	GSE10239_MEMORY_VS_KLRG1HIGH_EFF_CD8_TCELL_DN, GSE10239_MEMORY_VS_KLRG1HIGH_EFF_CD8_TCELL_DN GSE37532_WT_VS_PPARG_KO_LN_TCONV_DN, GSE37532_WT_VS_PPARG_KO_LN_TCONV_DN GSE40273_XBP1_KO_VS_WT_TREG_UP, GSE40273_XBP1_KO_VS_WT_TREG_UP GSE17721_POLVIC_VS_PAM3CSK4_1H_BMDC_UP, GSE17721_POLYIC_VS_PAM3CSK4_1H_BMDC_UP GSE19772_HCMV_INFL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_DN, GSE19772_HCMV_INFL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_DN GSE43863_TH1_VS_LY6C_LOW_CXCR5NEG_EFFECTOR_CD4_TCELL_DN, GSE43863_TH1_VS_LY6C_LOW_CXCR5NEG_EFFECTOR_CD4_TCELL_DN GSE39152_BRAIN_VS_SPLEEN_CD103_NEG_MEMORY_CD8_TCELL_UP, GSE39152_BRAIN_VS_SPLEEN_CD103_NEG_MEMORY_CD8_TCELL_DN GSE3686_ID3_TRANSDUCED_VS_ID3_KO_CD8_TCELL_DN, GSE23568_ID3_TRANSDUCED_VS_ID3_KO_CD8_TCELL_DN GSE36476_CTRL_VS_TSST_ACT_72H_MEMORY_CD4_TCELL_YOUNG_UP, GSE36476_CTRL_VS_TSST_ACT_72H_MEMORY_CD4_TCELL_YOUNG_UP
	GSE42021_TREG_PLN_VS_CD24INT_TREG_THYMUS_UP, GSE42021_TREG_PLN_VS_CD24INT_TREG_THYMUS_UP GSE15930_STIM_VS_STIM_AND_IL12_72H_CD8_T_CELL_UP, GSE15930_STIM_VS_STIM_AND_IL12_72H_CD8_T_CELL_UP GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_IL12_CD8_TCELL_DN, GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_IL12_CD8_TCELL_DN GSE3982_NEUTROPHIL_VS_NKCELL_UP, GSE3982_NEUTROPHIL_VS_NKCELL_UP GSE4535_BM_DERIVED_DC_VS_FOLLICULAR_DC_UP, GSE4535_BM_DERIVED_DC_VS_FOLLICULAR_DC_UP GSE411_100MIN_VS_400MIN_IL6_STIM_SOCS3_KO_MACROPHAGE_UP, GSE411_100MIN_VS_400MIN_IL6_STIM_SOCS3_KO_MACROPHAGE_UP GSE39820_CTRL_VS_TGFBETA1_IL6_CD4_TCELL_DN, GSE39820_CTRL_VS_TGFBETA1_IL6_CD4_TCELL_DN GSE11057_NAIVE_VS_MEMORY_CD4_TCELL_DN, GSE11057_NAIVE_VS_MEMORY_CD4_TCELL_DN JIANG_HYPOXIA_NORMAL, JIANG_HYPOXIA_NORMAL
	GSE30083_SP2_VS_SP4_THYMOCYTE_DN, GSE30083_SP2_VS_SP4_THYMOCYTE_DN GSE17721_CTRL_VS_CPG_0.5H_BMDC_UP, GSE17721_CTRL_VS_CPG_0.5H_BMDC_UP GSE27786_CD4_TCELL_VS_NKCELL_DN, GSE27786_CD4_TCELL_VS_NKCELL_DN GSE15330_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_UP, GSE15330_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_UP GSE15310_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_UP, GSE15330_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_UP GSE1721_0.5H_VS_24H_PAM3CSK4_BMDC_DN, GSE17721_0.5H_VS_24H_PAM3CSK4_BMDC_DN GSE40068_CXCR5NEG_BCL6NEG_CD4_TCELL_VS_CXCR5POS_BCL6NEG_TFH_DN, GSE40068_CXCR5NEG_BCL6NEG_CD4_TCELL_VS_CXCR5POS_BCL6NEG_TFH_DN GOLDRATH_NAIVE_VS_MEMORY_CD8_TCELL_DN, GOLDRATH_NAIVE_VS_MEMORY_CD8_TCELL_DN GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_DN, GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_DN GSE411_UNSTIM_VS_100MIN_IL6_STIM_SOCS3_KO_MACROPHAGE_DN, GSE411_UNSTIM_VS_100MIN_IL6_STIM_SOCS3_KO_MACROPHAGE_DN MODULE_239, MODULE_239 GSE33425_CD161_HIGH_VS_INT_CD8_TCELL_UP, GSE33425_CD161_HIGH_VS_INT_CD8_TCELL_UP
	GSE33425_CD161_HIGH_VS_INT_CD8_TCELL_UP, GSE33425_CD161_HIGH_VS_INT_CD8_TCELL_UP GSE30083_SP3_VS_SP4_THYMOCYTE_DN, GSE30083_SP3_VS_SP4_THYMOCYTE_DN GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCYTE_UP, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCYTE_UP GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_DN, GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_DN GSE42021_TCONV_PLN_VS_TREG_PRECURSORS_THYMUS_DN, GSE42021_TCONV_PLN_VS_TREG_PRECURSORS_THYMUS_DN GSE24634_TREG_VS_TCONV_POST_DAY3_IL4_CONVERSION_DN, GSE24634_TREG_VS_TCONV_POST_DAY3_IL4_CONVERSION_DN GSE33425_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_DN, GSE33425_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_DN GSE3395_TGFB_IL6_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_DN, GSE3365_CTRL_VS_LPS_INJECTED_DUSP1_KO_SPLENOCYTES_UP GSE34142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCYTE_FETAL_UP, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCYTE_FETAL_UP
	GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN SCHERER_PBMC_APSV_WETVAX_AGE_18_32YO_50_TO_60DY_UP, SCHERER_PBMC_APSV_WETVAX_AGE_18_32YO_50_TO_60DY_UP GSE43956_WT_VS_SGK1_KO_TH17_DIFFERENTIATED_CD4_TCELL_DN, GSE43956_WT_VS_SGK1_KO_TH17_DIFFERENTIATED_CD4_TCELL_DN GSE16450_IMMATURE_VS_MATURE_NEURON_CELL_LINE_12H_IFNA_STIM_DN, GSE16450_IMMATURE_VS_MATURE_NEURON_CELL_LINE_12H_IFNA_STIM_DN GSE29164_UNTREATED_VS_CD8_TCELL_TREATED_MELANOMA_DAY7_UP, GSE29164_UNTREATED_VS_CD8_TCELL_TREATED_MELANOMA_DAY7_UP GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_UP, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_UP GSE17186_MEMORY_VS_CD21HIGH_TRANSITIONAL_BCELL_DN, GSE17186_MEMORY_VS_CD21HIGH_TRANSITIONAL_BCELL_DN GSE40666_NAIVE_VS_EFFECTOR_CD8_TCELL_WITH_IFNA_STIM_90MIN_UP, GSE40666_NAIVE_VS_EFFECTOR_CD8_TCELL_WITH_IFNA_STIM_90MIN_UP
	GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_UP, GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_UP GSE38304_MYC_NEG_VS_POS_GC_BCELL_DN, GSE38304_MYC_NEG_VS_POS_GC_BCELL_DN GSE7852_TREG_VS_TCONV_UP, GSE7852_TREG_VS_TCONV_UP GSE19772_CTRL_VS_HCMV_INF_MONOCYTES_UP, GSE19772_CTRL_VS_HCMV_INF_MONOCYTES_UP GSE11057_NAIVE_VS_EFF_MEMORY_CD4_TCELL_DN, GSE11057_NAIVE_VS_EFF_MEMORY_CD4_TCELL_DN GSE17721_PAM3CSK4_VS_GADIQUIMOD_12H_BMDC_DN, GSE117721_PAM3CSK4_VS_GADIQUIMOD_12H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_12H_BMDC_DN, KAECH_DAY8_EFF_VS_MEMORY_CD8_TCELL_DN, KAECH_DAY8_EFF_VS_MEMORY_CD8_TCELL_DN, KAECH_DAY8_EFF_VS_MEMORY_CD8_TCELL_DN, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_FETAL_UP GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_FETAL_UP, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_FETAL_UP GSE19702_POLYIC_VS_PAM3CSK4_16H_BMDC_UP, GSE17721_POLYIC_VS_PAM3CSK4_16H_BMDC_UP GSE17721_POLYIC_VS_PAM3CSK4_16H_BMDC_UP, GSE17721_POLYIC_VS_PAM3CSK4_16H_BMDC_UP
	GSE37301_MULTIPOTENT_PROGENITOR_VS_RAG2_KO_NK_CELL_DN, GSE37301_MULTIPOTENT_PROGENITOR_VS_RAG2_KO_NK_CELL_DN GSE40274_HELIOS_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_HELIOS_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_DN, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_DN GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_MATURE_NEURON_CELL_LINE_DN, GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_MATURE_NEURON_CELL_LINE_DN GSE46143_CTRL_VS_LMP2A_TRANSDUCED_CD10_POS_GC_BCELL_DN, GSE46143_CTRL_VS_LMP2A_TRANSDUCED_CD10_POS_GC_BCELL_DN GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_DN, GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_DN GSE2935_UV_INACTIVATED_VS_LIVE_SENDAI_VIRUS_INF_MACROPHAGE_DN, GSE2935_UV_INACTIVATED_VS_LIVE_SENDAI_VIRUS_INF_MACROPHAGE_DN GSE9650_NAIVE_VS_MEMORY_CD8_TCELL_DN, GSE9650_NAIVE_VS_MEMORY_CD8_TCELL_DN GSE32034_UNTREATED_VS_ROSIGLIZATONE_TREATED_LY6C_HIGH_MONOCYTE_DN, GSE32034_UNTREATED_VS_ROSIGLIZATONE_TREATED_LY6C_HIGH_MONOCYTE_DN
	GSE7852_TREG_VS_TCONV_LN_UP, GSE7852_TREG_VS_TCONV_LN_UP GSE20366_CD103_KLRG1_DP_VS_DN_TREG_DN, GSE20366_CD103_KLRG1_DP_VS_DN_TREG_DN GSE13946_CTRL_VS_DSS_COLITIS_GD_TCELL_FROM_COLON_UP, GSE13946_CTRL_VS_DSS_COLITIS_GD_TCELL_FROM_COLON_UP GSE25087_TREG_VS_TCONV_FETUS_UP, GSE25087_TREG_VS_TCONV_FETUS_UP KAECH_NAIVE_VS_DAY15_EFF_CD8_TCELL_DN, KAECH_NAIVE_VS_DAY15_EFF_CD8_TCELL_DN GSE7852_LN_VS_FAT_TCONV_DN, GSE7852_LN_VS_FAT_TCONV_DN GSE7852_TREG_VS_TCONV_FAT_UP, GSE7852_TREG_VS_TCONV_FAT_UP GSE25087_TREG_VS_TCONV_FAT_UP, GSE25087_TREG_VS_TCONV_ADULT_UP GSE25087_TREG_VS_TCONV_ADULT_UP, GSE25087_TREG_VS_TCONV_ADULT_UP GSE33425_CD8_ALPHAALPHA_VS_ALPHABETA_CD16_HIGH_TCELL_UP, GSE33425_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_UP
	GSE6259_DEC205_POS_DC_VS_CD8_TCELL_UP, GSE6259_DEC205_POS_DC_VS_CD8_TCELL_UP GSE17721_0.5H_VS_12H_LPS_BMDC_DN, GSE17721_0.5H_VS_12H_LPS_BMDC_DN GSE3982_MAST_CELL_VS_TH2_UP, GSE3982_MAST_CELL_VS_TH2_UP GSE20366_TREG_VS_NAIVE_CD4_TCELL_HOMEOSTATIC_CONVERSION_DN, GSE20366_TREG_VS_NAIVE_CD4_TCELL_HOMEOSTATIC_CONVERSION_DN GSE27434_WT_VS_DNMT1_KO_TREG_DN, GSE27434_WT_VS_DNMT1_KO_TREG_DN GSE39916_B_CELL_SPLEEN_VS_PLASMA_CELL_BONE_MARROW_DN, GSE39916_B_CELL_SPLEEN_VS_PLASMA_CELL_BONE_MARROW_DN GSE7852_LN_VS_THYMUS_TCONV_UP, GSE7852_LN_VS_THYMUS_TCONV_UP GSE339_EX_VIVO_VS_IN_CULTURE_CD4POS_DC_DN, GSE339_EX_VIVO_VS_IN_CULTURE_CD4POS_DC_DN GSE39955_TH0_VS_TGFB_L6_TH17_ACT_CD4_TCELL_42H_UP, GSE43955_TH0_VS_TGFB_L6_TH17_ACT_CD4_TCELL_42H_UP GSE19923_HEB_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_DN, GSE19923_HEB_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_DN
	CUI_DEVELOPING_HEART_C9_B_T_CELL, CUI_DEVELOPING_HEART_C9_B_T_CELL  GSE15330_LYMPHOID_MULTIPOTENT_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_IKAROS_KO_DN, GSE15330_LYMPHOID_MULTIPOTENT_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_IKAROS_KO_DN  GSE339_CD4POS_VS_CD4CD8DN_DC_DN, GSE339_CD4POS_VS_CD4CD8DN_DC_DN  GSE11961_FOLLICULAR_BCELL_VS_PLASMA_CELL_DAY7_DN, GSE11961_FOLLICULAR_BCELL_VS_PLASMA_CELL_DAY7_DN  GSE3565_CTRL_VS_LPS_INJECTED_SPLENOCYTES_UP, GSE3565_CTRL_VS_LPS_INJECTED_SPLENOCYTES_UP  GSE3920_UNTREATED_VS_IFNA_TREATED_ENDOTHELIAL_CELL_UP, GSE3920_UNTREATED_VS_IFNA_TREATED_ENDOTHELIAL_CELL_UP  GSE13306_RA_VS_UNTREATED_MEM_CD4_TCELL_UP, GSE13306_RA_VS_UNTREATED_MEM_CD4_TCELL_UP  GSE3982_MAST_CELL_VS_TH1_UP, GSE3982_MAST_CELL_VS_TH1_UP  GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_DN, GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_DN
	GSE17721_0.5H_VS_24H_LPS_BMDC_DN, GSE17721_0.5H_VS_24H_LPS_BMDC_DN GSE37301_HEMATOPOIETIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_DN, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_DN GSE33374_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_DN, GSE33374_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_DN GSE17721_CTRL_VS_GARDIQUIMOD_6H_BMDC_DN, GSE17721_CTRL_VS_GARDIQUIMOD_6H_BMDC_DN GSE339_EX_VIVO_VS_IN_CULTURE_CD4CD8DN_DC_DN, GSE339_EX_VIVO_VS_IN_CULTURE_CD4CD8DN_DC_DN GCATTTG_MIR105, GCATTTG_MIR105 GSE1448_CTRL_VS_ANTI_VBETA5_DP_THYMOCYTE_DN, GSE1448_CTRL_VS_ANTI_VBETA5_DP_THYMOCYTE_DN GSE1386_NAIVE_VS_MEMORY_BCELL_DN, GSE11386_NAIVE_VS_MEMORY_BCELL_DN GSE1386_NAIVE_VS_MEMORY_BCELL_DP, GSE24142_DN2_VS_DN3_THYMOCYTE_UP GSE32255_WT_UNSTIM_VS_IMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_DN, GSE32255_WT_UNSTIM_VS_IMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_DN
	GSE2405_HEAT_KILLED_LYSATE_VS_LIVE_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_24H_UP, GSE2405_HEAT_KILLED_LYSATE_VS_LIVE_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_24H_UP RUBENSTEIN_SKELETAL_MUSCLE_FAP_CELLS, RUBENSTEIN_SKELETAL_MUSCLE_FAP_CELLS GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_DN, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_DN TRAVAGLINI_LUNG_MACROPHAGE_CELL, TRAVAGLINI_LUNG_MACROPHAGE_CELL GSE42021_CD24HI_TREG_VS_CD24HI_TCONV_THYMUS_DN, GSE42021_CD24HI_TREG_VS_CD24HI_TCONV_THYMUS_DN GSE25123_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_UP, GSE25123_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_UP GSE24142_ADULT_VS_FETAL_DN2_THYMOCYTE_UP, GSE24142_ADULT_VS_FETAL_DN2_THYMOCYTE_UP GSE7460_WT_VS_FOXP3_HET_ACT_WITH_TGFB_TCONV_DN, GSE7460_WT_VS_FOXP3_HET_ACT_WITH_TGFB_TCONV_DN GSE37301_MULTIPOTENT_PROGENITOR_VS_GRAN_MONO_PROGENITOR_UP, GSE37301_MULTIPOTENT_PROGENITOR_UP
	KAECH_NAIVE_VS_MEMORY_CD8_TCELL_DN, KAECH_NAIVE_VS_MEMORY_CD8_TCELL_DN  GSE26488_WT_VS_HDAC7_DELTAP_TG_OT2_THYMOCYTE_WITH_PEPTIDE_INJECTION_DN, GSE26488_WT_VS_HDAC7_DELTAP_TG_OT2_THYMOCYTE_WITH_PEPTIDE_INJECTION_DN  GSE13522_WT_VS_IFNAR_KO_SKING_T_CRUZI_Y_STRAIN_INF_UP, GSE13522_WT_VS_IFNAR_KO_SKING_T_CRUZI_Y_STRAIN_INF_UP  GSE7852_TREG_VS_TCONV_THYMUS_UP, GSE7852_TREG_VS_TCONV_THYMUS_UP  GSE13522_CTRL_VS_T_CRUZI_G_STRAIN_INF_SKIN_UP, GSE13522_CTRL_VS_T_CRUZI_G_STRAIN_INF_SKIN_UP  GSE27896_HDAC6_KO_VS_WT_TREG_DN, GSE27896_HDAC6_KO_VS_WT_TREG_DN  GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCYTE_DN, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCYTE_DN  GSE17721_CPG_VS_GARDIQUIMOD_24H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_24H_BMDC_DN  GROSS_HYPOXIA_VIA_ELK3_AND_HIF1A_UP, GROSS_HYPOXIA_VIA_ELK3_AND_HIF1A_UP, GGOSS_HYPOXIA_VIA_ELK3_AND_HIF1A_UP, GGOSS_HYP
	GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_UP, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_UP GSE13522_WT_VS_IFNAR_KO_SKIN_DN, GSE13522_WT_VS_IFNAR_KO_SKIN_DN GSE11961_PLASMA_CELL_DAYY_VS_MEMORY_BCELL_DAY40_UP, GSE11961_PLASMA_CELL_DAY7_VS_MEMORY_BCELL_DAY40_UP GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_UP, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_UP GSE339_EX_VIVO_VS_IN_CULTURE_CD8POS_DC_DN, GSE339_EX_VIVO_VS_IN_CULTURE_CD8POS_DC_DN GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_UP, GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_UP CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CDC25_DN, CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CDC25_DN GSE7219_UNSTIM_VS_LPS_AND_ANTI_CD40_STIM_DC_DN, GSE7219_UNSTIM_VS_LPS_AND_ANTI_CD40_STIM_DC_DN, GSE7213_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_DN, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_DN, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_DN, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_DN, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_DN, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_DN GSE2565_DUSP1_VS_WT_SPLENOCYTES_UP, GSE3565_DUSP1_VS_WT_SPLENOCYTES_UP
	MIR22_3P, MIR22_3P MIR6071, MIR6071 CHIARADONNA_NEOPLASTIC_TRANSFORMATION_KRAS_DN, CHIARADONNA_NEOPLASTIC_TRANSFORMATION_KRAS_DN HP_ABNORMALITY_OF_THE_LYMPH_NODES, HP_ABNORMALITY_OF_THE_LYMPH_NODES GSE21360_TERTIARY_VS_QUATERNARY_MEMORY_CD8_TCELL_DN, GSE21360_TERTIARY_VS_QUATERNARY_MEMORY_CD8_TCELL_DN GSE26030_UNSTIM_VS_RESTIM_TH17_DAY5_POST_POLARIZATION_UP, GSE26030_UNSTIM_VS_RESTIM_TH17_DAY5_POST_POLARIZATION_UP GAVIN_FOXP3_TARGETS_CLUSTER_P3, GAVIN_FOXP3_TARGETS_CLUSTER_P3 LANDIS_ERBB2_BREAST_TUMORS_324_UP, LANDIS_ERBB2_BREAST_TUMORS_324_UP ZNF133_TARGET_GENES, ZNF133_TARGET_GENES
	GSE4590_LARGE_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_UP, GSE4590_LARGE_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_UP GSE9650_NAIVE_VS_EXHAUSTED_CD8_TCELL_DN, GSE9650_NAIVE_VS_EXHAUSTED_CD8_TCELL_DN FAN_EMBRYONIC_CTX_BRAIN_EFFECTOR_T_CELL, FAN_EMBRYONIC_CTX_BRAIN_EFFECTOR_T_CELL GSE14415_NATURAL_TREG_VS_TCONV_UP, GSE14415_NATURAL_TREG_VS_TCONV_UP MIR200B_5P, MIR200B_5P GSE22886_NAIVE_CD8_TCELL_VS_NEUTROPHIL_DN, GSE22886_NAIVE_CD8_TCELL_VS_NEUTROPHIL_DN GSE17580_UNINFECTED_VS_S_MANSONI_INF_TEFF_DN, GSE17580_UNINFECTED_VS_S_MANSONI_INF_TEFF_DN GSE7460_CD8_TCELL_VS_CD4_TCELL_ACT_UP, GSE7460_CD8_TCELL_VS_CD4_TCELL_ACT_UP MIR200A_5P, MIR200A_5P MODULE_491, MODULE_491
	GSE30962_PRIMARY_VS_SECONDARY_CHRONIC_LCMV_INF_CD8_TCELL_DN, GSE30962_PRIMARY_VS_SECONDARY_CHRONIC_LCMV_INF_CD8_TCELL_DN SCHAEFFER_PROSTATE_DEVELOPMENT_6HR_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_6HR_UP GSE21033_CTRL_VS_POLYIC_STIM_DC_24H_UP, GSE21033_CTRL_VS_POLYIC_STIM_DC_24H_UP GSE7460_CTRL_VS_TGFB_TREATED_ACT_CD8_TCELL_DN, GSE7460_CTRL_VS_TGFB_TREATED_ACT_CD8_TCELL_DN IGLESIAS_E2F_TARGETS_UP, IGLESIAS_E2F_TARGETS_UP GSE37301_RAG2_KO_VS_RAG2_AND_ETS1_KO_NK_CELL_UP, GSE37301_RAG2_KO_VS_RAG2_AND_ETS1_KO_NK_CELL_UP GSE4590_SMALL_VS_LARGE_PRE_BCELL_DN, GSE4590_SMALL_VS_LARGE_PRE_BCELL_DN TRAVAGLINI_LUNG_NATURAL_KILLER_CELL, TRAVAGLINI_LUNG_NATURAL_KILLER_CELL GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_1_DN, GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_1_DN
	GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN, GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN BURTON_ADIPOGENESIS_9, BURTON_ADIPOGENESIS_9 MODULE_568, MODULE_568 GSE17721_CTRL_VS_CPG_4H_BMDC_DN, GSE17721_CTRL_VS_CPG_4H_BMDC_DN ICHIBA_GRAFT_VERSUS_HOST_DISEASE_35D_UP, ICHIBA_GRAFT_VERSUS_HOST_DISEASE_35D_UP GSE41176_UNSTIM_VS_ANTI_IGM_STIM_TAK1_KO_BCELL_24H_DN, GSE41176_UNSTIM_VS_ANTI_IGM_STIM_TAK1_KO_BCELL_24H_DN WAMUNYOKOLL_OVARIAN_CANCER_GRADES_1_2_UP, WAMUNYOKOLL_OVARIAN_CANCER_GRADES_1_2_UP ZHONG_PFC_C4_PTGDS_POS_OPC, ZHONG_PFC_C4_PTGDS_POS_OPC GNP2_MCL1, GNP2_MCL1
	GSEI7721_LPS_VS_POLYIC_2H_BMDC_UP, GSE17721_LPS_VS_POLYIC_2H_BMDC_UP CROMER_METASTASIS_UP, CROMER_METASTASIS_UP ROESSLER_LIVER_CANCER_METASTASIS_UP, ROESSLER_LIVER_CANCER_METASTASIS_UP GOCC_NEURON_SPINE, GOCC_NEURON_SPINE WIELAND_UP_BY_HBV_INFECTION, WIELAND_UP_BY_HBV_INFECTION GSE5589_LPS_AND_IL10_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_UP, GSE5589_LPS_AND_IL10_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_UP GSE43955_1H_VS_60H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN, GSE43955_1H_VS_60H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN MENON_FETAL_KIDNEY_6_COLLECTING_DUCT_CELLS, MENON_FETAL_KIDNEY_6_COLLECTING_DUCT_CELLS, MENON_FETAL_KIDNEY_6_COLLECTING_DUCT_CELS, MENON_FETAL_KIDNEY_F
	GOCC_RECYCLING_ENDOSOME_MEMBRANE, GOCC_RECYCLING_ENDOSOME_MEMBRANE TAVOR_CEBPA_TARGETS_UP, TAVOR_CEBPA_TARGETS_UP GSE26488_WT_VS_HDAC7_DELTAP_TG_OT2_THYMOCYTE_WITH_PEPTIDE_INJECTION_UP, GSE26488_WT_VS_HDAC7_DELTAP_TG_OT2_THYMOCYTE_WITH_PEPTIDE_INJECTION_UP GSE4590_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_UP, GSE4590_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_UP GSE21927_EL4_VS_MCA203_TUMOR_MONOCYTES_DN, GSE21927_EL4_VS_MCA203_TUMOR_MONOCYTES_DN GSE5099_CLASSICAL_M1_VS_ALTERNATIVE_M2_MACROPHAGE_DN, GSE5099_CLASSICAL_M1_VS_ALTERNATIVE_M2_MACROPHAGE_DN GSE339_CD4POS_VS_CD4CD8DN_DC_IN_CULTURE_UP, GSE339_CD4POS_VS_CD4CD8DN_DC_IN_CULTURE_UP BROWNE_HCMV_INFECTION_10HR_UP, BROWNE_HCMV_INFECTION_10HR_UP P53_DN.V1_DN, P53_DN.V1_DN
	GSE37301_PRO_BCELL_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_UP, GSE37301_PRO_BCELL_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_UP BOYLAN_MULTIPLE_MYELOMA_PCA1_UP, BOYLAN_MULTIPLE_MYELOMA_PCA1_UP MARKEY_RB1_CHRONIC_LOF_DN, MARKEY_RB1_CHRONIC_LOF_DN MIR1206, MIR1206 HP_HEPATOSPLENOMEGALY, HP_HEPATOSPLENOMEGALY YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_9, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_9 LE_EGR2_TARGETS_DN, LE_EGR2_TARGETS_DN GSE17721_LPS_VS_CPG_0.5H_BMDC_DN, GSE17721_LPS_VS_CPG_0.5H_BMDC_DN GSE17721_LPS_VS_CPG_0.5H_BMDC_DN, GSE17721_LPS_VS_CPG_0.5H_BMDC_DN GOBP_CORTICAL_CYTOSKELETON_ORGANIZATION, GOBP_CORTICAL_CYTOSKELETON_ORGANIZATION MIR3689A_5P_MIR3689B_5P_MIR3689E_MIR3689F, MIR3689B_5P_M
	GAJATE_RESPONSE_TO_TRABECTEDIN_UP, GAJATE_RESPONSE_TO_TRABECTEDIN_UP MEL18_DN.V1_DN, MEL18_DN.V1_DN GSE9650_EXHAUSTED_VS_MEMORY_CD8_TCELL_UP, GSE9650_EXHAUSTED_VS_MEMORY_CD8_TCELL_UP MIR21_5P, MIR21_5P GAVIN_FOXP3_TARGETS_CLUSTER_P2, GAVIN_FOXP3_TARGETS_CLUSTER_P2 MIR590_5P, MIR590_5P MIR3116, MIR3116 GOBP_CYTOKINETIC_PROCESS, GOBP_CYTOKINETIC_PROCESS GOBP_ODONTOGENESIS, GOBP_ODONTOGENESIS
	GOBP_PRESYNAPTIC_ENDOCYTOSIS, GOBP_PRESYNAPTIC_ENDOCYTOSIS GSE15733_BM_VS_SPLEEN_MEMORY_CD4_TCELL_UP, GSE15733_BM_VS_SPLEEN_MEMORY_CD4_TCELL_UP AMIT_EGF_RESPONSE_240_HELA, AMIT_EGF_RESPONSE_240_HELA GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_240_HELA GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_240_HELA GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_240_HELA GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_240_HELA GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_ACTIVITY, GMEMORY_CD4_FCB_ACTIVITY GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_ACTIVITY, GMEMORY_CD4_FCB_ACTIVITY GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_ACTIVITY, GMEMORY_CD4_FCB_ACTIVITY GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_FCB_ACTIVITY, GMEMORY_CD4_FCB_ACTIVITY GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_FCB_ACTIVITY, GMEMORY_CD4_FCB_ACTIVITY GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_FCB_ACTIVITY GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_FCB_ACTIVITY GMEMORY_CD4_FCB_AMIT_EGF_RESPONSE_FCB_ACTIVITY GMEMORY_CD4_FCB_ACTIVITY_CD4_FCB_ACTIVITY GMEMORY_CD4_FCB_ACTIVITY_CD4_FCB_ACTIVITY_CD4_FCB_ACTIVITY GMEMORY_CD4_FCB_ACTIVITY_C
	VALK_AMI_CLUSTER_15, VALK_AMI_CLUSTER_15 MIR6874_3P, MIR6874_3P ZHENG_CORD_BLOOD_C1_PUTATIVE_MEGAKARYOCYTE_PROGENITOR, ZHENG_CORD_BLOOD_C1_PUTATIVE_MEGAKARYOCYTE_PROGENITOR MIR4473, MIR4473 ZHONG_PFC_MAJOR_TYPES_OPC, ZHONG_PFC_MAJOR_TYPES_OPC DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_NK_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_NK_CELLS GOBP_NEGATIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS, GOBP_NEGATIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS CRX_NRL_DN.V1_DN, CRX_NRL_DN.V1_DN MIR134_3P, MIR134_3P
	MIR148B_5P, MIR148B_5P MIR8077, MIR8077 GOBP_NEGATIVE_REGULATION_OF_IMMUNE_RESPONSE, GOBP_NEGATIVE_REGULATION_OF_IMMUNE_RESPONSE BCAT_GDS748_UP, BCAT_GDS748_UP ROSS_AML_OF_FAB_M7_TYPE, ROSS_AML_OF_FAB_M7_TYPE RADAEVA_RESPONSE_TO_IFNA1_UP, RADAEVA_RESPONSE_TO_IFNA1_UP MIR4657, MIR4657 HP_INCREASED_CIRCULATING_ANTIBODY_LEVEL, HP_INCREASED_CIRCULATING_ANTIBODY_LEVEL HUMMERICH_SKIN_CANCER_PROGRESSION_UP, HUMMERICH_SKIN_CANCER_PROGRESSION_UP MCCLUNG_COCAINE_REWARD_5D, MCCLUNG_COCAINE_REWARD_5D
	BERENJENO_ROCK_SIGNALING_NOT_VIA_RHOA_DN, BERENJENO_ROCK_SIGNALING_NOT_VIA_RHOA_DN MIR4491, MIR4491 WOOD_EBV_EBNA1_TARGETS_DN, WOOD_EBV_EBNA1_TARGETS_DN MIR3187_3P, MIR3187_3P XIE_LT_HSC_SIPR3_OE_UP, XIE_LT_HSC_SIPR3_OE_UP ELLWOOD_MYC_TARGETS_DN, ELLWOOD_MYC_TARGETS_DN RHEIN_ALL_GLUCOCORTICOID_THERAPY_UP, RHEIN_ALL_GLUCOCORTICOID_THERAPY_UP INGRAM_SHH_TARGETS_UP, INGRAM_SHH_TARGETS_UP HP_ABNORMAL_PULMONARY_THORACIC_IMAGING_FINDING, HP_ABNORMAL_PULMONARY_THORACIC_IMAGING_FINDING SANA_TNF_SIGNALING_DN, SANA_TNF_SIGNALING_DN
	GOBP_REGULATION_OF_LIPASE_ACTIVITY, GOBP_REGULATION_OF_LIPASE_ACTIVITY BUSSLINGER_ESOPHAGEAL_LATE_SUPRABASAL_CELLS, BUSSLINGER_ESOPHAGEAL_LATE_SUPRABASAL_CELLS HOFFMANN_PRE_BI_TO_LARGE_PRE_BII_LYMPHOCYTE_DN, HOFFMANN_PRE_BI_TO_LARGE_PRE_BII_LYMPHOCYTE_DN GOBP_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY GNATENKO_PLATELET_SIGNATURE, GNATENKO_PLATELET_SIGNATURE MIR5089_3P, MIR5089_3P GOBP_RUFFLE_ORGANIZATION, GOBP_RUFFLE_ORGANIZATION GOMF_BETA_TUBULIN_BINDING, GOMF_BETA_TUBULIN_BINDING GOMF_DOONTOGENESIS_OF_DENTIN_CONTAINING_TOOTH, GOBP_ODONTOGENESIS_OF_DENTIN_CONTAINING_TOOTH
61_INT_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP	- HP_AUTOIMMUNE_ANTIBODY_POSITIVITY, HP_AUTOIMMUNE_ANTIBODY_POSITIVITY - GOMF_CELL_ADHESION_MEDIATOR_ACTIVITY, GOMF_CELL_ADHESION_MEDIATOR_ACTIVITY - APPEL_IMATINIB_RESPONSE, APPEL_IMATINIB_RESPONSE - LINDSTEDT_DENDRITIC_CELL_MATURATION_D, LINDSTEDT_DENDRITIC_CELL_MATURATION_D - MIR1288_5P, MIR1288_5P - PID_ARF6_TRAFFICKING_PATHWAY, PID_ARF6_TRAFFICKING_PATHWAY - REACTOME_ACTIVATION_OF_GENE_EXPRESSION_BY_SREBF_SREBP, REACTOME_ACTIVATION_OF_GENE_EXPRESSION_BY_SREBF_SREBP - HOSHIDA_LIVER_CANCER_LATE_RECURRENCE_UP, HOSHIDA_LIVER_CANCER_LATE_RECURRENCE_UP - GOMF_TUMOR_NECROSIS_FACTOR_RECEPTOR_SUPERFAMILY_BINDING, GOMF_TUMOR_NECROSIS_FACTOR_RECEPTOR_SUPERFAMILY_BINDING - REACTOME_TNFS_BIND_THEIR_PHYSIOLOGICAL_RECEPTORS, REACTOME_TNFS_BIND_THEIR_PHYSIOLOGICAL_RECEPTORS
	SALVADOR_MARTIN_PEDIATRIC_TBD_ANTI_TNF_THERAPY_NONRESPONDER_PRE_TREATMENT_UP, SALVADOR_MARTIN_PEDIATRIC_TBD_ANTI_TNF_THERAPY_NONRESPONDER_PRE_TREATMENT_UP GOBP_REGULATION_OF_POSTSYNAPTIC_MEMBRANE_NEUROTRANSMITTER_RECEPTOR_LEVELS, GOBP_REGULATION_OF_POSTSYNAPTIC_MEMBRANE_NEUROTRANSMITTER_RECEPTOR_LEVELS GSE10147_IL3_VS_IL3_AND_CPG_STIM_PDC_UP, GSE10147_IL3_VS_IL3_AND_CPG_STIM_PDC_UP DAVICIONI_RHABDOMYOSARCOMA_PAX_FOXO1_FUSION_UP, DAVICIONI_RHABDOMYOSARCOMA_PAX_FOXO1_FUSION_UP GROSS_ELK3_TARGETS_UP, GROSS_ELK3_TARGETS_UP HINATA_NFKB_TARGETS_LERATINOCYTE_UP, HINATA_NFKB_TARGETS_KERATINOCYTE_UP GOMF_PHOSPHOLIPID_TRANSPORTER_ACTIVITY, GOMF_PHOSPHOLIPID_TRANSPORTER_ACTIVITY GSE27786_LSK_VS_ERYTHROBLAST_DN, GSE27786_LSK_VS_ERYTHROBLAST_DN MCLACHLAN_DENTAL_CARIES_DN, MCLACHLAN_DENTAL_CARIES_DN GOBP_LYMPHOCYTE_APOPTOTIC_PROCESS, GOBP_LYMPHOCYTE_APOPTOTIC_PROCESS
	ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_1DY_UP, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_1DY_UP GOBP_LEUKOCYTE_APOPTOTIC_PROCESS, GOBP_LEUKOCYTE_APOPTOTIC_PROCESS BIOCARTA_HDAC_PATHWAY, BIOCARTA_HDAC_PATHWAY DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_MONOCYTES, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_MONOCYTES GOBP_T_CELL_MEDIATED_CYTOTOXICITY, GOBP_T_CELL_MEDIATED_CYTOTOXICITY LEE_LIVER_CANCER_E2F1_UP, LEE_LIVER_CANCER_E2F1_UP MIR8069, MIR8069 GOBP_POSITIVE_REGULATION_OF_LIPASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_LIPASE_ACTIVITY MODULE_265, MODULE_265 HP_ABNORMAL_MACROPHAGE_MORPHOLOGY, HP_ABNORMAL_MACROPHAGE_MORPHOLOGY
	GOBP_HETEROTYPIC_CELL_CELL_ADHESION, GOBP_HETEROTYPIC_CELL_CELL_ADHESION  GOCC_CELL_CELL_CONTACT_ZONE, GOCC_CELL_CELL_CONTACT_ZONE  GOBP_POSITIVE_REGULATION_OF_LIPID_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_LIPID_TRANSPORT  GOBP_REGULATION_OF_POSTSYNAPSE_ORGANIZATION, GOBP_REGULATION_OF_POSTSYNAPSE_ORGANIZATION  GOMF_CELL_CELL_ADHESION_MEDIATOR_ACTIVITY, GOMF_CELL_CELL_ADHESION_MEDIATOR_ACTIVITY  BORLAK_LIVER_CANCER_EGF_UP, BORLAK_LIVER_CANCER_EGF_UP  HP_POIKILOCYTOSIS, HP_POIKILOCYTOSIS  BOHN_PRIMARY_IMMUNODEFICIENCY_SYNDROM_DN, BOHN_PRIMARY_IMMUNODEFICIENCY_SYNDROM_DN  PLASARI_TGFBI_SIGNALING_VIA_NFIC_10HR_UP, PLASARI_TGFBI_SIGNALING_VIA_NFIC_10HR_UP  CHESLER_BRAIN_HIGHEST_EXPRESSION, CHESLER_BRAIN_HIGHEST_EXPRESSION
	LL_PBMC_MENOMUNE_A_C_Y_W_135_AGE_18_45YO_CORRELATED_WITH_ANTIBODY_RESPONSE_3DY_POSITIVE, LL_PBMC_MENOMUNE_A_C_Y_W_135_AGE_18_45YO_CORRELATED_WITH_ANTIBODY_RESPONSE_3DY_POSITIVE LEIN_NEURON_MARKERS, LEIN_NEURON_MARKERS GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_CYTOTOXICITY, GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_CYTOTOXICITY GOBP_EPITHELIAL_CELL_APOPTOTIC_PROCESS, GOBP_EPITHELIAL_CELL_APOPTOTIC_PROCESS GOBP_LIPID_TRANSLOCATION, GOBP_LIPID_TRANSLOCATION HP_ABNORMAL_CIRCULATING_IGM_LEVEL, HP_ABNORMAL_CIRCULATING_IGM_LEVEL GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_IMMUNITY, GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_IMMUNITY GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_CELL_CELL_ADHESION, GOBP_NEGATIVE_REGULATION_OF_LEUKOCYTE_CELL_CELL_ADHESION GOMF_TUMOR_NECROSIS_FACTOR_RECEPTOR_BINDING, GOMF_TUMOR_NECROSIS_FACTOR_RECEPTOR_BINDING
	GOBP_REGULATION_OF_MEMBRANE_LIPID_DISTRIBUTION, GOBP_REGULATION_OF_MEMBRANE_LIPID_DISTRIBUTION GOBP_NEGATIVE_REGULATION_OF_LYMPHOCYTE_MEDIATED_IMMUNITY, GOBP_NEGATIVE_REGULATION_OF_LYMPHOCYTE_MEDIATED_IMMUNITY GOBP_POSITIVE_REGULATION_OF_PEPTIDE_SECRETION, GOBP_POSITIVE_REGULATION_OF_PEPTIDE_SECRETION GOBP_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_MONOCYTE_DN, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_MONOCYTE_DN GOBP_POSITIVE_REGULATION_OF_LIPID_LOCALIZATION, GOBP_POSITIVE_REGULATION_OF_LIPID_LOCALIZATION GOMF_CORECEPTOR_ACTIVITY, GOMF_CORECEPTOR_ACTIVITY MORI_EMU_MYC_LYMPHOMA_BY_ONSET_TIME_DN, MORI_EMU_MYC_LYMPHOMA_BY_ONSET_TIME_DN GOBP_NEGROTIC_CELL_DEATH, GOBP_NECROTIC_CELL_DEATH
	MOOTHA_GLYCOGEN_METABOLISM, MOOTHA_GLYCOGEN_METABOLISM WOTTON_RUNX_TARGETS_DN, WOTTON_RUNX_TARGETS_DN GOMF_CHEMOKINE_BINDING, GOMF_CHEMOKINE_BINDING GOMF_CHEMOKINE_BINDING, GOMF_CHEMOKINE_BINDING GOBP_MEMBRANE_PROTEIN_PROTEOLYSIS, GOBP_MEMBRANE_PROTEIN_PROTEOLYSIS GOBP_REGULATION_OF_T_CELL_MEDIATED_CYTOTOXICITY, GOBP_REGULATION_OF_T_CELL_MEDIATED_CYTOTOXICITY RORIE_TARGETS_OF_EWSR1_FLI1_FUSION_UP, RORIE_TARGETS_OF_EWSR1_FLI1_FUSION_UP GOBP_TOLL_LIKE_RECEPTOR_3_SIGNALING_PATHWAY, GOBP_TOLL_LIKE_RECEPTOR_3_SIGNALING_PATHWAY WANG_RESPONSE_TO_BEXARCOTENE_DN, WANG_RESPONSE_TO_BEXARCOTENE_DN PID_ECADHERIN_STABILIZATION_PATHWAY, PID_ECADHERIN_STABILIZATION_PATHWAY GNF2_IL2RB, GNF2_IL2RB
	REACTOME_APOPTOTIC_CLEAVAGE_OF_CELLULAR_PROTEINS, REACTOME_APOPTOTIC_CLEAVAGE_OF_CELLULAR_PROTEINS PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_3, PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_3 MIR3657, MIR3657 GOBP_POSITIVE_REGULATION_OF_MONONUCLEAR_CELL_MIGRATION, GOBP_POSITIVE_REGULATION_OF_MONONUCLEAR_CELL_MIGRATION REACTOME_THROMBIN_SIGNALLING_THROUGH_PROTEINASE_ACTIVATED_RECEPTORS_PARS, REACTOME_THROMBIN_SIGNALLING_THROUGH_PROTEINASE_ACTIVATED_RECEPTORS_PARS, GOMF_INSULIN_LIKE_GROWTH_FACTOR_RECEPTOR_BINDING ZHAN_VARIABLE_EARLY_DIFFERENTIATION_GENES_UP, ZHAN_VARIABLE_EARLY_DIFFERENTIATION_GENES_UP, ZHAN_VARIABLE_EARLY_DIFFERENTIATION_GENES_UP GOBP_REGULATION_OF_CHOLESTEROL_METABOLIC_PROCESS, GOBP_REGULATION_OF_CHOLESTEROL_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_ADAPTIVE_IMMUNE_RESPONSE
	MIR7151_5P, MIR7151_5P GOBP_REGULATION_OF_ERAD_PATHWAY, GOBP_REGULATION_OF_ERAD_PATHWAY KORKOLA_YOLK_SAC_TUMOR, KORKOLA_YOLK_SAC_TUMOR BOWIE_RESPONSE_TO_TAMOXIFEN, BOWIE_RESPONSE_TO_TAMOXIFEN KAZMIN_PBMC_P_FALCIPARUM_RTSS_AS01_AGE_UNKNOWN_CORRELATED_WITH_PROTECTION_56DY_NEGATIVE, KAZMIN_PBMC_P_FALCIPARUM_RTSS_AS01_AGE_UNKNOWN_CORRELATED_WITH_PROTECTION_56DY_NEGATIVE HP_OSTEOPETROSIS, HP_OSTEOPETROSIS NAKAYA_PBMC_FLUAD_MALE_AGE_14_27YO_1D_POSTBOOST_VS_0D_PREIMM_MF59_ADJUVANTED_1DY_GENES_IN_BTM_M40_AND_M53_DN, NAKAYA_PBMC_FLUAD_MALE_AGE_14_27YO_1D_POSTBOOST_VS_0D_PREIMM_MF59_ADJUVANTED_1DY_GENES_OPPROGRAMMED_NECROTIC_CELL_DEATH GOBP_PROGRAMMED_NECROTIC_CELL_DEATH, GOBP_PROGRAMMED_NECROTIC_CELL_DEATH GOMF_CADHERIN_BINDING_INVOLVED_IN_CELL_CELL_ADHESION, GOMF_CADHERIN_BINDING_INVOLVED_IN_CELL_CELL_ADHESION KRASNOSELSKAYA_ILF3_TARGETS_UP, KRASNOSELSKAYA_ILF3_TARGETS_UP
	HP_POLYURIA, HP_POLYURIA MODULE_346, MODULE_346 MODULE_408, MODULE_408 REACTOME_DISSOLUTION_OF_FIBRIN_CLOT, REACTOME_DISSOLUTION_OF_FIBRIN_CLOT PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_5, PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_5 GOBP_SMOOTH_MUSCLE_CONTRACTION, GOBP_SMOOTH_MUSCLE_CONTRACTION OHGUCHI_LIVER_HNF4A_TARGETS_DN, OHGUCHI_LIVER_HNF4A_TARGETS_DN TRAVAGLINI_LUNG_CD8_MEMORY_EFFECTOR_T_CELL, TRAVAGLINI_LUNG_CD8_MEMORY_EFFECTOR_T_CELL GAZIN_EPIGENETIC_SILENCING_BY_KRAS, GAZIN_EPIGENETIC_SILENCING_BY_KRAS
	GOCC_FILOPODIUM_MEMBRANE, GOCC_FILOPODIUM_MEMBRANE WP_OSTEOBLAST_SIGNALING, WP_OSTEOBLAST_SIGNALING HP_PULMONARY_HEMORRHAGE GOBP_NEGATIVE_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY KEGG_GLYCOSPHINGOLIPID_BIOSYNTHESIS_LACTO_AND_NEOLACTO_SERIES, KEGG_GLYCOSPHINGOLIPID_BIOSYNTHESIS_LACTO_AND_NEOLACTO_SERIES GOBP_REGULATION_OF_SYNAPTIC_VESICLE_RECYCLING, GOBP_REGULATION_OF_SYNAPTIC_VESICLE_RECYCLING LANDIS_ERBB2_BREAST_TUMORS_65_UP, LANDIS_ERBB2_BREAST_TUMORS_65_UP  REACTOME_CROSS_PRESENTATION_OF_PARTICULATE_EXOGENOUS_ANTIGENS_PHAGOSOMES, REACTOME_CROSS_PRESENTATION_OF_PARTICULATE_EXOGENOUS_ANTIGENS_PHAGOSOMES GOBP_T_CELL_MIGRATION, GOBP_T_CELL_MIGRATION MCMURRAY_TP53_HRAS_COOPERATION_RESPONSE_UP, MCMURRAY_TP53_HRAS_COOPERATION_RESPONSE_UP
	MODULE_172, MODULE_172  LEE_LIVER_CANCER_DENA_UP, LEE_LIVER_CANCER_DENA_UP  BUSSLINGER_DUODENAL_EC_CELLS, BUSSLINGER_DUODENAL_EC_CELLS  GOBP_NEGATIVE_REGULATION_OF_CELL_KILLING, GOBP_NEGATIVE_REGULATION_OF_CELL_KILLING  GOBP_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GOBP_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS  KUWANO_RNA_STABILIZED_BY_NO, KUWANO_RNA_STABILIZED_BY_NO  GOMF_INTRAMEMBRANE_LIPID_TRANSPORTER_ACTIVITY, GOMF_INTRAMEMBRANE_LIPID_TRANSPORTER_ACTIVITY  GOBP_PLASMA_MEMBRANE_PHOSPHOLIPID_SCRAMBLING, GOBP_PLASMA_MEMBRANE_PHOSPHOLIPID_SCRAMBLING  MODULE_209, MODULE_209  GOBP_CELLULAR_RESPONSE_TO_MURAMYL_DIPEPTIDE, GOBP_CELLULAR_RESPONSE_TO_MURAMYL_DIPEPTIDE
	REACTOME_RHO_GTPASES_ACTIVATE_ROCKS, REACTOME_RHO_GTPASES_ACTIVATE_ROCKS RAY_TARGETS_OF_P210_BCR_ABL_FUSION_UP, RAY_TARGETS_OF_P210_BCR_ABL_FUSION_UP GOBP_NEGATIVE_REGULATION_OF_T_CELL_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_T_CELL_APOPTOTIC_PROCESS GOBP_ESTABLISHMENT_OF_EPITHELIAL_CELL_POLARITY, GOBP_ESTABLISHMENT_OF_EPITHELIAL_CELL_POLARITY GOMF_SERINE_TYPE_ENDOPEPTIDASE_INHIBITOR_ACTIVITY, GOMF_SERINE_TYPE_ENDOPEPTIDASE_INHIBITOR_ACTIVITY VALK_AML_WITH_CEBPA, VALK_AML_WITH_CEBPA WORSCHECH_TUMOR_REJECTION_UP, WORSCHECH_TUMOR_REJECTION_UP ZHENG_FOXP3_TARGETS_UP, ZHENG_FOXP3_TARGETS_UP GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_ENDOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_L_VIA_ER_PATHWAY, GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_ENDOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_L_VIA_I
	GOBP_POSITIVE_REGULATION_OF_ERAD_PATHWAY, GOBP_POSITIVE_REGULATION_OF_ERAD_PATHWAY PID_INTEGRIN5_PATHWAY, PID_INTEGRIN5_PATHWAY GOBP_MUSCLE_CELL_MIGRATION, GOBP_MUSCLE_CELL_MIGRATION GOBP_MUSCLE_CELL_MIGRATION, GOBP_MUSCLE_CELL_MIGRATION GOBP_POSITIVE_REGULATION_OF_BONE_RESORPTION, GOBP_POSITIVE_REGULATION_OF_BONE_RESORPTION GOBP_POSITIVE_REGULATION_OF_HORMONE_SECRETION, GOBP_POSITIVE_REGULATION_OF_HORMONE_SECRETION, HP_ANKYLOSIS, HP_ANKYLOSIS GOBP_POSITIVE_REGULATION_OF_FOCAL_ADHESION_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_FOCAL_ADHESION_ASSEMBLY GOCC_PERINUCLEAR_ENDOPLASMIC_RETICULUM, GOCC_PERINUCLEAR_ENDOPLASMIC_RETICULUM MIKKELSEN_PARTIALLY_REPROGRAMMED_TO_PLURIPOTENCY, MIKKELSEN_PARTIALLY_REPROGRAMMED_TO_PLURIPOTENCY KEGG_GLYCOSPHINGOLIPID_BIOSYNTHESIS_GLOBO_SERIES, KEGG_GLYCOSPHINGOLIPID_BIOSYNTHESIS_GLOBO_SERIES
	REACTOME_ERYTHROCYTES_TAKE_UP_OXYGEN_AND_RELEASE_CARBON_DIOXIDE, REACTOME_ERYTHROCYTES_TAKE_UP_OXYGEN_AND_RELEASE_CARBON_DIOXIDE  GOCC_ZONULA_ADHERENS, GOCC_ZONULA_ADHERENS  HP_EXTRAMEDULLARY_HEMATOPOIESIS, HP_EXTRAMEDULLARY_HEMATOPOIESIS  BOWIE_RESPONSE_TO_EXTRACELLULAR_MATRIX, BOWIE_RESPONSE_TO_EXTRACELLULAR_MATRIX  GOBP_REGULATION_OF_SYNAPTIC_VESICLE_ENDOCYTOSIS, GOBP_REGULATION_OF_SYNAPTIC_VESICLE_ENDOCYTOSIS  BIOCARTA_PGC1A_PATHWAY, BIOCARTA_PGC1A_PATHWAY  GOBP_POSITIVE_REGULATION_OF_MONOCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MONOCYTE_DIFFERENTIATION  MODULE_293, MODULE_293  GOBP_RESPONSE_TO_LIGHT_INTENSITY, GOBP_RESPONSE_TO_LIGHT_INTENSITY  GOBP_OVULATION_CYCLE_PROCESS, GOBP_OVULATION_CYCLE_PROCESS
	GOMF_1_PHOSPHATIDYLINOSITOL_BINDING, GOMF_1_PHOSPHATIDYLINOSITOL_BINDING GOMF_1_PHOSPHATIDYLINOSITOL_BINDING, GOMF_1_PHOSPHATIDYLINOSITOL_BINDING GOBP_REGULATION_OF_PHOSPHOLIPID_TRANSPORT, GOBP_REGULATION_OF_PHOSPHOLIPID_TRANSPORT HP_RHEUMATOID_FACTOR_POSITIVE, HP_RHEUMATOID_FACTOR_POSITIVE HP_AMYLOIDOSIS, HP_AMYLOIDOSIS  DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_COLUMNAR_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_COLUMNAR_CELLS BIOCARTA_MONOCYTE_PATHWAY, BIOCARTA_MONOCYTE_PATHWAY PID_SYNDECAN_4_PATHWAY, PID_SYNDECAN_4_PATHWAY VERRECCHIA_RESPONSE_TO_TGFB1_C5, VERRECCHIA_RESPONSE_TO_TGFB1_C5 GOBP_DRUG_TRANSMEMBRANE_TRANSPORT, GOBP_DRUG_TRANSMEMBRANE_TRANSPORT
	GOBP_REGULATION_OF_ACTIVATION_OF_JANUS_KINASE_ACTIVITY, GOBP_REGULATION_OF_ACTIVATION_OF_JANUS_KINASE_ACTIVITY GOBP_REGULATION_OF_T_CELL_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_T_CELL_APOPTOTIC_PROCESS GOBP_PARACRINE_SIGNALING, GOBP_PARACRINE_SIGNALING GOBP_POSITIVE_REGULATION_OF_ACTIVATION_OF_JANUS_KINASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_ACTIVATION_OF_JANUS_KINASE_ACTIVITY GOBP_REGULATION_OF_VOLTAGE_GATED_SODIUM_CHANNEL_ACTIVITY, GOBP_REGULATION_OF_VOLTAGE_GATED_SODIUM_CHANNEL_ACTIVITY GOMF_ASPARTIC_TYPE_ENDOPEPTIDASE_INHIBITOR_ACTIVITY, GOMF_ASPARTIC_TYPE_ENDOPEPTIDASE_INHIBITOR_ACTIVITY GOBP_POSITIVE_REGULATION_OF_PEPTIDE_HORMONE_SECRETION, GOBP_POSITIVE_REGULATION_OF_PEPTIDE_HORMONE_SECRETION MAHADEVAN_IMATINIB_RESISTANCE_DN, MAHADEVAN_IMATINIB_RESISTANCE_DN RICHERT_PBMC_HIV_LIPO_5_AGE_37_48YO_STIMULATED_VS_UNSTIMULATED_14W_TOP_FUNCTIONAL_NETWORK_UP GOMF_MYOSIN_V_BINDING, GOMF_MYOSIN_V_BINDING
	GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY, GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY  GOCC_DENDRITIC_SPINE_HEAD, GOCC_DENDRITIC_SPINE_HEAD  GOBP_CELLULAR_RESPONSE_TO_LIPOPROTEIN_PARTICLE_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_LIPOPROTEIN_PARTICLE_STIMULUS  GOBP_NEGATIVE_REGULATION_OF_VASOCONSTRICTION, GOBP_NEGATIVE_REGULATION_OF_VASOCONSTRICTION  GOBP_REGULATION_OF_GLYCOGEN_CATABOLIC_PROCESS, GOBP_REGULATION_OF_GLYCOGEN_CATABOLIC_PROCESS  REACTOME_ERYTHROCYTES_TAKE_UP_CARBON_DIOXIDE_AND_RELEASE_OXYGEN, REACTOME_ERYTHROCYTES_TAKE_UP_CARBON_DIOXIDE_AND_RELEASE_OXYGEN  GOBP_CD8_POSITIVE_ALPHA_BETA_T_CELL_PROLIFERATION, GOBP_CD8_POSITIVE_ALPHA_BETA_T_CELL_PROLIFERATION  GOBP_MEMBRANE_DEPOLARIZATION_DURING_ACTION_POTENTIAL, GOBP_MEMBRANE_DEPOLARIZATION_DURING_ACTION_POTENTIAL  DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_HORIZONTAL_BASAL_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_HORIZONTAL_BASAL_CELLS
	MIKKELSEN_IPS_LCP_WITH_H3K27ME3, MIKKELSEN_IPS_LCP_WITH_H3K27ME3 HP_PALMOPLANTAR_HYPERHIDROSIS, HP_PALMOPLANTAR_HYPERHIDROSIS RUAN_RESPONSE_TO_TNF_UP, RUAN_RESPONSE_TO_TNF_UP GOBP_PARANODAL_JUNCTION_ASSEMBLY, GOBP_PARANODAL_JUNCTION_ASSEMBLY GOMF_C_X3_C_CHEMOKINE_BINDING, GOMF_C_X3_C_CHEMOKINE_BINDING HP_INTERPHALANGEAL_JOINT_EROSIONS, HP_INTERPHALANGEAL_JOINT_EROSIONS TERAMOTO_OPN_TARGETS_CLUSTER_8, TERAMOTO_OPN_TARGETS_CLUSTER_8 GOBP_RESPONSE_TO_FOLLICLE_STIMULATING_HORMONE, GOBP_RESPONSE_TO_FOLLICLE_STIMULATING_HORMONE BIOCARTA_EOSINOPHILS_PATHWAY, BIOCARTA_EOSINOPHILS_PATHWAY YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_3, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_3
	HP_ABNORMAL_FOVEAL_MORPHOLOGY, HP_ABNORMAL_FOVEAL_MORPHOLOGY  GOBP_REGULATION_OF_ENTRY_OF_BACTERIUM_INTO_HOST_CELL, GOBP_REGULATION_OF_ENTRY_OF_BACTERIUM_INTO_HOST_CELL  GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_MIGRATION, GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_MIGRATION  WALLACE_PROSTATE_CANCER_DN, WALLACE_PROSTATE_CANCER_DN  HP_GLOSSITIS, HP_GLOSSITIS  GOMF_ARYLESTERASE_ACTIVITY, GOMF_ARYLESTERASE_ACTIVITY  GOMF_MHC_CLASS_I_PROTEIN_COMPLEX_BINDING, GOMF_MHC_CLASS_I_PROTEIN_COMPLEX_BINDING  REACTOME_ACROSOME_REACTION_AND_SPERM_OOCYTE_MEMBRANE_BINDING, REACTOME_ACROSOME_REACTION_AND_SPERM_OOCYTE_MEMBRANE_BINDING  GOBP_NEGATIVE_REGULATION_OF_NEUROBLAST_PROLIFERATION_GOBP_NEGATIVE_REGULATION_OF_NEUROBLAST_PROLIFERATION
	CASTELLANO_NRAS_TARGETS_DN, CASTELLANO_NRAS_TARGETS_DN CEBPE_TARGET_GENES, CEBPE_TARGET_GENES GOBP_PROTEIN_LOCALIZATION_TO_POSTSYNAPSE, GOBP_PROTEIN_LOCALIZATION_TO_POSTSYNAPSE HP_SYMPHALANGISM_AFFECTING_THE_PHALANGES_OF_THE_HAND, HP_SYMPHALANGISM_AFFECTING_THE_PHALANGES_OF_THE_HAND HP_PREMATURE_LOSS_OF_PRIMARY_TEETH, HP_PREMATURE_LOSS_OF_PRIMARY_TEETH GOBP_INTERLEUKIN_8_PRODUCTION, GOBP_INTERLEUKIN_8_PRODUCTION GOBP_REGULATION_OF_PROGRAMMED_NECROTIC_CELL_DEATH, GOBP_REGULATION_OF_PROGRAMMED_NECROTIC_CELL_DEATH GOBP_REGULATION_OF_NATURAL_KILLER_CELL_CHEMOTAXIS, GOBP_REGULATION_OF_NATURAL_KILLER_CELL_CHEMOTAXIS, GOBP_REGULATION_OF_NATURAL_KILLER_CELL_CHEMOTAXIS HP_ELBOW_ANKYLOSIS, HP_ELBOW_ANKYLOSIS HP_ABNORMALITY_OF_FOVEAL_PIGMENTATION, HP_ABNORMALITY_OF_FOVEAL_PIGMENTATION CORD_CORD_CORD_UNICTION_ASSEMBLY_CORD_CAP_UNICTION_CORD_CAP_UNICTION_ASSEMBLY_CORD_CAP_UNICTION_CORD
	GOBP_GAP_JUNCTION_ASSEMBLY, GOBP_GAP_JUNCTION_ASSEMBLY GOBP_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE, GOBP_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_SUSTAINED_IN_MONOCYTE_DN, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_SUSTAINED_IN_MONOCYTE_DN ISHIKAWA_STING_SIGNALING, ISHIKAWA_STING_SIGNALING CROONQUIST_IL6_DEPRIVATION_UP, CROONQUIST_IL6_DEPRIVATION_UP LA_MEN1_TARGETS, LA_MEN1_TARGETS GOBP_CARBON_DIOXIDE_TRANSPORT, GOBP_CARBON_DIOXIDE_TRANSPORT GOBP_DENDRITE_SELF_AVOIDANCE, GOBP_DENDRITE_SELF_AVOIDANCE GOCC_SPERMATOPROTEASOME_COMPLEX, GOCC_SPERMATOPROTEASOME_COMPLEX
	GOMF_NEUREGULIN_BINDING, GOMF_NEUREGULIN_BINDING GOMF_EXTRACELLULAR_MATRIX_PROTEIN_BINDING, GOMF_EXTRACELLULAR_MATRIX_PROTEIN_BINDING, HP_IMPAIRED_USE_OF_NONVERBAL_BEHAVIORS, HP_IMPAIRED_USE_OF_NONVERBAL_BEHAVIORS GIAROLA_SILVA_BLOOD_PANDEMRIX_AGE_21_51YO_3DY_UP, GIAROLA_SILVA_BLOOD_PANDEMRIX_AGE_21_51YO_3DY_UP GOBP_POSITIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_ACTIVITY HP_SWAN_NECK_LIKE_DEFORMITIES_OF_THE_FINGERS, HP_SWAN_NECK_LIKE_DEFORMITIES_OF_THE_FINGERS GOBP_MYOTUBE_DIFFERENTIATION_INVOLVED_IN_SKELETAL_MUSCLE_REGENERATION, GOBP_MYOTUBE_DIFFERENTIATION_INVOLVED_IN_SKELETAL_MUSCLE_REGENERATION YAMANAKA_GLIOBLASTOMA_SURVIVAL_DN, YAMANAKA_GLIOBLASTOMA_SURVIVAL_DN GOBP_POSITIVE_REGULATION_OF_T_CELL_APOPTOTIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_T_CELL_APOPTOTIC_PROCESS WP_PLATELETMEDIATED_INTERACTIONS_WITH_VASCULAR_AND_CIRCULATING_CELLS, WP_PLATELETMEDIATED_INTERACTIONS_WITH_VASCULAR_AND_CIRCULATING_CELLS
	GOBP_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE, GOBP_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE MURAKAMI_UV_RESPONSE_1HR_DN, MURAKAMI_UV_RESPONSE_1HR_DN HOFFMAN_CLOCK_TARGETS_DN, HOFFMAN_CLOCK_TARGETS_DN SOUCEK_MYC_TARGETS, SOUCEK_MYC_TARGETS BASSO_HAIRY_CELL_LEUKEMIA_DN, BASSO_HAIRY_CELL_LEUKEMIA_DN SHARMA_ASTROCYTOMA_WITH_NFI_SYNDROM, SHARMA_ASTROCYTOMA_WITH_NFI_SYNDROM HP_PROXIMAL_SYMPHALANGISM, HP_PROXIMAL_SYMPHALANGISM GOBP_ASYMMETRIC_NEUROBLAST_DIVISION, GOBP_ASYMMETRIC_NEUROBLAST_DIVISION GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_CHEMOTAXIS, GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_CHEMOTAXIS
	GOMF_MHC_CLASS_IB_RECEPTOR_ACTIVITY, GOMF_MHC_CLASS_IB_RECEPTOR_ACTIVITY GOBP_REGULATION_OF_MEMBRANE_DEPOLARIZATION_DURING_ACTION_POTENTIAL, GOBP_REGULATION_OF_MEMBRANE_DEPOLARIZATION_DURING_ACTION_POTENTIAL DEBOSSCHER_NFKB_TARGETS_REPRESSED_BY_GLUCOCORTICOIDS, DEBOSSCHER_NFKB_TARGETS_REPRESSED_BY_GLUCOCORTICOIDS HP_INCREASED_CIRCULATING_RENIN_LEVEL, HP_INCREASED_CIRCULATING_RENIN_LEVEL REACTOME_ACYL_CHAIN_REMODELLING_OF_PS, REACTOME_ACYL_CHAIN_REMODELLING_OF_PS HP_NAIL_PITS, HP_NAIL_PITS HP_HYPOLIPIDEMIA, HP_HYPOLIPIDEMIA HP_HYPOLIPIDEMIA, HP_HYPOCHLOREMIA HP_HYPOCHLOREMIA, HP_HYPOCHLOREMIA GAO_LARGE_INTESTINE_24W_C4_PROM1LOW_PROGENITOR, GAO_LARGE_INTESTINE_24W_C4_PROM1LOW_PROGENITOR GOBP_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_ACTIVITY
	GOBP_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_ACTIVITY chr12q22, chr12q22 GOBP_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY_INVOLVED_IN_HEART_PROCESS GOBP_MYOBLAST_FUSION_INVOLVED_IN_SKELETAL_MUSCLE_REGENERATION, GOBP_MYOBLAST_FUSION_INVOLVED_IN_SKELETAL_MUSCLE_REGENERATION GOBP_NEGATIVE_REGULATION_BY_HOST_OF_SYMBIONT_MOLECULAR_FUNCTION, GOBP_NEGATIVE_REGULATION_BY_HOST_OF_SYMBIONT_MOLECULAR_FUNCTION GOMF_CCR1_CHEMOKINE_RECEPTOR_BINDING, GOMF_CCR1_CHEMOKINE_RECEPTOR_BINDING GOBP_REGULATION_OF_MEMBRANE_DEPOLARIZATION_DURING_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL, GOBP_REGULATION_OF_MEMBRANE_DEPOLARIZATION_DURING_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL GOBP_NEGATIVE_REGULATION_OF_IRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE GOBP_FOAM_CELL_DIFFERENTIATION, GOBP_FOAM_CELL_DIFFERENTIATION GOBP_REGULATION_OF_MACROPHAGE_DERIVED_FOAM_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_MACROPHAGE_DERIVED_FOAM_CELL_DIFFERENTIATION KEGG_RETINOL_METABOLISM, KEGG_RETINOL_METABOLISM
	KEGG_RETINOL_METABOLISM, KEGG_RETINOL_METABOLISM  KIM_GLIS2_TARGETS_DN, KIM_GLIS2_TARGETS_DN  GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY, GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY  GOBP_POSITIVE_REGULATION_OF_CARDIAC_MUSCLE_CONTRACTION, GOBP_POSITIVE_REGULATION_OF_CARDIAC_MUSCLE_CONTRACTION  HP_HYPOTRIGLYCERIDEMIA, HP_HYPOTRIGLYCERIDEMIA  GOBP_POSITIVE_REGULATION_OF_STRIATED_MUSCLE_CONTRACTION, GOBP_POSITIVE_REGULATION_OF_STRIATED_MUSCLE_CONTRACTION  HP_ABNORMAL_CIRCULATING_RENIN, HP_ABNORMAL_CIRCULATING_RENIN  HP_FETAL_POLYURIA, HP_FETAL_POLYURIA  GOBP_NEGATIVE_REGULATION_OF_INTERLEUKIN_8_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_INTERLEUKIN_8_PRODUCTION