

ACID_DEPRIVATION, KRIGE_AMINO_ACID_DEPRIVATION

MTOR_UP.N4.V1_UP, MTOR_UP.N4.V1_UP
TOOKER_GEMCITABINE_RESISTANCE_DN, TOOKER_GEMCITABINE_RESISTANCE_DN
GSE3039_NKT_CELL_VS_B2_BCELL_DN, GSE3039_NKT_CELL_VS_B2_BCELL_DN
ZHAN_MULTIPLE_MYELOMA_CD1_VS_CD2_UP, ZHAN_MULTIPLE_MYELOMA_CD1_VS_CD2_UP
GSE45365_HEALTHY_VS_MCMV_INFECTION_BCELL_IFNAR_KO_DN, GSE45365_HEALTHY_VS_MCMV_INFECTION_BCELL_IFNAR_KO_DN
ZHAN_MULTIPLE_MYELOMA_CD1_UP, ZHAN_MULTIPLE_MYELOMA_CD1_UP
SABATES_COLORECTAL_ADENOMA_UP, SABATES_COLORECTAL_ADENOMA_UP
HELLEBREKERS_SILENCED_DURING_TUMOR_ANGIOGENESIS, HELLEBREKERS_SILENCED_DURING_TUMOR_ANGIOGENESIS
KERLEY_RESPONSE_TO_CISPLATIN_UP, KERLEY_RESPONSE_TO_CISPLATIN_UP
GSE12392_WT_VS_IFNB_KO_CD8A_POS_SPLEEN_DC_DN, GSE12392_WT_VS_IFNB_KO_CD8A_POS_SPLEEN_DC_DN
CTIP_DN.V1_UP, CTIP_DN.V1_UP
REACTOME_TRANSPORT_OF_INORGANIC_CATIONS_ANIONS_AND_AMINO_ACIDS_OLIGOPEPTIDES, REACTOME_TRANSPORT_OF_INORGANIC_CATIONS_ANIONS_AND_AMINO_
KANG_FLUOROURACIL_RESISTANCE_DN, KANG_FLUOROURACIL_RESISTANCE_DN
GO_POSITIVE_REGULATION_OF_CHEMOTAXIS, GO_POSITIVE_REGULATION_OF_CHEMOTAXIS
DCA_UP.V1_UP, DCA_UP.V1_UP
GSE40666_STAT1_KO_VS_STAT4_KO_CD8_TCELL_WITH_IFNA_STIM_90MIN_DN, GSE40666_STAT1_KO_VS_STAT4_KO_CD8_TCELL_WITH_IFNA_STIM_90MIN_DN
GO_AMINO_ACID_TRANSMEMBRANE_TRANSPORT, GO_AMINO_ACID_TRANSMEMBRANE_TRANSPORT
GO_PYRIDOXAL_PHOSPHATE_BINDING, GO_PYRIDOXAL_PHOSPHATE_BINDING
KEGG_GLYCINE_SERINE_AND_THREONINE_METABOLISM, KEGG_GLYCINE_SERINE_AND_THREONINE_METABOLISM
GO_REGULATION_OF_INTERLEUKIN_4_PRODUCTION, GO_REGULATION_OF_INTERLEUKIN_4_PRODUCTION
MOHANKUMAR_TLX1_TARGETS_DN, MOHANKUMAR_TLX1_TARGETS_DN
LY_AGING_MIDDLE_UP, LY_AGING_MIDDLE_UP
CHO_NR4A1_TARGETS, CHO_NR4A1_TARGETS
GO_NEGATIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_IN_ABSENCE_OF_LIGAND, GO_NEGATIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_IN_ABSENCE_OF_LIGAND
GSE40274_IRF4_VS_FOXP3_AND_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_IRF4_VS_FOXP3_AND_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GO_RESPONSE_TO_DEXAMETHASONE, GO_RESPONSE_TO_DEXAMETHASONE
GO_DICARBOXYLIC_ACID_TRANSPORT, GO_DICARBOXYLIC_ACID_TRANSPORT
BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_48HR_DN, BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_48HR_DN
SMID_BREAST_CANCER_RELAPSE_IN_PLEURA_DN, SMID_BREAST_CANCER_RELAPSE_IN_PLEURA_DN
GO_REGULATION_OF_MESENCHYMAL_CELL_PROLIFERATION, GO_REGULATION_OF_MESENCHYMAL_CELL_PROLIFERATION
GO_NEGATIVE_REGULATION_OF_BEHAVIOR, GO_NEGATIVE_REGULATION_OF_BEHAVIOR
GO_EYE_PHOTORECEPTOR_CELL_DEVELOPMENT, GO_EYE_PHOTORECEPTOR_CELL_DEVELOPMENT
GO_NEGATIVE_REGULATION_OF_ASTROCYTE_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_ASTROCYTE_DIFFERENTIATION
GO_EYE_PHOTORECEPTOR_CELL_DIFFERENTIATION, GO_EYE_PHOTORECEPTOR_CELL_DIFFERENTIATION
GO_POSITIVE_REGULATION_OF_KIDNEY_DEVELOPMENT, GO_POSITIVE_REGULATION_OF_KIDNEY_DEVELOPMENT
GO_MODIFIED_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GO_MODIFIED_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
BIOCARTA_STEM_PATHWAY, BIOCARTA_STEM_PATHWAY
GO_DETECTION_OF_MECHANICAL_STIMULUS, GO_DETECTION_OF_MECHANICAL_STIMULUS
GO_ION_ANTIPORTER_ACTIVITY, GO_ION_ANTIPORTER_ACTIVITY
GO_NEGATIVE_REGULATION_OF_TRANSCRIPTION_REGULATORY_REGION_DNA_BINDING, GO_NEGATIVE_REGULATION_OF_TRANSCRIPTION_REGULATORY_REGION_DNA_BIN
GO_LEUCINE_ZIPPER_DOMAIN_BINDING, GO_LEUCINE_ZIPPER_DOMAIN_BINDING
SU_LIVER, SU_LIVER
GO_RECEPTOR_AGONIST_ACTIVITY, GO_RECEPTOR_AGONIST_ACTIVITY
GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_4H_ACT_CD4_TCELL_DN, GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_4H_ACT_CD4_TCELL_DN