
GSE7764_IL15_NK_CELL_24H_VS_SPLENOCYTE_UP, GSE7764_IL15_NK_CELL_24H_VS_SPLENOCYTE_UP
GSE40274_CTRL_VS_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE20754_WT_VS_TCF1_KO_MEMORY_CD8_TCELL_UP, GSE20754_WT_VS_TCF1_KO_MEMORY_CD8_TCELL_UP
GSE20715_WT_VS_TLR4_KO_LUNG_DN, GSE20715_WT_VS_TLR4_KO_LUNG_DN
GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_DN, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_DN
REACTOME_PEROXISOMAL_PROTEIN_IMPORT, REACTOME_PEROXISOMAL_PROTEIN_IMPORT
QI_HYPOXIA, QI_HYPOXIA
GSE36527_CD62L_HIGH_CD69_NEG_VS_CD62L_LOW_CD69_POS_TREG_KLRG1_NEG_DN, GSE36527_CD62L_HIGH_CD69_NEG_VS_CD62L_LOW_CD69_POS_TREG_KLRG1_NEG_
MIR6771_3P, MIR6771_3P
GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_RRNA, GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_RRNA
GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_DN, GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_DN
GSE29614_DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMC_UP, GSE29614_DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMC_UP
HP_DECREASED_SKULL_OSSIFICATION, HP_DECREASED_SKULL_OSSIFICATION
GOBP_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_IMPORT_ACROSS_PLASMA_MEMBRANE
REACTOME_CHOLESTEROL_BIOSYNTHESIS, REACTOME_CHOLESTEROL_BIOSYNTHESIS
NIKOLSKY_BREAST_CANCER_16Q24_AMPLICON, NIKOLSKY_BREAST_CANCER_16Q24_AMPLICON
WENG_POR_TARGETS_GLOBAL_UP, WENG_POR_TARGETS_GLOBAL_UP
GOBP_OLEFINIC_COMPOUND_METABOLIC_PROCESS, GOBP_OLEFINIC_COMPOUND_METABOLIC_PROCESS
MIR4300, MIR4300
GOCC_RNA_POLYMERASE_II_CORE_COMPLEX, GOCC_RNA_POLYMERASE_II_CORE_COMPLEX
REACTOME_PIWI_INTERACTING_RNA_PIRNA_BIOGENESIS, REACTOME_PIWI_INTERACTING_RNA_PIRNA_BIOGENESIS
GOBP_AMINO_ACID_TRANSMEMBRANE_TRANSPORT, GOBP_AMINO_ACID_TRANSMEMBRANE_TRANSPORT
WENG_POR_TARGETS_LIVER_UP, WENG_POR_TARGETS_LIVER_UP
MODULE_234, MODULE_234
GOBP_AMINO_ACID_IMPORT, GOBP_AMINO_ACID_IMPORT
REACTOME_TRANSPORT_OF_INORGANIC_CATIONS_ANIONS_AND_AMINO_ACIDS_OLIGOPEPTIDES, REACTOME_TRANSPORT_OF_INORGANIC_CATIONS_ANIONS_AND_
HEIDENBLAD_AMPLICON_8Q24_DN, HEIDENBLAD_AMPLICON_8Q24_DN
GOMF_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
MIR6798_5P, MIR6798_5P
SERVITJA_ISLET_HNF1A_TARGETS_DN, SERVITJA_ISLET_HNF1A_TARGETS_DN
GOMF_SULFUR_COMPOUND_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_SULFUR_COMPOUND_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOMF_L_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_L_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_NEGATIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION, GOBP_NEGATIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION
GNF2_TM4SF2, GNF2_TM4SF2
REACTOME_COLLAGEN_DEGRADATION, REACTOME_COLLAGEN_DEGRADATION
GOBP_CELLULAR_RESPONSE_TO_PH, GOBP_CELLULAR_RESPONSE_TO_PH
HP_SELECTIVE_TOOTH_AGENESIS, HP_SELECTIVE_TOOTH_AGENESIS
HP_AGENESIS_OF_INCISOR, HP_AGENESIS_OF_INCISOR
GOBP_AZOLE_TRANSMEMBRANE_TRANSPORT, GOBP_AZOLE_TRANSMEMBRANE_TRANSPORT
URS_ADIPOCYTE_DIFFERENTIATION_DN, URS_ADIPOCYTE_DIFFERENTIATION_DN
GOBP_PYRIMIDINE_CONTAINING_COMPOUND_TRANSMEMBRANE_TRANSPORT, GOBP_PYRIMIDINE_CONTAINING_COMPOUND_TRANSMEMBRANE_TRANSPORT
REACTOME_ASSEMBLY_OF_COLLAGEN_FIBRILS_AND_OTHER_MULTIMERIC_STRUCTURES, REACTOME_ASSEMBLY_OF_COLLAGEN_FIBRILS_AND_OTHER_MULTIMERIC_S
ZNF529_TARGET_GENES, ZNF529_TARGET_GENES
GOMF_AZOLE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_AZOLE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
NAKAMURA_ADIPOGENESIS_LATE_DN, NAKAMURA_ADIPOGENESIS_LATE_DN
FRIDMAN_IMMORTALIZATION_DN, FRIDMAN_IMMORTALIZATION_DN
HP_SPARSE_EYELASHES, HP_SPARSE_EYELASHES
GOBP_SERINE_FAMILY_AMINO_ACID_CATABOLIC_PROCESS, GOBP_SERINE_FAMILY_AMINO_ACID_CATABOLIC_PROCESS
REACTOME_AMINO_ACID_TRANSPORT_ACROSS_THE_PLASMA_MEMBRANE, REACTOME_AMINO_ACID_TRANSPORT_ACROSS_THE_PLASMA_MEMBRANE
SPIRA_SMOKERS_LUNG_CANCER_DN, SPIRA_SMOKERS_LUNG_CANCER_DN
NAKAMURA_ADIPOGENESIS_EARLY_DN, NAKAMURA_ADIPOGENESIS_EARLY_DN
GOBP_LONG_CHAIN_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS, GOBP_LONG_CHAIN_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS
WP_TYPE_I_COLLAGEN_SYNTHESIS_IN_THE_CONTEXT_OF_OSTEOGENESIS_IMPERFECTA, WP_TYPE_I_COLLAGEN_SYNTHESIS_IN_THE_CONTEXT_OF_OSTEOGENESIS_IMP
GOBP_RESPONSE_TO_PH, GOBP_RESPONSE_TO_PH
HP_BROAD_PHALANX_OF_THE_TOES, HP_BROAD_PHALANX_OF_THE_TOES
HP_RADIAL_DEVIATION_OF_THE_HAND_OR_OF_FINGERS_OF_THE_HAND, HP_RADIAL_DEVIATION_OF_THE_HAND_OR_OF_FINGERS_OF_THE_HAND
GOMF_GAMMA_TUBULIN_BINDING, GOMF_GAMMA_TUBULIN_BINDING
HAY_BONE_MARROW_CD34_POS_MKP, HAY_BONE_MARROW_CD34_POS_MKP
GOMF_BASIC_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_BASIC_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
MIR6845_5P, MIR6845_5P
HP_FLARED_METAPHYSIS, HP_FLARED_METAPHYSIS
MIR1470, MIR1470
GOBP_NEUROEPITHELIAL_CELL_DIFFERENTIATION, GOBP_NEUROEPITHELIAL_CELL_DIFFERENTIATION
REACTOME_COLLAGEN_FORMATION, REACTOME_COLLAGEN_FORMATION
STEGER_ADIPOGENESIS_DN, STEGER_ADIPOGENESIS_DN
GOBP_POTASSIUM_ION_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_POTASSIUM_ION_IMPORT_ACROSS_PLASMA_MEMBRANE
MODULE_141, MODULE_141
GOBP_MUSCLE_FIBER_DEVELOPMENT, GOBP_MUSCLE_FIBER_DEVELOPMENT
GOBP_EPITHELIAL_CELL_FATE_COMMITMENT, GOBP_EPITHELIAL_CELL_FATE_COMMITMENT
GOBP_METANEPHRIC_TUBULE_FORMATION, GOBP_METANEPHRIC_TUBULE_FORMATION
_VS_KLRG1HIGH_EFF_CD8_TCELL_UP, GSE10239_KLRG1INT_VS_KLRG1HIGH_EFF_CD8_TCELL_UP