ALCALAY_AML_BY_NPM1_LOCALIZATION_UP, ALCALAY_AML_BY_NPM1_LOCALIZATION_UP

GSE30971_WBP7_HET_VS_KO_MACROPHAGE_2H_LPS_STIM_DN, GSE30971_WBP7_HET_VS_KO_MACROPHAGE_2H_LPS_STIM_DN
MIKKELSEN_IPS_WITH_HCP_H3K27ME3, MIKKELSEN_IPS_WITH_HCP_H3K27ME3
AKT_UP.V1_UP, AKT_UP.V1_UP
PID_INTEGRIN1_PATHWAY, PID_INTEGRIN1_PATHWAY
ROSS_AML_WITH_PML_RARA_FUSION, ROSS_AML_WITH_PML_RARA_FUSION
TTTNNANAGCYR_UNKNOWN, TTTNNANAGCYR_UNKNOWN
DACOSTA_ERCC3_ALLELE_XPCS_VS_TTD_UP, DACOSTA_ERCC3_ALLELE_XPCS_VS_TTD_UP
GO_DEFENSE_RESPONSE_TO_GRAM_POSITIVE_BACTERIUM, GO_DEFENSE_RESPONSE_TO_GRAM_POSITIVE_BACTERIUM
GO_NEGATIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION
LIEN_BREAST_CARCINOMA_METAPLASTIC_VS_DUCTAL_DN, LIEN_BREAST_CARCINOMA_METAPLASTIC_VS_DUCTAL_DN
MODULE_139, MODULE_139
GSE45365_BCELL_VS_CD8_TCELL_DN, GSE45365_BCELL_VS_CD8_TCELL_DN
GO_METANEPHRIC_NEPHRON_DEVELOPMENT, GO_METANEPHRIC_NEPHRON_DEVELOPMENT
JI_CARCINOGENESIS_BY_KRAS_AND_STK11_DN, JI_CARCINOGENESIS_BY_KRAS_AND_STK11_DN
GO_TRANSFORMING_GROWTH_FACTOR_BETA_BINDING, GO_TRANSFORMING_GROWTH_FACTOR_BETA_BINDING
MEISSNER_NPC_HCP_WITH_H3K27ME3, MEISSNER_NPC_HCP_WITH_H3K27ME3
WANG_THOC1_TARGETS_DN, WANG_THOC1_TARGETS_DN
GATA_Q6, GATA_Q6
GO_EXCRETION, GO_EXCRETION
GO_METANEPHROS_DEVELOPMENT, GO_METANEPHROS_DEVELOPMENT
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_15, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_15
REACTOME_REGULATION_OF_INSULIN_LIKE_GROWTH_FACTOR_IGF_ACTIVITY_BY_INSULIN_LIKE_GROWTH_FACTOR_BINDING_PROTEINS_IGFBPS, REACTOME_REGULATION_OF_INSULIN_LIKE_GROWTH_FACTOR_IGF_ACTIVITY_BY_INSULIN_LIKE_GROWTH_FACTOR_BINI
MATZUK_IMPLANTATION_AND_UTERINE, MATZUK_IMPLANTATION_AND_UTERINE
GO_NERVE_DEVELOPMENT, GO_NERVE_DEVELOPMENT
LEIN_CHOROID_PLEXUS_MARKERS, LEIN_CHOROID_PLEXUS_MARKERS
GO_METANEPHRIC_EPITHELIUM_DEVELOPMENT, GO_METANEPHRIC_EPITHELIUM_DEVELOPMENT
GO_POSITIVE_REGULATION_OF_KIDNEY_DEVELOPMENT, GO_POSITIVE_REGULATION_OF_KIDNEY_DEVELOPMENT
GO_ENDOCARDIAL_CUSHION_DEVELOPMENT, GO_ENDOCARDIAL_CUSHION_DEVELOPMENT
VERRECCHIA_DELAYED_RESPONSE_TO_TGFB1, VERRECCHIA_DELAYED_RESPONSE_TO_TGFB1
GO_CARDIAC_CELL_DEVELOPMENT, GO_CARDIAC_CELL_DEVELOPMENT
DAWSON_METHYLATED_IN_LYMPHOMA_TCL1, DAWSON_METHYLATED_IN_LYMPHOMA_TCL1
LIU_CDX2_TARGETS_UP, LIU_CDX2_TARGETS_UP
GO_CRANIAL_NERVE_DEVELOPMENT, GO_CRANIAL_NERVE_DEVELOPMENT
GO_INNER_EAR_MORPHOGENESIS, GO_INNER_EAR_MORPHOGENESIS
GO_EMBRYONIC_SKELETAL_SYSTEM_DEVELOPMENT, GO_EMBRYONIC_SKELETAL_SYSTEM_DEVELOPMENT
HOXA3_01, HOXA3_01
GO_PHARYNGEAL_SYSTEM_DEVELOPMENT, GO_PHARYNGEAL_SYSTEM_DEVELOPMENT