	GSE29618_MONOCYTE_VS_MDC_UP, GSE29618_MGGSE22886_DAY0_VS_DAY7_MONOCYTE_IN_CULT	ONOCYTE_VS_MDC_UP URE_UP, GSE22886_DAY0_VS_DAY7_MONOCYTE_IN_CULTURE_UP
	GSE19888_ADENOSINE_A3R_INH_PRETREAT_ANI GSE13411_IGM_VS_SWITCHED_MEMORY_BCELL_ GSE19888_CTRL_VS_A3R_INHIBITOR_TREATED_M GSE42088_UNINF_VS_LEISHMANIA_INF_DC_24H_	
	GSE37416_12H_VS_24H_F_TULARENSIS_LVS_NEUT GSE36826_NORMAL_VS_STAPH_AUREUS_INF_IL1 GSE9988_ANTI_TREM1_VS_LOW_LPS_MONOCYTE GSE34515_CD16_POS_MONOCYTE_VS_DC_DN, GS GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_24I GSE12366_GC_VS_MEMORY_BCELL_DN, GSE12366	TROPHIL_DN, GSE37416_12H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN .R_KO_SKIN_UP, GSE36826_NORMAL_VS_STAPH_AUREUS_INF_IL1R_KO_SKIN_UP E_UP, GSE9988_ANTI_TREM1_VS_LOW_LPS_MONOCYTE_UP E34515_CD16_POS_MONOCYTE_VS_DC_DN H_CD4_TCELL_UP, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_24H_CD4_TCELL_UP
	GSE42021_TCONV_PLN_VS_CD24HI_TCONV_THYNGSE34515_CD16_NEG_MONOCYTE_VS_DC_UP, GSGSE42021_CD24HI_VS_CD24INT_TREG_THYMUS_IGSE36826_NORMAL_VS_STAPH_AUREUS_INF_SKIGSE32986_CURDLAN_LOWDOSE_VS_GMCSF_ANIGSE12366_GC_VS_NAIVE_BCELL_DN, GSE12366_GG	DN, GSE42021_CD24HI_VS_CD24INT_TREG_THYMUS_DN IN_DN, GSE36826_NORMAL_VS_STAPH_AUREUS_INF_SKIN_DN D_CURDLAN_LOWDOSE_STIM_DC_UP, GSE32986_CURDLAN_LOWDOSE_VS_GMCSF_AND_CURDLAN_LOWDOSE_STIM_DC_UP C_VS_NAIVE_BCELL_DN
	GSE10325_MYELOID_VS_LUPUS_MYELOID_DN, GSGSE41176_WT_VS_TAK1_KO_ANTI_IGM_STIM_BCSGSE20P.V1_UP, ERB2_UP.V1_UP GSE25088_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_VS_ACT_CD4_TCESTITE GSE9960_HEALTHY_VS_GRAM_NEG_SEPSIS_PBMG	ELL_6H_DN, GSE41176_WT_VS_TAK1_KO_ANTI_IGM_STIM_BCELL_6H_DN .ITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_UP, GSE25088_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_UP LL_48H_DN, GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_DN C_DN, GSE9960_HEALTHY_VS_GRAM_NEG_SEPSIS_PBMC_DN
	GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_I GSE1460_INTRATHYMIC_T_PROGENITOR_VS_NA GSE42021_TREG_PLN_VS_CD24INT_TREG_THYMU GSE2770_UNTREATED_VS_IL12_TREATED_ACT_C GSE28737_BCL6_HET_VS_BCL6_KO_MARGINAL_Z GSE30971_WBP7_HET_VS_KO_MACROPHAGE_4H_	UP, GSE3982_MAC_VS_CENT_MEMORY_CD4_TCELL_UP DN, GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_DN IVE_CD4_TCELL_ADULT_BLOOD_DN, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_DN IS_DN, GSE42021_TREG_PLN_VS_CD24INT_TREG_THYMUS_DN ID4_TCELL_2H_UP, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_2H_UP IONE_BCELL_DN, GSE28737_BCL6_HET_VS_BCL6_KO_MARGINAL_ZONE_BCELL_DN LPS_STIM_DN, GSE30971_WBP7_HET_VS_KO_MACROPHAGE_4H_LPS_STIM_DN KO_ANTI_CD3_STIM_DP_THYMOCYTES_DN, GSE21546_SAP1A_KO_VS_SAP1A_KO_AND_ELK1_KO_ANTI_CD3_STIM_DP_THYMOCYTES_DN
	GSE26030_UNSTIM_VS_RESTIM_TH1_DAY15_POST GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCE GSE39556_UNTREATED_VS_3H_POLYIC_INJ_MOU GSE44732_UNSTIM_VS_IL27_STIM_IMATURE_DC_I GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_G GSE7831_1H_VS_4H_CPG_STIM_PDC_UP, GSE7831_	T_POLARIZATION_UP, GSE26030_UNSTIM_VS_RESTIM_TH1_DAY15_POST_POLARIZATION_UP ELL_2H_DN, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_DN USE_NK_CELL_DN, GSE39556_UNTREATED_VS_3H_POLYIC_INJ_MOUSE_NK_CELL_DN DN, GSE44732_UNSTIM_VS_IL27_STIM_IMATURE_DC_DN CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP _1H_VS_4H_CPG_STIM_PDC_UP
	GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_H GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCEN GSE19198_CTRL_VS_IL21_TREATED_TCELL_1H_DN GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_ GSE28737_WT_VS_BCL6_HET_FOLLICULAR_BCELN GSE40685_TREG_VS_FOXP3_KO_TREG_PRECURSO	_HSC_UP, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_HSC_UP IIGHDOSE_STIM_DC_UP, GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_UP LL_ADULT_BLOOD_DN, GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_DN N, GSE19198_CTRL_VS_IL21_TREATED_TCELL_1H_DN _6H_UP, GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_6H_UP L_DN, GSE28737_WT_VS_BCL6_HET_FOLLICULAR_BCELL_DN DR_DN, GSE40685_TREG_VS_FOXP3_KO_TREG_PRECURSOR_DN
	GSE7764_IL15_NK_CELL_24H_VS_SPLENOCYTE_DEGSE13411_NAIVE_VS_SWITCHED_MEMORY_BCELGSE9988_LOW_LPS_VS_ANTI_TREM1_AND_LPS_MGSE22229_RENAL_TRANSPLANT_IMMUNOSUPP_	LINE_DN, GSE8384_CTRL_VS_B_ABORTUS_4H_MAC_CELL_LINE_DN IN, GSE7764_IL15_NK_CELL_24H_VS_SPLENOCYTE_DN L_DN, GSE73411_NAIVE_VS_SWITCHED_MEMORY_BCELL_DN MONOCYTE_DN, GSE9988_LOW_LPS_VS_ANTI_TREM1_AND_LPS_MONOCYTE_DN THERAPY_VS_HEALTHY_PBMC_DN, GSE22229_RENAL_TRANSPLANT_IMMUNOSUPP_THERAPY_VS_HEALTHY_PBMC_DN M_DN, GSE7509_DC_VS_MONOCYTE_WITH_FCGRIIB_STIM_DN I1_UP
	GSE3982_MAST_CELL_VS_MAC_DN, GSE3982_MAS	CELL_DAY3_DN, GSE24634_IL4_VS_CTRL_TREATED_NAIVE_CD4_TCELL_DAY3_DN ST_CELL_VS_MAC_DN CELL_UP, GSE3982_EOSINOPHIL_VS_EFF_MEMORY_CD4_TCELL_UP /S_PDC_UP 3568_ID3_KO_VS_WT_CD8_TCELL_DN
	GSE41867_NAIVE_VS_DAY6_LCMV_EFFECTOR_CEGSE14000_UNSTIM_VS_4H_LPS_DC_DN, GSE14000_HALLMARK_HEME_METABOLISM, HALLMARK_EGSE2405_0H_VS_3H_A_PHAGOCYTOPHILUM_STIMESE21360_NAIVE_VS_SECONDARY_MEMORY_CD	D8_TCELL_UP, GSE41867_NAIVE_VS_DAY6_LCMV_EFFECTOR_CD8_TCELL_UP _UNSTIM_VS_4H_LPS_DC_DN HEME_METABOLISM M_NEUTROPHIL_UP, GSE2405_0H_VS_3H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_UP 18_TCELL_DN, GSE21360_NAIVE_VS_SECONDARY_MEMORY_CD8_TCELL_DN LPSLIKE_STIM_DC_3H_UP, GSE4748_LPS_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP
	GSE3982_MAC_VS_BCELL_UP, GSE3982_MAC_VS_I GSE2706_2H_VS_8H_R848_STIM_DC_UP, GSE2706_2 GSE22886_CTRL_VS_LPS_24H_DC_DN, GSE22886_C GSE29618_BCELL_VS_MDC_UP, GSE29618_BCELL_V EGFR_UP.V1_UP, EGFR_UP.V1_UP GSE21670_STAT3_KO_VS_WT_CD4_TCELL_IL6_TRI	BCELL_UP 2H_VS_8H_R848_STIM_DC_UP CTRL_VS_LPS_24H_DC_DN
	MODULE_321, MODULE_321 BROCKE_APOPTOSIS_REVERSED_BY_IL6, BROCKE GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL10_STIN GSE19198_CTRL_VS_IL21_TREATED_TCELL_6H_DN GSE12366_NAIVE_VS_MEMORY_BCELL_DN, GSE12 GSE26030_UNSTIM_VS_RESTIM_TH1_DAY15_POST	E_APOPTOSIS_REVERSED_BY_IL6 M_MACROPHAGE_45MIN_DN, GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_DN N, GSE19198_CTRL_VS_IL21_TREATED_TCELL_6H_DN 2366_NAIVE_VS_MEMORY_BCELL_DN I_POLARIZATION_DN, GSE26030_UNSTIM_VS_RESTIM_TH1_DAY15_POST_POLARIZATION_DN
	GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_W GSE29618_BCELL_VS_PDC_DAY7_FLU_VACCINE_U GSE7509_UNSTIM_VS_IFNA_STIM_IMMATURE_DC GSE40273_XBP1_KO_VS_WT_TREG_DN, GSE40273_X GSE33425_CD8_ALPHAALPHA_VS_ALPHABETA_C GSE7219_WT_VS_NIK_NFKB2_KO_DC_UP, GSE7219	CD161_HIGH_TCELL_UP, GSE33425_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_UP 9_WT_VS_NIK_NFKB2_KO_DC_UP
	PETROVA_ENDOTHELIUM_LYMPHATIC_VS_BLOGSE19198_1H_VS_24H_IL21_TREATED_TCELL_DN, SMIRNOV_CIRCULATING_ENDOTHELIOCYTES_INGSE40666_WT_VS_STAT1_KO_CD8_TCELL_UP, GSEGSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_GSE19198_6H_VS_24H_IL21_TREATED_TCELL_UP,	N_CANCER_UP, SMIRNOV_CIRCULATING_ENDOTHELIOCYTES_IN_CANCER_UP E40666_WT_VS_STAT1_KO_CD8_TCELL_UP _12H_UP, GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_12H_UP GSE19198_6H_VS_24H_IL21_TREATED_TCELL_UP
	GSE17721_POLYIC_VS_GARDIQUIMOD_16H_BMD0	_48H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_48H_CD8_T_CELL_DN C_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_16H_BMDC_UP 5_DN, GSE42021_CD24HI_VS_CD24INT_TCONV_THYMUS_DN CD4_TCELL_VS_BCELL_DN
	GSE2585_AIRE_KO_VS_WT_CD80_LOW_MTEC_DN GSE17721_POLYIC_VS_GARDIQUIMOD_2H_BMDC GSE28737_BCL6_HET_VS_BCL6_KO_FOLLICULAR_ GSE2706_UNSTIM_VS_8H_R848_DC_UP, GSE2706_U LTE2_UP.V1_UP, LTE2_UP.V1_UP GSE22140_GERMFREE_VS_SPF_ARTHRITIC_MOUS	I, GSE2585_AIRE_KO_VS_WT_CD80_LOW_MTEC_DN C_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_2H_BMDC_DN BCELL_DN, GSE28737_BCL6_HET_VS_BCL6_KO_FOLLICULAR_BCELL_DN
	GSE6259_FLT3L_INDUCED_33D1_POS_DC_VS_CD8 GSE3039_NKT_CELL_VS_ALPHAALPHA_CD8_TCE GSE10856_CTRL_VS_TNFRSF6B_IN_MACROPHAGE GSE22886_DAY1_VS_DAY7_MONOCYTE_IN_CULT ALCALAY_AML_BY_NPM1_LOCALIZATION_UP, A	_AB_KNOCKIN_TCELL_2H_UP, GSE36888_UNTREATED_VS_IL2_TREATED_STAT5_AB_KNOCKIN_TCELL_2H_UP 3_TCELL_UP, GSE6259_FLT3L_INDUCED_33D1_POS_DC_VS_CD8_TCELL_UP 6LL_UP, GSE3039_NKT_CELL_VS_ALPHAALPHA_CD8_TCELL_UP 6_UP, GSE10856_CTRL_VS_TNFRSF6B_IN_MACROPHAGE_UP 6_UP, GSE10856_CTRL_VS_TNFRSF6B_IN_MACROPHAGE_UP 6_URE_UP, GSE22886_DAY1_VS_DAY7_MONOCYTE_IN_CULTURE_UP 6_URE_UP, GSE22886_DAY1_VS_DAY7_MONOCYTE_IN_CULTURE_UP 6_UCALAY_AML_BY_NPM1_LOCALIZATION_UP 6_TCELL_DN, GSE13738_TCR_VS_BYSTANDER_ACTIVATED_CD4_TCELL_DN
	GSE7852_TREG_VS_TCONV_FAT_UP, GSE7852_TRE GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_ GSE23321_CENTRAL_MEMORY_VS_NAIVE_CD8_T	EG_VS_TCONV_FAT_UP _17H_DN, GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_17H_DN _CELL_UP, GSE23321_CENTRAL_MEMORY_VS_NAIVE_CD8_TCELL_UP _BA_DC_DN, GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8A_DC_DN _15_0H_VS_24H_OZONE_LUNG_UP NKT_CELL_VS_B2_BCELL_DN
	GSE24142_DN2_VS_DN3_THYMOCYTE_FETAL_UP_GSE12845_PRE_GC_VS_DARKZONE_GC_TONSIL_EGO_POSITIVE_REGULATION_OF_VASCULATURE_GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WGSE46606_IRF4_KO_VS_WT_CD40L_IL2_IL5_1DAY_	, GSE24142_DN2_VS_DN3_THYMOCYTE_FETAL_UP BCELL_UP, GSE12845_PRE_GC_VS_DARKZONE_GC_TONSIL_BCELL_UP _DEVELOPMENT, GO_POSITIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT BP7_KO_4H_UP, GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_KO_4H_UP _STIMULATED_BCELL_DN, GSE46606_IRF4_KO_VS_WT_CD40L_IL2_IL5_1DAY_STIMULATED_BCELL_DN CD4_TCELL_DN, GSE20500_CTRL_VS_RETINOIC_ACID_TREATED_CD4_TCELL_DN
	GSE17721_12H_VS_24H_PAM3CSK4_BMDC_DN, GSGSE3982_BASOPHIL_VS_NKCELL_UP, GSE3982_BARAF_UP.V1_UP, RAF_UP.V1_UPGSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_GSE14769_UNSTIM_VS_40MIN_LPS_BMDM_DN, GSGSE37605_C57BL6_VS_NOD_FOXP3_FUSION_GFP_T	SE17721_12H_VS_24H_PAM3CSK4_BMDC_DN SOPHIL_VS_NKCELL_UP _CD4_TCELL_2H_DN, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_2H_DN SE14769_UNSTIM_VS_40MIN_LPS_BMDM_DN TREG_DN, GSE37605_C57BL6_VS_NOD_FOXP3_FUSION_GFP_TREG_DN
	ZHU_CMV_ALL_DN, ZHU_CMV_ALL_DN GSE1460_DP_THYMOCYTE_VS_NAIVE_CD4_TCELL GAUSSMANN_MLL_AF4_FUSION_TARGETS_F_UF GSE26488_WT_VS_VP16_TRANSGENIC_HDAC7_KO	RECEPTOR_ACTIVITY 10D_DN, TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_10D_DN L_ADULT_BLOOD_DN, GSE1460_DP_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_DN P, GAUSSMANN_MLL_AF4_FUSION_TARGETS_F_UP D_DOUBLE_POSITIVE_THYMOCYTE_DN, GSE26488_WT_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN
	GSE7460_TCONV_VS_TREG_THYMUS_UP, GSE7460 BROWNE_HCMV_INFECTION_24HR_DN, BROWNI GSE2128_CTRL_VS_MIMETOPE_NEGATIVE_SELEC GSE36476_YOUNG_VS_OLD_DONOR_MEMORY_CI CHIARADONNA_NEOPLASTIC_TRANSFORMATIO	
	SARRIO_EPITHELIAL_MESENCHYMAL_TRANSITIONDER_CDH1_TARGETS_1_UP, ONDER_CDH1_TARGETS_1_UP.V1_UP	CED_TREG_IL2_CULTURE_UP, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_UP ION_DN, SARRIO_EPITHELIAL_MESENCHYMAL_TRANSITION_DN ARGETS_1_UP IBP7_KO_2H_UP, GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_KO_2H_UP
	GSE21774_CD62L_POS_CD56_BRIGHT_VS_CD62L_N E2F1_UP.V1_DN, E2F1_UP.V1_DN GSE13484_3H_UNSTIM_VS_YF17D_VACCINE_STIM CHIANG_LIVER_CANCER_SUBCLASS_CTNNB1_D GSE5142_CTRL_VS_HTERT_TRANSDUCED_CD8_TG	_UP, GSE17974_0H_VS_4H_IN_VITRO_ACT_CD4_TCELL_UP NEG_CD56_DIM_NK_CELL_UP, GSE21774_CD62L_POS_CD56_BRIGHT_VS_CD62L_NEG_CD56_DIM_NK_CELL_UP 1_PBMC_DN, GSE13484_3H_UNSTIM_VS_YF17D_VACCINE_STIM_PBMC_DN N, CHIANG_LIVER_CANCER_SUBCLASS_CTNNB1_DN CELL_EARLY_PASSAGE_CLONE_DN, GSE5142_CTRL_VS_HTERT_TRANSDUCED_CD8_TCELL_EARLY_PASSAGE_CLONE_DN UP, GSE24142_ADULT_VS_FETAL_DN2_THYMOCYTE_UP
	GSE42021_CD24INT_VS_CD24LOW_TCONV_THYM GSE27670_CTRL_VS_LMP1_TRANSDUCED_GC_BCI GSE42021_TCONV_PLN_VS_CD24LO_TCONV_THY GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_WI GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_ MODULE_491, MODULE_491	IUS_DN, GSE42021_CD24INT_VS_CD24LOW_TCONV_THYMUS_DN ELL_DN, GSE27670_CTRL_VS_LMP1_TRANSDUCED_GC_BCELL_DN 'MUS_DN, GSE42021_TCONV_PLN_VS_CD24LO_TCONV_THYMUS_DN ITH_EGR2_DELETED_UP, GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_WITH_EGR2_DELETED_UP _MEMORY_BCELL_DAY40_DN, GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_MEMORY_BCELL_DAY40_DN
	GSE2706_R848_VS_LPS_8H_STIM_DC_DN, GSE2706_ GSE36888_UNTREATED_VS_IL2_TREATED_STAT5_ MODULE_122, MODULE_122 GSE21360_SECONDARY_VS_QUATERNARY_MEMO GSE3982_DC_VS_MAC_LPS_STIM_DN, GSE3982_DC GSE6259_FLT3L_INDUCED_33D1_POS_DC_VS_CD4	_AB_KNOCKIN_TCELL_6H_UP, GSE36888_UNTREATED_VS_IL2_TREATED_STAT5_AB_KNOCKIN_TCELL_6H_UP ORY_CD8_TCELL_UP, GSE21360_SECONDARY_VS_QUATERNARY_MEMORY_CD8_TCELL_UP
	GSE32423_CTRL_VS_IL4_MEMORY_CD8_TCELL_UI KATSANOU_ELAVL1_TARGETS_UP, KATSANOU_ GSE22886_NAIVE_BCELL_VS_BLOOD_PLASMA_CF GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN, C GSE14000_UNSTIM_VS_4H_LPS_DC_TRANSLATED	ELL_UP, GSE22886_NAIVE_BCELL_VS_BLOOD_PLASMA_CELL_UP GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN D_RNA_DN, GSE14000_UNSTIM_VS_4H_LPS_DC_TRANSLATED_RNA_DN
	GSE11924_TH1_VS_TH17_CD4_TCELL_DN, GSE1192 GSE360_CTRL_VS_T_GONDII_MAC_DN, GSE360_CTVANHARANTA_UTERINE_FIBROID_DN, VANHARANTA_UTERINE_FIBROID_DN, VANHARANTA_UTERINE_FIBROID_DN, GSE17721_PAM3CSK4_VS_CPG_12H_BMDC_DN, GSE17721_PAM3CSK4_VS_CPG_12H_BMDC_DN, GSECONS_EARCH GSE3982_EOSINOPHIL_VS_BASOPHIL_UP, GSE3982_GSE20366_TREG_VS_TCONV_DN, GSE20366_TREG_	TRL_VS_T_GONDII_MAC_DN RANTA_UTERINE_FIBROID_DN SE17721_PAM3CSK4_VS_CPG_12H_BMDC_DN RLY_UP, VECCHI_GASTRIC_CANCER_ADVANCED_VS_EARLY_UP 2_EOSINOPHIL_VS_BASOPHIL_UP
	MODULE_345, MODULE_345 LEF1_UP.V1_DN, LEF1_UP.V1_DN GSE20715_WT_VS_TLR4_KO_24H_OZONE_LUNG_U GSE32034_UNTREATED_VS_ROSIGLIZATONE_TRE GSE14415_NATURAL_TREG_VS_TCONV_UP, GSE14 GSE11961_MEMORY_BCELL_DAY7_VS_GERMINAL	L_CENTER_BCELL_DAY7_DN, GSE11961_MEMORY_BCELL_DAY7_VS_GERMINAL_CENTER_BCELL_DAY7_DN
	GSE43863_TH1_VS_LY6C_INT_CXCR5POS_MEMOR	8_CTRL_VS_LPS_STIM_DC_3H_DN _TH2_UP
	MODULE_436, MODULE_436	GETS_UP URA_ADIPOGENESIS_LATE_UP
	MARKEY_RB1_CHRONIC_LOF_DN, MARKEY_RB1_	_CHRONIC_LOF_DN P, GSE3982_CENT_MEMORY_CD4_TCELL_VS_TH2_UP E_HCMV_INFECTION_18HR_DN MEN ENANCE_OF_LOCATION
	GSE17721_PAM3CSK4_VS_GADIQUIMOD_0.5H_BM REACTOME_ACTIVATED_TLR4_SIGNALLING, REACTOME_ACTIVATED_TLR4_SIGNALLING, REACTOME_ACTIVATED_TLRA_SI	C_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_12H_BMDC_UP HAGE_45MIN_DN, GSE5589_LPS_VS_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_DN BCELL_UP, GSE28449_WT_VS_LRF_KO_GERMINAL_CENTER_BCELL_UP
	FRASOR_RESPONSE_TO_ESTRADIOL_DN, FRASOR_GSE17721_CTRL_VS_LPS_2H_BMDC_DN, GSE17721_GSE360_T_GONDII_VS_M_TUBERCULOSIS_DC_DNGSE46606_IRF4_KO_VS_WT_UNSTIM_BCELL_DN, GO_PLATELET_DEGRANULATION, GO_PLATELET_GO_CELL_REDOX_HOMEOSTASIS, GO_CELL_RED	R_RESPONSE_TO_ESTRADIOL_DN _CTRL_VS_LPS_2H_BMDC_DN N, GSE360_T_GONDII_VS_M_TUBERCULOSIS_DC_DN GSE46606_IRF4_KO_VS_WT_UNSTIM_BCELL_DN T_DEGRANULATION
	GSE45739_UNSTIM_VS_ACD3_ACD28_STIM_WT_C WESTON_VEGFA_TARGETS, WESTON_VEGFA_TA GO_GLYCERALDEHYDE_3_PHOSPHATE_METABO REACTOME_PHASE1_FUNCTIONALIZATION_OF_ REACTOME_MYD88_MAL_CASCADE_INITIATED_ VART_KSHV_INFECTION_ANGIOGENIC_MARKEI	D4_TCELL_UP, GSE45739_UNSTIM_VS_ACD3_ACD28_STIM_WT_CD4_TCELL_UP ARGETS DLIC_PROCESS, GO_GLYCERALDEHYDE_3_PHOSPHATE_METABOLIC_PROCESS _COMPOUNDS, REACTOME_PHASE1_FUNCTIONALIZATION_OF_COMPOUNDS _ON_PLASMA_MEMBRANE, REACTOME_MYD88_MAL_CASCADE_INITIATED_ON_PLASMA_MEMBRANE RS_DN, VART_KSHV_INFECTION_ANGIOGENIC_MARKERS_DN
	GSE6269_HEALTHY_VS_STAPH_AUREUS_INF_PBM GSE16385_ROSIGLITAZONE_IL4_VS_IFNG_TNF_ST GSE46606_IRF4HIGH_VS_WT_CD40L_IL2_IL5_DAY HELLEBREKERS_SILENCED_DURING_TUMOR_AN RODRIGUES_NTN1_TARGETS_DN, RODRIGUES_N SEKI_INFLAMMATORY_RESPONSE_LPS_UP, SEKI_	_INFLAMMATORY_RESPONSE_LPS_UP
	P53_DN.V2_UP, P53_DN.V2_UP GSE1925_CTRL_VS_3H_IFNG_STIM_IFNG_PRIMED PARK_TRETINOIN_RESPONSE_AND_PML_RARA_ GO_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_3 BMI1_DN.V1_UP, BMI1_DN.V1_UP	PROCESS, GO_REACTIVE_OXYGEN_SPECIES_METABOLIC_PROCESS
S_MDC_DAY7_FLU_VACCINE_UP, GSE29618_MONOCYTE_VS_MDC_DAY7_FLU_VACCINE_UP	GO_T_CELL_MEDIATED_IMMUNITY, GO_T_CELL_BOYLAN_MULTIPLE_MYELOMA_PCA1_UP, BOYL_GSE3982_NKCELL_VS_TH2_UP, GSE3982_NKCELL_GSE20366_TREG_VS_NAIVE_CD4_TCELL_DN, GSE20366_TREG_VS_NAIVE_CD4_TCELL_DN, GSE2036_TCELL_DN, GSE2036_TCELL_DN, GSE2036_TCELL_DN, GSE2036_TCELL_DN, GSE2036_TCELL_DN, GSE205_TCELL_DN, GSE205_TCELL_TC	MAC_DN, GSE360_T_GONDII_VS_B_MALAYI_HIGH_DOSE_MAC_DN _MEDIATED_IMMUNITY .AN_MULTIPLE_MYELOMA_PCA1_UP _VS_TH2_UP 20366_TREG_VS_NAIVE_CD4_TCELL_DN
	ALTEMEIER_RESPONSE_TO_LPS_WITH_MECHAN WINZEN_DEGRADED_VIA_KHSRP, WINZEN_DEG	SCRIPTION_IN_RESPONSE_TO_STRESS, GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_IN_RESPONSE_TO_STRESS ECKNER_TARGETS_OF_MIRLET7A3_UP JIST_NRAS_SIGNALING_UP
	GSE36891_UNSTIM_VS_POLYIC_TLR3_STIM_PERIT	S, GO_CELLULAR_RESPONSE_TO_OXYGEN_LEVELS TONEAL_MACROPHAGE_UP, GSE36891_UNSTIM_VS_POLYIC_TLR3_STIM_PERITONEAL_MACROPHAGE_UP _IGLK_UP, MAGRANGEAS_MULTIPLE_MYELOMA_IGLL_VS_IGLK_UP
	GSE3039_CD4_TCELL_VS_ALPHABETA_CD8_TCELLLAN_LIPA_TARGETS_3M, LIAN_LIPA_TARGETS_3 GSE39820_IL1B_IL6_VS_IL1B_IL6_IL23A_TREATED_ GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROF GO_FAT_CELL_DIFFERENTIATION, GO_FAT_CELI	_CD4_TCELL_DN, GSE39820_IL1B_IL6_VS_IL1B_IL6_IL23A_TREATED_CD4_TCELL_DN PHAGE_180MIN_DN, GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_180MIN_DN
	LANDIS_ERBB2_BREAST_PRENEOPLASTIC_DN, LA KEGG_ADIPOCYTOKINE_SIGNALING_PATHWAY RADMACHER_AML_PROGNOSIS, RADMACHER_A	TH1_VS_TH2_48H_ACT_UP TION_MEDIATED_BY_TLR4_SIGNALING_REPERTOIRE, REACTOME_NFKB_AND_MAP_KINASES_ACTIVATION_MEDIATED_BY_TLR4_SIGNALING_REPERTOIRE ANDIS_ERBB2_BREAST_PRENEOPLASTIC_DN 7, KEGG_ADIPOCYTOKINE_SIGNALING_PATHWAY
	JECHLINGER_EPITHELIAL_TO_MESENCHYMAL_T MODULE_241, MODULE_241 GSE22886_CD8_VS_CD4_NAIVE_TCELL_DN, GSE22	DID_CANCER_CLUSTER_1 IVE_CD4_TCELL_CORD_BLOOD_DN, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN TRANSITION_UP, JECHLINGER_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_UP
	MOREAUX_B_LYMPHOCYTE_MATURATION_BY_TROSS_ACUTE_MYELOID_LEUKEMIA_CBF, ROSS_ASABATES_COLORECTAL_ADENOMA_UP, SABATES GO_COLLAGEN_BINDING, GO_COLLAGEN_BINDING, GO_COLLAGEN_BINDING, APPEL_IMATINIB_IMODULE_291, MODULE_291 OSADA_ASCL1_TARGETS_DN, OSADA_TARGETS_DN, OSADA_TARGETS_DN, OSADA_TARGETS_DN, OSADA_TARGETS_DN, OSADA_TARGETS_DN, OSADA_TARGET	ES_COLORECTAL_ADENOMA_UP DING RESPONSE
	MODULE_410, MODULE_410 BOYLAN_MULTIPLE_MYELOMA_PCA3_UP, BOYL GSE4748_CYANOBACTERIUM_LPSLIKE_VS_LPS_A REACTOME_TRAF6_MEDIATED_INDUCTION_OF_ GO_SECRETORY_GRANULE_MEMBRANE, GO_SEC GSE25123_CTRL_VS_IL4_STIM_PPARG_KO_MACRO	AN_MULTIPLE_MYELOMA_PCA3_UP ND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_DN, GSE4748_CYANOBACTERIUM_LPSLIKE_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_DN NFKB_AND_MAP_KINASES_UPON_TLR7_8_OR_9_ACTIVATION, REACTOME_TRAF6_MEDIATED_INDUCTION_OF_NFKB_AND_MAP_KINASES_UPON_TLR7_8_OR_9_ACTIVATION
	CHUANG_OXIDATIVE_STRESS_RESPONSE_UP, CH GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCE WILENSKY_RESPONSE_TO_DARAPLADIB, WILEN KANNAN_TP53_TARGETS_UP, KANNAN_TP53_TA TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_OPID_SYNDECAN_2	HUANG_OXIDATIVE_STRESS_RESPONSE_UP ELL_52H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_52H_UP ISKY_RESPONSE_TO_DARAPLADIB ARGETS_UP 6HR_DN, TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_6HR_DN
	REACTOME_GLYCOSAMINOGLYCAN_METABOLE OHGUCHI_LIVER_HNF4A_TARGETS_UP, OHGUCE GNF2_ITGAL, GNF2_ITGAL	ING_UP P, GAUSSMANN_MLL_AF4_FUSION_TARGETS_D_UP ISM, REACTOME_GLYCOSAMINOGLYCAN_METABOLISM HI_LIVER_HNF4A_TARGETS_UP
	GSE32423_MEMORY_VS_NAIVE_CD8_TCELL_IL7_I GO_DEFENSE_RESPONSE_TO_FUNGUS, GO_DEFE DELACROIX_RAR_TARGETS_UP, DELACROIX_RA GO_PATTERN_RECOGNITION_RECEPTOR_SIGNA KOBAYASHI_EGFR_SIGNALING_6HR_DN, KOBAY VALK_AML_CLUSTER_5, VALK_AML_CLUSTER_5	R_TARGETS_UP ALING_PATHWAY, GO_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY (ASHI_EGFR_SIGNALING_6HR_DN
	WESTON_VEGFA_TARGETS_3HR, WESTON_VEGF MODULE_96, MODULE_96 BHAT_ESR1_TARGETS_NOT_VIA_AKT1_DN, BHAT YAO_TEMPORAL_RESPONSE_TO_PROGESTERON MODULE_171, MODULE_171	, ABRAHAM_ALPC_VS_MULTIPLE_MYELOMA_UP FA_TARGETS_3HR T_ESR1_TARGETS_NOT_VIA_AKT1_DN TE_CLUSTER_1, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_1
	GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_0.5 CHIARADONNA_NEOPLASTIC_TRANSFORMATIC GO_AMINOGLYCAN_METABOLIC_PROCESS, GO_	AST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_VS_ACT_IN_MAST_CELL_DN 5H_CD4_TCELL_UP, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_0.5H_CD4_TCELL_UP ON_KRAS_CDC25_DN, CHIARADONNA_NEOPLASTIC_TRANSFORMATION_KRAS_CDC25_DN _AMINOGLYCAN_METABOLIC_PROCESS RY_RESPONSE, GO_POSITIVE_REGULATION_OF_INFLAMMATORY_RESPONSE
	LIANG_SILENCED_BY_METHYLATION_2, LIANG_TORCHIA_TARGETS_OF_EWSR1_FLI1_FUSION_TOGO_SPHINGOLIPID_METABOLIC_PROCESS, GO_SGSE22886_NEUTROPHIL_VS_DC_UP, GSE22886_NEGO_INTRINSIC_COMPONENT_OF_EXTERNAL_SILGO_HUMORAL_IMMUNE_RESPONSE, GO_HUMOGO_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS.	SILENCED_BY_METHYLATION_2 DP20_UP, TORCHIA_TARGETS_OF_EWSR1_FLI1_FUSION_TOP20_UP SPHINGOLIPID_METABOLIC_PROCESS EUTROPHIL_VS_DC_UP DE_OF_PLASMA_MEMBRANE, GO_INTRINSIC_COMPONENT_OF_EXTERNAL_SIDE_OF_PLASMA_MEMBRANE DRAL_IMMUNE_RESPONSE CESS, GO_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS
	TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_GO_PROTEOGLYCAN_BINDING, GO_PROTEOGLY	IA_DN, GUTIERREZ_CHRONIC_LYMPHOCYTIC_LEUKEMIA_DN _SUSTAINDED_IN_ERYTHROCYTE_UP, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_SUSTAINDED_IN_ERYTHROCYTE_UP (CAN_BINDING _INVOLVED_IN_IMMUNE_RESPONSE, GO_REGULATION_OF_CYTOKINE_PRODUCTION_INVOLVED_IN_IMMUNE_RESPONSE _UV_RESPONSE_CLUSTER_G2
	GSE22025_TGFB1_VS_TGFB1_AND_PROGESTERONGSE360_L_DONOVANI_VS_L_MAJOR_MAC_DN, G	TION, GO_REGULATION_OF_INTERLEUKIN_6_PRODUCTION T_ALPHA_GRANULE _CYTOTOXIC_MODULE
	GSE25502_WT_VS_KLF13_KO_THYMIC_MEMORY_ PID_AVB3_OPN_PATHWAY, PID_AVB3_OPN_PAT	LIKE_CD8_TCELL_DN, GSE25502_WT_VS_KLF13_KO_THYMIC_MEMORY_LIKE_CD8_TCELL_DN THWAY N, GSE6269_FLU_VS_STREP_PNEUMO_INF_PBMC_DN UKOCYTE_MEDIATED_IMMUNITY
	NUMATA_CSF3_SIGNALING_VIA_STAT3, NUMATA ATM_DN.V1_UP, ATM_DN.V1_UP MCDOWELL_ACUTE_LUNG_INJURY_UP, MCDOWGO_POSITIVE_REGULATION_OF_TRANSCRIPTIONGSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_N	FA_CSF3_SIGNALING_VIA_STAT3 VELL_ACUTE_LUNG_INJURY_UP N_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_STRESS, GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONS MYD88_KO_MACROPHAGE_DN, GSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN E_PRODUCTION, GO_REGULATION_OF_MACROPHAGE_CYTOKINE_PRODUCTION
	DAVIES_MULTIPLE_MYELOMA_VS_MGUS_DN, DEGENTILE_UV_LOW_DOSE_DN, GENTILE_UV_LOW_CHEN_ETV5_TARGETS_SERTOLI, CHEN_ETV5_TAGSE27859_MACROPHAGE_VS_DC_DN, GSE27859_NGO_REGULATION_OF_CELLULAR_EXTRAVASATEWORSCHECH_TUMOR_REJECTION_UP, WORSCHE	AVIES_MULTIPLE_MYELOMA_VS_MGUS_DN V_DOSE_DN \RGETS_SERTOLI MACROPHAGE_VS_DC_DN ION, GO_REGULATION_OF_CELLULAR_EXTRAVASATION
	GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_F GNF2_RAB7L1, GNF2_RAB7L1 GERHOLD_ADIPOGENESIS_UP, GERHOLD_ADIPO BENNETT_SYSTEMIC_LUPUS_ERYTHEMATOSUS, GO_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC GO_NEGATIVE_REGULATION_OF_EPITHELIAL_C	POST_CORONAVIRUS_BRAIN_UP, GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_POST_CORONAVIRUS_BRAIN_UP DGENESIS_UP BENNETT_SYSTEMIC_LUPUS_ERYTHEMATOSUS C_PROCESS, GO_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS CELL_APOPTOTIC_PROCESS, GO_NEGATIVE_REGULATION_OF_EPITHELIAL_CELL_APOPTOTIC_PROCESS
	GO_ACUTE_PHASE_RESPONSE, GO_ACUTE_PHASE GO_ANCHORED_COMPONENT_OF_MEMBRANE, GO_RESPONSE_TO_FUNGUS, GO_RESPONSE_TO_ GO_MONOCYTE_CHEMOTAXIS, GO_MONOCYTE_ NIELSEN_GIST_AND_SYNOVIAL_SARCOMA_DN, LINDVALL_IMMORTALIZED_BY_TERT_DN, LINDVALL_IMMORTALIZED_BY_TERT_DN, LINDVALL_SARCOME_G_ALPHA_I_SIGNALLING_EVENTS, I	GO_ANCHORED_COMPONENT_OF_MEMBRANE FUNGUS _CHEMOTAXIS NIELSEN_GIST_AND_SYNOVIAL_SARCOMA_DN VALL_IMMORTALIZED_BY_TERT_DN REACTOME_G_ALPHA_I_SIGNALLING_EVENTS
	MODULE_208, MODULE_208 MANTOVANI_NFKB_TARGETS_UP, MANTOVANI	EETS_UP AMASHITA_LIVER_CANCER_STEM_CELL_UP IULTIPLE_MYELOMA_CD1_UP ONE_MARROW_UP, GSE29949_MICROGLIA_BRAIN_VS_MONOCYTE_BONE_MARROW_UP
	CHIANG_LIVER_CANCER_SUBCLASS_UNANNOT DCA_UP.V1_UP, DCA_UP.V1_UP GNF2_SPI1, GNF2_SPI1	
	GO_INTERLEUKIN_1_MEDIATED_SIGNALING_PA	PATHWAY 1_NEIGHBORS IHWAY 12H_ACT_CD4_TCELL_DN, GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_12H_ACT_CD4_TCELL_DN ATHWAY, GO_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY
	GO_UNSATURATED_FATTY_ACID_METABOLIC_I LEE_EARLY_T_LYMPHOCYTE_DN, LEE_EARLY_T_ SAMOLS_TARGETS_OF_KHSV_MIRNAS_DN, SAMO	UM_RESPONSE_40_MCF10A JLAN_UV_RESPONSE_NORMAL_DN C_PROCESS, GO_REACTIVE_NITROGEN_SPECIES_METABOLIC_PROCESS PROCESS, GO_UNSATURATED_FATTY_ACID_METABOLIC_PROCESS _LYMPHOCYTE_DN OLS_TARGETS_OF_KHSV_MIRNAS_DN
	WANG_CLASSIC_ADIPOGENIC_TARGETS_OF_PPA	RESPONSE_TO_INTERFERON_GAMMA LONG_CHAIN_FATTY_ACID_BINDING
	BROWNE_HCMV_INFECTION_1HR_UP, BROWNE_IL15_UP.V1_DN, IL15_UP.V1_DN GO_MAP_KINASE_PHOSPHATASE_ACTIVITY, GO GO_PLATELET_ALPHA_GRANULE_MEMBRANE, G BIOCARTA_PPARA_PATHWAY, BIOCARTA_PPAR GSE13306_RA_VS_UNTREATED_TREG_DN, GSE133	D_MAP_KINASE_PHOSPHATASE_ACTIVITY GO_PLATELET_ALPHA_GRANULE_MEMBRANE RA_PATHWAY B06_RA_VS_UNTREATED_TREG_DN
	MARTINELLI_IMMATURE_NEUTROPHIL_DN, MAGSE1460_DP_THYMOCYTE_VS_NAIVE_CD4_TCELTGO_ANCHORED_COMPONENT_OF_EXTERNAL_SMCBRYAN_PUBERTAL_BREAST_6_7WK_DN, MCBAKT_UP.V1_DN, AKT_UP.V1_DN RASHI_RESPONSE_TO_IONIZING_RADIATION_1, AMIT_EGF_RESPONSE_40_HELA, AMIT_EGF_RESPONSE_40_H	RETINELLI_IMMATURE_NEUTROPHIL_DN L_CORD_BLOOD_DN, GSE1460_DP_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN SIDE_OF_PLASMA_MEMBRANE, GO_ANCHORED_COMPONENT_OF_EXTERNAL_SIDE_OF_PLASMA_MEMBRANE SRYAN_PUBERTAL_BREAST_6_7WK_DN RASHI_RESPONSE_TO_IONIZING_RADIATION_1 PONSE_40_HELA
	GSE15659_NAIVE_VS_PTPRC_NEG_CD4_TCELL_UIGRAHAM_CML_QUIESCENT_VS_NORMAL_QUIESCENT_VS_NORMAL_QUIESCENT_VS_NORMAL_QUIESCENT_VS_NORMAL_QUIESCENT_VS_NOTER_CDH1_TAL_ONDER_CDH1_TARGETS_3_UP, ONDER_CDH1_TAL_NAKAMURA_METASTASIS_MODEL_DN, NAKAMURA_METASTASIS_MODEL_DN, NAKAMURA_REGULATION_OF_RESPONSE_TO_VBOHN_PRIMARY_IMMUNODEFICIENCY_SYNDRO	P, GSE15659_NAIVE_VS_PTPRC_NEG_CD4_TCELL_UP SCENT_DN, GRAHAM_CML_QUIESCENT_VS_NORMAL_QUIESCENT_DN , LINDSTEDT_DENDRITIC_CELL_MATURATION_A ARGETS_3_UP IURA_METASTASIS_MODEL_DN WOUNDING, GO_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING DM_DN, BOHN_PRIMARY_IMMUNODEFICIENCY_SYNDROM_DN
	GO_REGULATION_OF_INTERLEUKIN_12_PRODUCTION GO_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GO_LAMININ_BINDING, GO_LAMININ_BINDING GO_POSITIVE_REGULATION_OF_CHEMOTAXIS, COUNTY GNF2_PTX3, GNF2_PTX3 CADWELL_ATG16L1_TARGETS_UP, CADWELL_ATGO_POSITIVE_REGULATION_OF_BLOOD_CIRCU	CTION, GO_REGULATION_OF_INTERLEUKIN_12_PRODUCTION GO_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS GO_POSITIVE_REGULATION_OF_CHEMOTAXIS TG16L1_TARGETS_UP LATION, GO_POSITIVE_REGULATION_OF_BLOOD_CIRCULATION
	DELLA_RESPONSE_TO_TSA_AND_BUTYRATE, DEGO_CYTOPLASMIC_SEQUESTERING_OF_TRANSCEBIOCARTA_LYM_PATHWAY, BIOCARTA_LYM_PAKUMAMOTO_RESPONSE_TO_NUTLIN_3A_UP, KUDORN_ADENOVIRUS_INFECTION_48HR_DN, DOIVISALA_AGING_LYMPHOCYTE_DN, VISALA_AGING_LYMPHOCYTE_DN, VISALA_LYMPHOCYTE_DN, VISALA_LYMPHOCYT	ELLA_RESPONSE_TO_TSA_AND_BUTYRATE CRIPTION_FACTOR, GO_CYTOPLASMIC_SEQUESTERING_OF_TRANSCRIPTION_FACTOR ATHWAY JMAMOTO_RESPONSE_TO_NUTLIN_3A_UP RN_ADENOVIRUS_INFECTION_48HR_DN
	HEDVAT_ELF4_TARGETS_UP, HEDVAT_ELF4_TARGO_AMINOGLYCAN_CATABOLIC_PROCESS, GO_GO_POSITIVE_REGULATION_OF_EXOCYTOSIS, GO_REGULATION_OF_G_PROTEIN_COUPLED_REGO_HYALURONIC_ACID_BINDING, GO_HYALURGO_LOW_DENSITY_LIPOPROTEIN_PARTICLE, GO	RGETS_UP _AMINOGLYCAN_CATABOLIC_PROCESS O_POSITIVE_REGULATION_OF_EXOCYTOSIS ECEPTOR_PROTEIN_SIGNALING_PATHWAY, GO_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_PROTEIN_SIGNALING_PATHWAY RONIC_ACID_BINDING
	TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_CLU_TUMOR_ANGIOGENESIS_UP, LU_TUMOR_ANGO_PHOSPHOLIPASE_C_ACTIVITY, GO_PHOSPHOGILDEA_METASTASIS, GILDEA_METASTASISGO_NUCLEOSIDE_SALVAGE, GO_NUCLEOSIDE_SALVAGE	3D_DN, TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_3D_DN IGIOGENESIS_UP OLIPASE_C_ACTIVITY SALVAGE IMUNE_RESPONSE, GO_CYTOKINE_PRODUCTION_INVOLVED_IN_IMMUNE_RESPONSE
	PASINI_SUZ12_TARGETS_UP, PASINI_SUZ12_TARGETS_UP, PASINI_SUZ12_TARGETS_UP, PASINI_SUZ12_TARGETS_UP.V1_UP, MAHARE PDGF_UP.V1_DN, PDGF_UP.V1_DN GO_BASIC_AMINO_ACID_TRANSPORT, GO_BASIC_YAMASHITA_METHYLATED_IN_PROSTATE_CANGET MODULE_407, MODULE_407 JNK_DN.V1_DN, JNK_DN.V1_DN	GETS_UP .DEVAN_RESPONSE_TO_MP470_UP C_AMINO_ACID_TRANSPORT ICER, YAMASHITA_METHYLATED_IN_PROSTATE_CANCER
	HOFFMAN_CLOCK_TARGETS_UP, HOFFMAN_CLAPID_FRA_PATHWAY, PID_FRA_PATHWAY GO_PENTOSE_METABOLIC_PROCESS, GO_PENTO chr4q22, chr4q22 FOURNIER_ACINAR_DEVELOPMENT_EARLY_UP, GO_PYRIMIDINE_NUCLEOSIDE_CATABOLIC_PRO GO_IMMUNOGLOBULIN_BINDING, GO_IMMUNO	OSE_METABOLIC_PROCESS , FOURNIER_ACINAR_DEVELOPMENT_EARLY_UP OCESS, GO_PYRIMIDINE_NUCLEOSIDE_CATABOLIC_PROCESS
	MODULE_375, MODULE_375 MOROSETTI_FACIOSCAPULOHUMERAL_MUSCUL BIOCARTA_EPONFKB_PATHWAY, BIOCARTA_EPO	ARK_BILE_ACID_METABOLISM _OXYGEN_SPECIES, GO_REGULATION_OF_RESPONSE_TO_REACTIVE_OXYGEN_SPECIES LAR_DISTROPHY_DN, MOROSETTI_FACIOSCAPULOHUMERAL_MUSCULAR_DISTROPHY_DN ONFKB_PATHWAY
	GO_FATTY_ACID_BINDING, GO_FATTY_ACID_BINGSE19923_E2A_KO_VS_HEB_AND_E2A_KO_DP_THEVERRECCHIA_RESPONSE_TO_TGFB1_C3, VERRECTED RIGGINS_TAMOXIFEN_RESISTANCE_UP, RIGGINS_GSE29949_CD8_NEG_DC_SPLEEN_VS_CD8_POS_DGO_FUCOSE_METABOLIC_PROCESS, GO_FUCOSE_MEINHOLD_OVARIAN_CANCER_LOW_GRADE_U	NDING HYMOCYTE_DN, GSE19923_E2A_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_DN CHIA_RESPONSE_TO_TGFB1_C3 S_TAMOXIFEN_RESISTANCE_UP C_SPLEEN_UP, GSE29949_CD8_NEG_DC_SPLEEN_VS_CD8_POS_DC_SPLEEN_UP S_METABOLIC_PROCESS JP, MEINHOLD_OVARIAN_CANCER_LOW_GRADE_UP
	GO_NEGATIVE_REGULATION_OF_IMMUNE_RESIGN_LONG_CHAIN_FATTY_ACID_METABOLIC_PRESIGNED TO	PONSE, GO_NEGATIVE_REGULATION_OF_IMMUNE_RESPONSE ROCESS, GO_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS 2_AND_IL17A_SIGNALING PROCESS, GO_MULTICELLULAR_ORGANISM_METABOLIC_PROCESS BDULRAHMAN_KIDNEY_CANCER_VHL_DN
		IN_METABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_GLYCOPROTEIN_METABOLIC_PROCESS