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CTACTGT_MIR199A, CTACTGT_MIR199A
                                                                                               TGTGTGA MIR377, TGTGTGA MIR377
                                                                                                GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_UP, GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_UP
                                                                                               GSE22282_HYPOXIA_VS_NORMOXIA_MYELOID_DC_DN, GSE22282_HYPOXIA_VS_NORMOXIA_MYELOID_DC_DN
                                                                                               MIR425_5P, MIR425_5P
                                                                                               GSE30083_SP3_VS_SP4_THYMOCYTE_UP, GSE30083_SP3_VS_SP4_THYMOCYTE_UP
                                                                                               GSE24210_IL35_TREATED_VS_UNTREATED_TCONV_CD4_TCELL_DN, GSE24210_IL35_TREATED_VS_UNTREATED_TCONV_CD4_TCELL_DN
                                                                                               MIR4653_5P, MIR4653_5P
                                                                                               MIR3921, MIR3921
                                                                                               MIR5699_5P, MIR5699_5P
                                                                                               GAO_ESOPHAGUS_25W_C3_FGFR1LOW_EPITHELIAL_CELLS, GAO_ESOPHAGUS_25W_C3_FGFR1LOW_EPITHELIAL_CELLS
                                                                                               GSE46242_CTRL_VS_EGR2_DELETED_TH1_CD4_TCELL_DN, GSE46242_CTRL_VS_EGR2_DELETED_TH1_CD4_TCELL_DN
                                                                                                MIR7852 3P, MIR7852 3P
                                                                                                GOBP_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
                                                                                                MIR517_5P, MIR517_5P
                                                                                                GSE27786_BCELL_VS_ERYTHROBLAST_DN, GSE27786_BCELL_VS_ERYTHROBLAST_DN
                                                                                                MIR6856_5P, MIR6856_5P
                                                                                                WP_ARRHYTHMOGENIC_RIGHT_VENTRICULAR_CARDIOMYOPATHY, WP_ARRHYTHMOGENIC_RIGHT_VENTRICULAR_CARDIOMYOPATHY
                                                                                                KEGG_ARRHYTHMOGENIC_RIGHT_VENTRICULAR_CARDIOMYOPATHY_ARVC, KEGG_ARRHYTHMOGENIC_RIGHT_VENTRICULAR_CARDIOMYOPATHY_ARVC
                                                                                                RTTTNNNYTGGM_UNKNOWN, RTTTNNNYTGGM_UNKNOWN
                                                                                                HP_ABNORMALITIES_OF_PLACENTA_OR_UMBILICAL_CORD, HP_ABNORMALITIES_OF_PLACENTA_OR_UMBILICAL_CORD
                                                                                                GU_PDEF_TARGETS_UP, GU_PDEF_TARGETS_UP
                                                                                                GOBP N TERMINAL PROTEIN AMINO ACID MODIFICATION, GOBP N TERMINAL PROTEIN AMINO ACID MODIFICATION
                                                                                                KEGG_BASAL_CELL_CARCINOMA, KEGG_BASAL_CELL_CARCINOMA
                                                                                                GOMF_AMINOPEPTIDASE_ACTIVITY, GOMF_AMINOPEPTIDASE_ACTIVITY
                                                                                                LIU_SMARCA4_TARGETS, LIU_SMARCA4_TARGETS
                                                                                                GOBP_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II, GOBP_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II
                                                                                                MIR2909, MIR2909
                                                                                                MIR3169, MIR3169
                                                                                                MIR4776_5P, MIR4776_5P
                                                                                                GOBP_PEPTIDYL_METHIONINE_MODIFICATION, GOBP_PEPTIDYL_METHIONINE_MODIFICATION
                                                                                                GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_1H_ACT_CD4_TCELL_DN, GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_1H_ACT_CD4_TCELL_DN
                                                                                                GOMF_METALLOEXOPEPTIDASE_ACTIVITY, GOMF_METALLOEXOPEPTIDASE_ACTIVITY
                                                                                                GOCC_MULTIVESICULAR_BODY, GOCC_MULTIVESICULAR_BODY
_VS_NAIVE_CD8_TCELL_IL7_IL4_DN, GSE32423_MEMORY_VS_NAIVE_CD8_TCELL_IL7_IL4_DN
                                                                                                GOCC_TRANSCRIPTIONALLY_ACTIVE_CHROMATIN, GOCC_TRANSCRIPTIONALLY_ACTIVE_CHROMATIN
                                                                                                GOBP_POSITIVE_REGULATION_OF_VIRAL_RELEASE_FROM_HOST_CELL, GOBP_POSITIVE_REGULATION_OF_VIRAL_RELEASE_FROM_HOST_CELL
                                                                                                GOBP_NUCLEOSOME_POSITIONING, GOBP_NUCLEOSOME_POSITIONING
                                                                                                HP_PEDAL_EDEMA, HP_PEDAL_EDEMA
                                                                                                chr7q31, chr7q31
                                                                                                WP_REGULATION_OF_WNTBCATENIN_SIGNALING_BY_SMALL_MOLECULE_COMPOUNDS, WP_REGULATION_OF_WNTBCATENIN_SIGNALING_BY_SMALL_MOLECULE_COMPOUNDS
                                                                                                HP_URETEROPELVIC_JUNCTION_OBSTRUCTION, HP_URETEROPELVIC_JUNCTION_OBSTRUCTION
                                                                                                HP_ABNORMAL_ARM_SPAN, HP_ABNORMAL_ARM_SPAN
                                                                                                HP URETERAL STENOSIS, HP URETERAL STENOSIS
                                                                                                CTIP_DN.V1_DN, CTIP_DN.V1_DN
                                                                                                GOCC POSTSYNAPTIC CYTOSOL, GOCC POSTSYNAPTIC CYTOSOL
                                                                                                HP_URETHRAL_VALVE, HP_URETHRAL_VALVE
                                                                                                HP_ABNORMALITY_OF_THE_2ND_METACARPAL, HP_ABNORMALITY_OF_THE_2ND_METACARPAL
                                                                                                GOBP NEGATIVE REGULATION OF MYOBLAST DIFFERENTIATION, GOBP NEGATIVE REGULATION OF MYOBLAST DIFFERENTIATION
                                                                                                GOCC MANCHETTE, GOCC MANCHETTE
                                                                                                GOBP NEGATIVE REGULATION OF CARDIOCYTE DIFFERENTIATION, GOBP NEGATIVE REGULATION OF CARDIOCYTE DIFFERENTIATION
                                                                                                GOMF_CO_RECEPTOR_BINDING, GOMF_CO_RECEPTOR_BINDING
                                                                                                HP_APLASIA_HYPOPLASIA_OF_THE_2ND_METACARPAL, HP_APLASIA_HYPOPLASIA_OF_THE_2ND_METACARPAL
                                                                                                GOBP_REGULATION_OF_TRANSCRIPTION_BY_GLUCOSE, GOBP_REGULATION_OF_TRANSCRIPTION_BY_GLUCOSE
                                                                                                GOBP_ADENOHYPOPHYSIS_DEVELOPMENT, GOBP_ADENOHYPOPHYSIS_DEVELOPMENT
                                                                                                GOBP_TRANSMEMBRANE_RECEPTOR_PROTEIN_TYROSINE_PHOSPHATASE_SIGNALING_PATHWAY, GOBP_TRANSMEMBRANE_RECEPTOR_PROTEIN_TYROSINE_PHOSPHATASE_SIGNALING_PATHWAY
                                                                                               GOBP_PHOTOTRANSDUCTION_VISIBLE_LIGHT, GOBP_PHOTOTRANSDUCTION_VISIBLE_LIGHT
                                                                                               GOBP_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION
                                                                                               GOBP_NEGATIVE_REGULATION_OF_INTERLEUKIN_13_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_INTERLEUKIN_13_PRODUCTION
                                                                                               NADELLA_PRKAR1A_TARGETS_DN, NADELLA_PRKAR1A_TARGETS_DN
                                                                                               GOBP_POSITIVE_REGULATION_OF_PHOSPHOLIPASE_A2_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PHOSPHOLIPASE_A2_ACTIVITY
                                                                                               GOMF_ESTROGEN_RECEPTOR_ACTIVITY, GOMF_ESTROGEN_RECEPTOR_ACTIVITY
                                                                                               GOBP_TANGENTIAL_MIGRATION_FROM_THE_SUBVENTRICULAR_ZONE_TO_THE_OLFACTORY_BULB, GOBP_TANGENTIAL_MIGRATION_FROM_THE_SUBVENTRICULAR_ZONE_TO_THE_OLFACTORY_BULB
                                                                                               GOBP_APOPTOTIC_PROCESS_INVOLVED_IN_HEART_MORPHOGENESIS, GOBP_APOPTOTIC_PROCESS_INVOLVED_IN_HEART_MORPHOGENESIS
                                                                                               GOBP_APOPTOTIC_PROCESS_IN_BONE_MARROW_CELL, GOBP_APOPTOTIC_PROCESS_IN_BONE_MARROW_CELL
                                                                                               GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_HEART_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVO
                                                                                               GOBP_NEGATIVE_REGULATION_OF_APOPTOTIC_PROCESS_INVOLVED_IN_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_APOPTOTIC_PROCESS_INVOLVED_IN_DEVELOPMENT
                                                                                               GOBP_PRIMITIVE_STREAK_FORMATION, GOBP_PRIMITIVE_STREAK_FORMATION
                                                                                               GOBP_NEGATIVE_REGULATION_OF_WNT_SIGNALING_PATHWAY_INVOLVED_IN_HEART_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_WNT_SIGNALING_PATHWAY_INVOLVED_IN_HEART_DEVELOPMENT
                                                                                                GOBP_REGULATION_OF_HEART_MORPHOGENESIS, GOBP_REGULATION_OF_HEART_MORPHOGENESIS
                                                                                               chr1p12, chr1p12
                                                                                                chr5q15, chr5q15
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GSE18893\_CTRL\_VS\_TNF\_TREATED\_TREG\_2H\_UP, GSE18893\_CTRL\_VS\_TNF\_TREATED\_TREG\_2H\_UP

GSE32423 MEMORY\_VS\_NAIVE\_CD8\_TCELL\_DN, GSE32423 MEMORY\_VS\_NAIVE\_CD8\_TCELL\_DN

MIR1207\_3P, MIR1207\_3P

GSE32423\_IL7\_VS\_IL7\_IL4\_NAIVE\_CD8\_TCELL\_DN, GSE32423\_IL7\_VS\_IL7\_IL4\_NAIVE\_CD8\_TCELL\_DN