JUNCTION_ASSEMBLY, GO_CELL_JUNCTION_ASSEMBLY
GNF2_CDH3, GNF2_CDH3
KRAS.50_UP.V1_DN, KRAS.50_UP.V1_DN
KRAS.300_UP.V1_DN, KRAS.300_UP.V1_DN
GCM_PTPRU, GCM_PTPRU
GO_STRUCTURAL_CONSTITUENT_OF_CYTOSKELETON, GO_STRUCTURAL_CONSTITUENT_OF_CYTOSKELETON
HUPER_BREAST_BASAL_VS_LUMINAL_UP, HUPER_BREAST_BASAL_VS_LUMINAL_UP
GO_GLYCOPROTEIN_COMPLEX, GO_GLYCOPROTEIN_COMPLEX
REACTOME_APOPTOTIC_EXECUTION_PHASE, REACTOME_APOPTOTIC_EXECUTION_PHASE
KORKOLA_EMBRYONIC_CARCINOMA_VS_SEMINOMA_UP, KORKOLA_EMBRYONIC_CARCINOMA_VS_SEMINOMA_UP
GCM_MYCL1, GCM_MYCL1
GO_RECEPTOR_SIGNALING_COMPLEX_SCAFFOLD_ACTIVITY, GO_RECEPTOR_SIGNALING_COMPLEX_SCAFFOLD_ACTIVITY
GNF2_SERPINB5, GNF2_SERPINB5
GSE23502_WT_VS_HDC_KO_MYELOID_DERIVED_SUPPRESSOR_CELL_COLON_TUMOR_DN, GSE23502_WT_VS_HDC_KO_MYELOID_DERIVED_SUPPRESSOR_CELL_C
HUMMERICH_MALIGNANT_SKIN_TUMOR_UP, HUMMERICH_MALIGNANT_SKIN_TUMOR_UP
GO_MICROTUBULE_PLUS_END, GO_MICROTUBULE_PLUS_END
CAR_WBSCR22, CAR_WBSCR22
GO_INTERCALATED_DISC, GO_INTERCALATED_DISC
GO_CELL_SUBSTRATE_JUNCTION_ASSEMBLY, GO_CELL_SUBSTRATE_JUNCTION_ASSEMBLY
YAO_HOXA10_TARGETS_VIA_PROGESTERONE_DN, YAO_HOXA10_TARGETS_VIA_PROGESTERONE_DN
GO_CELL_COMMUNICATION_INVOLVED_IN_CARDIAC_CONDUCTION, GO_CELL_COMMUNICATION_INVOLVED_IN_CARDIAC_CONDUCTION
GO_CELL_CELL_CONTACT_ZONE, GO_CELL_CELL_CONTACT_ZONE
MODULE_297, MODULE_297
MODULE_154, MODULE_154
GO_MICROTUBULE_END, GO_MICROTUBULE_END
GO_REGULATION_OF_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL, GO_REGULATION_OF_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL
GO_INTERMEDIATE_FILAMENT_BASED_PROCESS, GO_INTERMEDIATE_FILAMENT_BASED_PROCESS
MODULE_357, MODULE_357
GO_REGULATION_OF_HEART_RATE_BY_CARDIAC_CONDUCTION, GO_REGULATION_OF_HEART_RATE_BY_CARDIAC_CONDUCTION
CAR_MLANA, CAR_MLANA
GO_SARCOMERE_ORGANIZATION, GO_SARCOMERE_ORGANIZATION
GO_CELL_DIFFERENTIATION_INVOLVED_IN_EMBRYONIC_PLACENTA_DEVELOPMENT, GO_CELL_DIFFERENTIATION_INVOLVED_IN_EMBRYONIC_PLACENTA_D
VANDESLUIS_COMMD1_TARGETS_GROUP_4_UP, VANDESLUIS_COMMD1_TARGETS_GROUP_4_UP
GO_LENS_FIBER_CELL_DIFFERENTIATION, GO_LENS_FIBER_CELL_DIFFERENTIATION
EDIATE_FILAMENT, GO_INTERMEDIATE_FILAMENT