LEF1_UP.V1_UP, LEF1_UP.V1_UP

P53_DN.V1_UP, P53_DN.V1_UP
PEREZ_TP53_AND_TP63_TARGETS, PEREZ_TP53_AND_TP63_TARGETS
BCAT.100_UP.V1_UP, BCAT.100_UP.V1_UP
KRAS.KIDNEY_UP.V1_UP, KRAS.KIDNEY_UP.V1_UP
GO_EAR_DEVELOPMENT, GO_EAR_DEVELOPMENT
MEISSNER_NPC_HCP_WITH_H3K4ME3_AND_H3K27ME3, MEISSNER_NPC_HCP_WITH_H3K4ME3_AND_H3K27ME3
GSE22025_UNTREATED_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_UP, GSE22025_UNTREATED_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_UP
HOOI_ST7_TARGETS_DN, HOOI_ST7_TARGETS_DN
HAMAI_APOPTOSIS_VIA_TRAIL_DN, HAMAI_APOPTOSIS_VIA_TRAIL_DN
GO_HEPARIN_BINDING, GO_HEPARIN_BINDING
DACOSTA_ERCC3_ALLELE_XPCS_VS_TTD_UP, DACOSTA_ERCC3_ALLELE_XPCS_VS_TTD_UP
VERHAAK_GLIOBLASTOMA_CLASSICAL, VERHAAK_GLIOBLASTOMA_CLASSICAL
IL21_UP.V1_UP, IL21_UP.V1_UP
NOTCH_DN.V1_DN, NOTCH_DN.V1_DN
ATF2_S_UP.V1_UP, ATF2_S_UP.V1_UP
GSE32986_GMCSF_AND_CURDLAN_LOWDOSE_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_DN, GSE32986_GMCSF_AND_CURDLAN_LOWDOSE_VS_GMCSF_AND_CURDLAN_HIGH
NOUSHMEHR_GBM_SILENCED_BY_METHYLATION, NOUSHMEHR_GBM_SILENCED_BY_METHYLATION
KRAS.LUNG.BREAST_UP.V1_DN, KRAS.LUNG.BREAST_UP.V1_DN
ISSAEVA_MLL2_TARGETS, ISSAEVA_MLL2_TARGETS
MOHANKUMAR_TLX1_TARGETS_DN, MOHANKUMAR_TLX1_TARGETS_DN
RICKMAN_HEAD_AND_NECK_CANCER_F, RICKMAN_HEAD_AND_NECK_CANCER_F
GO_MESENCHYME_DEVELOPMENT, GO_MESENCHYME_DEVELOPMENT
MODULE_139, MODULE_139
STEGER_ADIPOGENESIS_DN, STEGER_ADIPOGENESIS_DN
LEE_TARGETS_OF_PTCH1_AND_SUFU_DN, LEE_TARGETS_OF_PTCH1_AND_SUFU_DN
MODULE_385, MODULE_385
GSE36392_EOSINOPHIL_VS_MAC_IL25_TREATED_LUNG_UP, GSE36392_EOSINOPHIL_VS_MAC_IL25_TREATED_LUNG_UP
HOELZEL_NF1_TARGETS_DN, HOELZEL_NF1_TARGETS_DN
WILCOX_RESPONSE_TO_PROGESTERONE_DN, WILCOX_RESPONSE_TO_PROGESTERONE_DN
GO_DOPAMINE_METABOLIC_PROCESS, GO_DOPAMINE_METABOLIC_PROCESS
GO_NEGATIVE_CHEMOTAXIS, GO_NEGATIVE_CHEMOTAXIS
GO_VASCULAR_SMOOTH_MUSCLE_CELL_DIFFERENTIATION, GO_VASCULAR_SMOOTH_MUSCLE_CELL_DIFFERENTIATION
GO_OXYGEN_BINDING, GO_OXYGEN_BINDING
ANASTASSIOU_MULTICANCER_INVASIVENESS_SIGNATURE, ANASTASSIOU_MULTICANCER_INVASIVENESS_SIGNATURE
PTEN_DN.V1_DN, PTEN_DN.V1_DN
MODULE_190, MODULE_190
MODULE_329, MODULE_329
GO_INNER_EAR_MORPHOGENESIS, GO_INNER_EAR_MORPHOGENESIS
GO_REGULATION_OF_ORGAN_FORMATION, GO_REGULATION_OF_ORGAN_FORMATION
GO_CATECHOLAMINE_BIOSYNTHETIC_PROCESS, GO_CATECHOLAMINE_BIOSYNTHETIC_PROCESS
MODULE_387, MODULE_387
GSE24210_IL35_TREATED_VS_RESTING_TREG_DN, GSE24210_IL35_TREATED_VS_RESTING_TREG_DN
chr11p13, chr11p13
GSE2405_0H_VS_3H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_DN, GSE2405_0H_VS_3H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_DN
GSE13411_IGM_VS_SWITCHED_MEMORY_BCELL_UP, GSE13411_IGM_VS_SWITCHED_MEMORY_BCELL_UP
MODULE_131, MODULE_131
JI_CARCINOGENESIS_BY_KRAS_AND_STK11_DN, JI_CARCINOGENESIS_BY_KRAS_AND_STK11_DN
MODULE_557, MODULE_557
GO_REGULATION_OF_CARDIAC_MUSCLE_TISSUE_DEVELOPMENT, GO_REGULATION_OF_CARDIAC_MUSCLE_TISSUE_DEVELOPMENT
MODULE_497, MODULE_497
GO_NEGATIVE_REGULATION_OF_KIDNEY_DEVELOPMENT, GO_NEGATIVE_REGULATION_OF_KIDNEY_DEVELOPMENT
REACTOME_FGFR_LIGAND_BINDING_AND_ACTIVATION, REACTOME_FGFR_LIGAND_BINDING_AND_ACTIVATION
REACTOME_INTERACTION_BETWEEN_L1_AND_ANKYRINS, REACTOME_INTERACTION_BETWEEN_L1_AND_ANKYRINS