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GSE39820_CTRL_VS_TGFBETA3_IL6_CD4_TCELL_DN, GSE39820_CTRL_VS_TGFBETA3_IL6_CD4_TCELL_DN
GSE6674_CPG_VS_CPG_AND_ANTI_IGM_STIM_BCELL_DN, GSE6674_CPG_VS_CPG_AND_ANTI_IGM_STIM_BCELL_DN
GSE39820 CTRL VS TGFBETA1 IL6 CD4 TCELL DN, GSE39820 CTRL VS TGFBETA1 IL6 CD4 TCELL DN
GSE17721_LPS_VS_GARDIQUIMOD_12H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_12H_BMDC_DN
WP_EGFEGFR_SIGNALING_PATHWAY, WP_EGFEGFR_SIGNALING_PATHWAY
MIR302A_5P, MIR302A_5P
GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_UP, GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_UP
GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_DN, GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_DN
GSE17721_0.5H_VS_8H_POLYIC_BMDC_DN, GSE17721_0.5H_VS_8H_POLYIC_BMDC_DN
MIR217_5P, MIR217_5P
GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_UP, GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_UP
GSE17721_0.5H_VS_8H_LPS_BMDC_DN, GSE17721_0.5H_VS_8H_LPS_BMDC_DN
GSE17721_CTRL_VS_CPG_8H_BMDC_DN, GSE17721_CTRL_VS_CPG_8H_BMDC_DN
GSE39820_TGFBETA3_IL6_VS_TGFBETA3_IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA3_IL6_VS_TGFBETA3_IL6_IL23A_TREATED_CD4_TC
MIR337_3P, MIR337_3P
HP_NARROW_PALATE, HP_NARROW_PALATE
MIR6807_3P, MIR6807_3P
GSE13493_DP_VS_CD4INTCD8POS_THYMOCYTE_UP, GSE13493_DP_VS_CD4INTCD8POS_THYMOCYTE_UP
 GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCE
TOOKER_GEMCITABINE_RESISTANCE_DN, TOOKER_GEMCITABINE_RESISTANCE_DN
ATAGGAA_MIR202, ATAGGAA_MIR202
MIR4426, MIR4426
MIR5689, MIR5689
MIR708_3P, MIR708_3P
MIR509_5P, MIR509_5P
MIR4418, MIR4418
HP_HIGH_NARROW_PALATE, HP_HIGH_NARROW_PALATE
MIR509_3_5P, MIR509_3_5P
 ATGCAGT_MIR217, ATGCAGT_MIR217
HP_DEVIATION_OF_TOES, HP_DEVIATION_OF_TOES
 MIR8076, MIR8076
MIR433_3P, MIR433_3P
GSE14026_TH1_VS_TH17_DN, GSE14026_TH1_VS_TH17_DN
MIR6828_5P, MIR6828_5P
NAKAYA_MONOCYTE_FLUARIX_FLUVIRIN_AGE_18_50YO_7DY_UP, NAKAYA_MONOCYTE_FLUARIX_FLUVIRIN_AGE_18_50YO_7DY_UP
PANGAS_TUMOR_SUPPRESSION_BY_SMAD1_AND_SMAD5_DN, PANGAS_TUMOR_SUPPRESSION_BY_SMAD1_AND_SMAD5_DN
MIR4645_5P, MIR4645_5P
GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN
HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANGES_OF_THE_HAND, HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANGES_OF_THE_H
HP_CUTIS_MARMORATA, HP_CUTIS_MARMORATA
MIR374A_3P, MIR374A_3P
MIR6759_3P, MIR6759_3P
HP_LOWER_LIMB_MUSCLE_WEAKNESS, HP_LOWER_LIMB_MUSCLE_WEAKNESS
BOYAULT_LIVER_CANCER_SUBCLASS_G12_UP, BOYAULT_LIVER_CANCER_SUBCLASS_G12_UP
MIR361_5P, MIR361_5P
GCTTGAA MIR498, GCTTGAA MIR498
GSE41176_UNSTIM_VS_ANTI_IGM_STIM_TAK1_KO_BCELL_1H_DN, GSE41176_UNSTIM_VS_ANTI_IGM_STIM_TAK1_KO_BCELL_1H_DN
GOCC_CAVEOLA, GOCC_CAVEOLA
HP_EXODEVIATION, HP_EXODEVIATION
TBK1.DN.48HRS_DN, TBK1.DN.48HRS_DN
GOCC_PLASMA_MEMBRANE_RAFT, GOCC_PLASMA_MEMBRANE_RAFT
HP_ARTERIAL_STENOSIS, HP_ARTERIAL_STENOSIS
MIR518D_5P_MIR518F_5P_MIR520C_5P_MIR526A_5P, MIR518D_5P_MIR518F_5P_MIR520C_5P_MIR526A_5P
FALVELLA_SMOKERS_WITH_LUNG_CANCER, FALVELLA_SMOKERS_WITH_LUNG_CANCER
GSE4142_NAIVE_VS_MEMORY_BCELL_UP, GSE4142_NAIVE_VS_MEMORY_BCELL_UP
GOCC_ENDOPLASMIC_RETICULUM_QUALITY_CONTROL_COMPARTMENT, GOCC_ENDOPLASMIC_RETICULUM_QUALITY_CONTROL_COMPARTM
MIR518E 5P MIR519A 5P MIR519B 5P MIR519C 5P MIR522 5P MIR523 5P, MIR518E 5P MIR519A 5P MIR519B 5P MIR519C 5P MIR522 5P MIR523 5P
MIR216B_5P, MIR216B_5P
GOMF_PROTEIN_TYROSINE_KINASE_BINDING, GOMF_PROTEIN_TYROSINE_KINASE_BINDING
GSE369_SOCS3_KO_VS_IFNG_KO_LIVER_UP, GSE369_SOCS3_KO_VS_IFNG_KO_LIVER_UP
WP_TGFBETA_RECEPTOR_SIGNALLING_IN_SKELETAL_DYSPLASIAS, WP_TGFBETA_RECEPTOR_SIGNALLING_IN_SKELETAL_DYSPLASIAS
REACTOME_INTEGRATION_OF_ENERGY_METABOLISM, REACTOME_INTEGRATION_OF_ENERGY_METABOLISM
REACTOME_GLYCEROPHOSPHOLIPID_BIOSYNTHESIS, REACTOME_GLYCEROPHOSPHOLIPID_BIOSYNTHESIS
MIR938, MIR938
REACTOME_INTRA_GOLGI_TRAFFIC, REACTOME_INTRA_GOLGI_TRAFFIC
CCATCCA_MIR432, CCATCCA_MIR432
MIR1197, MIR1197
WP_TGFBETA_RECEPTOR_SIGNALING, WP_TGFBETA_RECEPTOR_SIGNALING
GOCC_ACTOMYOSIN, GOCC_ACTOMYOSIN
GOBP_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION, GOBP_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NU
 GOBP_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE, GOBP_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE
 GOBP_BONE_MINERALIZATION, GOBP_BONE_MINERALIZATION
GOBP_CARDIAC_VENTRICLE_DEVELOPMENT, GOBP_CARDIAC_VENTRICLE_DEVELOPMENT
HP_TRANSIENT_ISCHEMIC_ATTACK, HP_TRANSIENT_ISCHEMIC_ATTACK
GOBP_RESPONSE_TO_BMP, GOBP_RESPONSE_TO_BMP
GOBP_CELL_CELL_JUNCTION_ASSEMBLY, GOBP_CELL_CELL_JUNCTION_ASSEMBLY
 GOBP SYNAPTIC VESICLE PRIMING, GOBP SYNAPTIC VESICLE PRIMING
REACTOME_SIGNALING_BY_ERBB2, REACTOME_SIGNALING_BY_ERBB2
MIR4323, MIR4323
GOBP_KIDNEY_EPITHELIUM_DEVELOPMENT, GOBP_KIDNEY_EPITHELIUM_DEVELOPMENT
GOBP_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION, GOBP_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION
GOBP_EMBRYONIC_SKELETAL_SYSTEM_MORPHOGENESIS, GOBP_EMBRYONIC_SKELETAL_SYSTEM_MORPHOGENESIS
MIR4776_5P, MIR4776_5P
GCM_RING1, GCM_RING1
FXR_IR1_Q6, FXR_IR1_Q6
GOBP_EMBRYONIC_CRANIAL_SKELETON_MORPHOGENESIS, GOBP_EMBRYONIC_CRANIAL_SKELETON_MORPHOGENESIS
GOBP_SPECIFICATION_OF_SYMMETRY, GOBP_SPECIFICATION_OF_SYMMETRY
MIR6856 3P, MIR6856 3P
REACTOME_SIGNALING_BY_MODERATE_KINASE_ACTIVITY_BRAF_MUTANTS, REACTOME_SIGNALING_BY_MODERATE_KINASE_ACTIVITY_BRAF_
HP_ABNORMAL_5TH_FINGER_PHALANX_MORPHOLOGY, HP_ABNORMAL_5TH_FINGER_PHALANX_MORPHOLOGY
HP_AORTIC_DISSECTION, HP_AORTIC_DISSECTION
GOMF_MICROTUBULE_PLUS_END_BINDING, GOMF_MICROTUBULE_PLUS_END_BINDING
GOMF_TRANSMEMBRANE_RECEPTOR_PROTEIN_KINASE_ACTIVITY, GOMF_TRANSMEMBRANE_RECEPTOR_PROTEIN_KINASE_ACTIVITY
GOBP_CRANIAL_SKELETAL_SYSTEM_DEVELOPMENT, GOBP_CRANIAL_SKELETAL_SYSTEM_DEVELOPMENT
HP_INVERTED_NIPPLES, HP_INVERTED_NIPPLES
GSE41978_WT_VS_ID2_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_UP, GSE41978_WT_VS_ID2_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_UP
GOBP BONE MORPHOGENESIS, GOBP BONE MORPHOGENESIS
MIR4278, MIR4278
GOBP_REGULATION_OF_BONE_MINERALIZATION, GOBP_REGULATION_OF_BONE_MINERALIZATION
BIOCARTA_ARAP_PATHWAY, BIOCARTA_ARAP_PATHWAY
BIOCARTA_LIS1_PATHWAY, BIOCARTA_LIS1_PATHWAY
 GOBP_VENTRICULAR_SEPTUM_DEVELOPMENT, GOBP_VENTRICULAR_SEPTUM_DEVELOPMENT
GOBP_BIOMINERALIZATION, GOBP_BIOMINERALIZATION
 HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANGES_OF_THE_TOES, HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANGES_OF_THE_TO
 GOBP_OUTFLOW_TRACT_MORPHOGENESIS, GOBP_OUTFLOW_TRACT_MORPHOGENESIS
 HP_DISTAL_LOWER_LIMB_AMYOTROPHY, HP_DISTAL_LOWER_LIMB_AMYOTROPHY
HP_ABDOMINAL_AORTIC_ANEURYSM, HP_ABDOMINAL_AORTIC_ANEURYSM
 GOBP_OUTFLOW_TRACT_SEPTUM_MORPHOGENESIS, GOBP_OUTFLOW_TRACT_SEPTUM_MORPHOGENESIS
 GOBP_NEGATIVE_REGULATION_OF_SMOOTHENED_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_SMOOTHENED_SIGNALING_PAT
 HP_DESCENDING_THORACIC_AORTA_ANEURYSM, HP_DESCENDING_THORACIC_AORTA_ANEURYSM
 GOBP_ACTIN_FILAMENT_DEPOLYMERIZATION, GOBP_ACTIN_FILAMENT_DEPOLYMERIZATION
XU_AKT1_TARGETS_6HR, XU_AKT1_TARGETS_6HR
 HP_ABNORMALITY OF THE DISTAL PHALANGES OF THE TOES, HP_ABNORMALITY OF THE DISTAL PHALANGES OF THE TOES
GOCC_CATENIN_COMPLEX, GOCC_CATENIN_COMPLEX
REACTOME_CELL_CELL_JUNCTION_ORGANIZATION, REACTOME_CELL_CELL_JUNCTION_ORGANIZATION
HP_VASCULAR_TORTUOSITY, HP_VASCULAR_TORTUOSITY
REACTOME_ADHERENS_JUNCTIONS_INTERACTIONS, REACTOME_ADHERENS_JUNCTIONS_INTERACTIONS
HP_ABNORMAL_2ND_FINGER_MORPHOLOGY, HP_ABNORMAL_2ND_FINGER_MORPHOLOGY
ZNF488_TARGET_GENES, ZNF488_TARGET_GENES
GOBP_REGULATION_OF_CARDIAC_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_REGULATION_OF_CARDIAC_EPITHELIAL_TO_MESENCH
HP_LOWER_LIMB_AMYOTROPHY, HP_LOWER_LIMB_AMYOTROPHY
GOBP_CARDIAC_SEPTUM_MORPHOGENESIS, GOBP_CARDIAC_SEPTUM_MORPHOGENESIS
GOBP_LUNG_ALVEOLUS_DEVELOPMENT, GOBP_LUNG_ALVEOLUS_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_BONE_MINERALIZATION, GOBP_POSITIVE_REGULATION_OF_BONE_MINERALIZATION
HP_HEMIMEGALENCEPHALY, HP_HEMIMEGALENCEPHALY
GOBP_CHONDROCYTE_DEVELOPMENT, GOBP_CHONDROCYTE_DEVELOPMENT
WIKMAN_ASBESTOS_LUNG_CANCER_UP, WIKMAN_ASBESTOS_LUNG_CANCER_UP
REACTOME_YAP1_AND_WWTR1_TAZ_STIMULATED_GENE_EXPRESSION, REACTOME_YAP1_AND_WWTR1_TAZ_STIMULATED_GENE_EXPRESSION
GOMF_ACTIVIN_BINDING, GOMF_ACTIVIN_BINDING
GOBP_POSITIVE_REGULATION_OF_FOCAL_ADHESION_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_FOCAL_ADHESION_ASSEMBLY
MIR23A_5P, MIR23A_5P
REACTOME_AKT_PHOSPHORYLATES_TARGETS_IN_THE_NUCLEUS, REACTOME_AKT_PHOSPHORYLATES_TARGETS_IN_THE_NUCLEUS
BREUHAHN_GROWTH_FACTOR_SIGNALING_IN_LIVER_CANCER, BREUHAHN_GROWTH_FACTOR_SIGNALING_IN_LIVER_CANCER
GOBP_REGULATION_OF_SYNAPTIC_VESICLE_PRIMING, GOBP_REGULATION_OF_SYNAPTIC_VESICLE_PRIMING
HP_ASCENDING_AORTIC_DISSECTION, HP_ASCENDING_AORTIC_DISSECTION
GOMF_TRANSFORMING_GROWTH_FACTOR_BETA_BINDING, GOMF_TRANSFORMING_GROWTH_FACTOR_BETA_BINDING
GOBP_REGULATION_OF_NEUROBLAST_PROLIFERATION, GOBP_REGULATION_OF_NEUROBLAST_PROLIFERATION
MIR100_5P_MIR99A_5P_MIR99B_5P, MIR100_5P_MIR99A_5P_MIR99B_5P
GOBP_APICAL_JUNCTION_ASSEMBLY, GOBP_APICAL_JUNCTION_ASSEMBLY
GOBP_ERBB2_SIGNALING_PATHWAY, GOBP_ERBB2_SIGNALING_PATHWAY
BIOCARTA_TOB1_PATHWAY, BIOCARTA_TOB1_PATHWAY
PID_ECADHERIN_KERATINOCYTE_PATHWAY, PID_ECADHERIN_KERATINOCYTE_PATHWAY
GOBP_PROTEIN_LIPID_COMPLEX_ASSEMBLY, GOBP_PROTEIN_LIPID_COMPLEX_ASSEMBLY
STAMBOLSKY_RESPONSE_TO_VITAMIN_D3_DN, STAMBOLSKY_RESPONSE_TO_VITAMIN_D3_DN
HP_2_3_FINGER_SYNDACTYLY, HP_2_3_FINGER_SYNDACTYLY
GOMF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY, GOMF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINI
GOBP_POSITIVE_REGULATION_OF_BIOMINERALIZATION, GOBP_POSITIVE_REGULATION_OF_BIOMINERALIZATION
GOBP_ARTERY_MORPHOGENESIS, GOBP_ARTERY_MORPHOGENESIS
HP_APLASIA_HYPOPLASIA_OF_THE_PHALANGES_OF_THE_2ND_FINGER, HP_APLASIA_HYPOPLASIA_OF_THE_PHALANGES_OF_THE_2ND_FINGEI
GOMF_TRANSFORMING_GROWTH_FACTOR_BETA_RECEPTOR_BINDING, GOMF_TRANSFORMING_GROWTH_FACTOR_BETA_RECEPTOR_BINDING
 HP_SKIN_EROSION, HP_SKIN_EROSION
HP_KETONURIA, HP_KETONURIA
GOBP_PROTEIN_LIPID_COMPLEX_SUBUNIT_ORGANIZATION, GOBP_PROTEIN_LIPID_COMPLEX_SUBUNIT_ORGANIZATION
GOMF_MYOSIN_HEAVY_CHAIN_BINDING, GOMF_MYOSIN_HEAVY_CHAIN_BINDING
HP_SHORT_MIDDLE_PHALANX_OF_TOE, HP_SHORT_MIDDLE_PHALANX_OF_TOE
HP_DUPLICATION_OF_RENAL_PELVIS, HP_DUPLICATION_OF_RENAL_PELVIS
HP_ABNORMALITY_OF_THE_MIDDLE_PHALANX_OF_THE_2ND_FINGER, HP_ABNORMALITY_OF_THE_MIDDLE_PHALANX_OF_THE_2ND_FINGER
WP_BMP_SIGNALING_PATHWAY_IN_EYELID_DEVELOPMENT, WP_BMP_SIGNALING_PATHWAY_IN_EYELID_DEVELOPMENT
GOBP_NEPHRIC_DUCT_MORPHOGENESIS, GOBP_NEPHRIC_DUCT_MORPHOGENESIS
GOBP_MAINTENANCE_OF_SYNAPSE_STRUCTURE, GOBP_MAINTENANCE_OF_SYNAPSE_STRUCTURE
GOCC_HEMIDESMOSOME, GOCC_HEMIDESMOSOME
HP_ABNORMALITY_OF_THE_PHALANGES_OF_THE_2ND_FINGER, HP_ABNORMALITY_OF_THE_PHALANGES_OF_THE_2ND_FINGER
HP_ATRIAL_STANDSTILL, HP_ATRIAL_STANDSTILL
GOBP_POSITIVE_REGULATION_OF_CARTILAGE_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_CARTILAGE_DEVELOPMENT
GOBP_RECEPTOR_GUANYLYL_CYCLASE_SIGNALING_PATHWAY, GOBP_RECEPTOR_GUANYLYL_CYCLASE_SIGNALING_PATHWAY
GOBP_EMBRYONIC_SKELETAL_JOINT_MORPHOGENESIS, GOBP_EMBRYONIC_SKELETAL_JOINT_MORPHOGENESIS
GOBP_REGULATION_OF_ARTERY_MORPHOGENESIS, GOBP_REGULATION_OF_ARTERY_MORPHOGENESIS
GOBP_PHARYNGEAL_SYSTEM_DEVELOPMENT, GOBP_PHARYNGEAL_SYSTEM_DEVELOPMENT
HP_PROGRESSIVE_VISUAL_FIELD_DEFECTS, HP_PROGRESSIVE_VISUAL_FIELD_DEFECTS
HP_EPIGASTRIC_PAIN, HP_EPIGASTRIC_PAIN
HP_ABNORMALITY_OF_THE_DISTAL_PHALANX_OF_THE_HALLUX, HP_ABNORMALITY_OF_THE_DISTAL_PHALANX_OF_THE_HALLUX
GOMF_BMP_BINDING, GOMF_BMP_BINDING
REACTOME_ACETYLCHOLINE_NEUROTRANSMITTER_RELEASE_CYCLE, REACTOME_ACETYLCHOLINE_NEUROTRANSMITTER_RELEASE_CYCLE
GOBP POSITIVE REGULATION OF CARDIAC MUSCLE CONTRACTION, GOBP POSITIVE REGULATION OF CARDIAC MUSCLE CONTRACTION
GOBP_TOLERANCE_INDUCTION, GOBP_TOLERANCE_INDUCTION
GOCC_ZONULA_ADHERENS, GOCC_ZONULA_ADHERENS
HP_APLASIA_HYPOPLASIA_OF_THE_2ND_FINGER, HP_APLASIA_HYPOPLASIA_OF_THE_2ND_FINGER
HP_ABNORMALITY_OF_THE_MIDDLE_PHALANGES_OF_THE_TOES, HP_ABNORMALITY_OF_THE_MIDDLE_PHALANGES_OF_THE_TOES
HP_ABNORMALITY_OF_ENDOCRINE_PANCREAS_PHYSIOLOGY, HP_ABNORMALITY_OF_ENDOCRINE_PANCREAS_PHYSIOLOGY
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GSE39820_TGFBETA1_IL6_VS_TGFBETA1_IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_IL6_VS_TGFBETA1_IL6_IL23A_TREATED_CD4_TC

_TGFBETA3_IN_IL6_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_TREATED_CD4_TCELL_UP