

ORY_BCELL_VS_PLASMA_CELL_UP, GSE13411_SWITCHED_MEMORY_BCELL_VS_PLASMA_CELL_UP

GSE17301_CTRL_VS_48H_IFNA2_STIM_CD8_TCELL_UP, GSE17301_CTRL_VS_48H_IFNA2_STIM_CD8_TCELL_UP
GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_UP
GSE13411_PLASMA_CELL_VS_MEMORY_BCELL_DN, GSE13411_PLASMA_CELL_VS_MEMORY_BCELL_DN
GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_DN, GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_DN
GSE17301_IFNA2_VS_IFNA5_STIM_ACD3_ACD28_ACT_CD8_TCELL_UP, GSE17301_IFNA2_VS_IFNA5_STIM_ACD3_ACD28_ACT_CD8_TCELL_UP
GSE13411_IGM_MEMORY_BCELL_VS_PLASMA_CELL_UP, GSE13411_IGM_MEMORY_BCELL_VS_PLASMA_CELL_UP
GSE17974_0.5H_VS_72H_IL4_AND_ANTIL_IL12_ACT_CD4_TCELL_UP, GSE17974_0.5H_VS_72H_IL4_AND_ANTIL_IL12_ACT_CD4_TCELL_UP
GSE40273_EOS_KO_VS_WT_TREG_DN, GSE40273_EOS_KO_VS_WT_TREG_DN
GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN
GSE34515_CD16_POS_MONOCYTE_VS_DC_DN, GSE34515_CD16_POS_MONOCYTE_VS_DC_DN
GSE12366_GC_VS_NAIVE_BCELL_DN, GSE12366_GC_VS_NAIVE_BCELL_DN
GSE6259_FLT3L_INDUCED_VS_WT_SPLENIC_DC_33D1_POS_UP, GSE6259_FLT3L_INDUCED_VS_WT_SPLENIC_DC_33D1_POS_UP
GSE29618_BCELL_VS_PDC_UP, GSE29618_BCELL_VS_PDC_UP
GSE20727_CTRL_VS_H2O2_TREATED_DC_DN, GSE20727_CTRL_VS_H2O2_TREATED_DC_DN
GSE41867_MEMORY_VS_EXHAUSTED_CD8_TCELL_DAY30_LCMV_UP, GSE41867_MEMORY_VS_EXHAUSTED_CD8_TCELL_DAY30_LCMV_UP
GSE13484_12H_VS_3H_YF17D_VACCINE_STIM_PBMG_UP, GSE13484_12H_VS_3H_YF17D_VACCINE_STIM_PBMG_UP
GSE24574_BCL6_HIGH_VS_LOW_TFH_CD4_TCELL_UP, GSE24574_BCL6_HIGH_VS_LOW_TFH_CD4_TCELL_UP
GSE33425_CD161_HIGH_VS_INT_CD8_TCELL_UP, GSE33425_CD161_HIGH_VS_INT_CD8_TCELL_UP
GSE39556_CD8A_DC_VS_NK_CELL_DN, GSE39556_CD8A_DC_VS_NK_CELL_DN
GSE2770_UNTREATED_VS_ACT_CD4_TCELL_2H_UP, GSE2770_UNTREATED_VS_ACT_CD4_TCELL_2H_UP
GSE22935_WT_VS_MYD88_KO_MACROPHAGE_48H_MBOVIS_BCG_STIM_DN, GSE22935_WT_VS_MYD88_KO_MACROPHAGE_48H_MBOVIS_BCG_STIM_DN
GSE22886_NAIVE_BCELL_VS_BM_PLASMA_CELL_UP, GSE22886_NAIVE_BCELL_VS_BM_PLASMA_CELL_UP
GSE29618_BCELL_VS_PDC_DAY7_FLU_VACCINE_UP, GSE29618_BCELL_VS_PDC_DAY7_FLU_VACCINE_UP
GSE20727_CTRL_VS_H2O2_TREATED_DC_UP, GSE20727_CTRL_VS_H2O2_TREATED_DC_UP
GSE33374_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_DN, GSE33374_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_DN
GSE36527_CD69_NEG_VS_POS_TREG_CD62L_LOS_KLRG1_NEG_UP, GSE36527_CD69_NEG_VS_POS_TREG_CD62L_LOS_KLRG1_NEG_UP
GSE15930_STIM_VS_STIM_AND_IL12_48H_CD8_T_CELL_UP, GSE15930_STIM_VS_STIM_AND_IL12_48H_CD8_T_CELL_UP
GSE13411_NAIVE_BCELL_VS_PLASMA_CELL_UP, GSE13411_NAIVE_BCELL_VS_PLASMA_CELL_UP
GSE29618_BCELL_VS_MONOCYTE_UP, GSE29618_BCELL_VS_MONOCYTE_UP
GSE23568_ID3_KO_VS_WT_CD8_TCELL_DN, GSE23568_ID3_KO_VS_WT_CD8_TCELL_DN
GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN, GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN
GSE3982_BCELL_VS_EFF_MEMORY_CD4_TCELL_UP, GSE3982_BCELL_VS_EFF_MEMORY_CD4_TCELL_UP
GSE3565_CTRL_VS_LPS_INJECTED_DUSP1_KO_SPLENOCYTES_DN, GSE3565_CTRL_VS_LPS_INJECTED_DUSP1_KO_SPLENOCYTES_DN
GSE17721_PAM3CSK4_VS_GADIQUIMOD_8H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_8H_BMDC_DN
GSE3982_MEMORY_CD4_TCELL_VS_BCELL_DN, GSE3982_MEMORY_CD4_TCELL_VS_BCELL_DN
GSE10500_ARTHRITIC_SYNOVIAL_FLUID_VS_HEALTHY_MACROPHAGE_DN, GSE10500_ARTHRITIC_SYNOVIAL_FLUID_VS_HEALTHY_MACROPHAGE_DN
GSE3565_DUSP1_VS_WT_SPLENOCYTES_DN, GSE3565_DUSP1_VS_WT_SPLENOCYTES_DN
GSE10325_LUPUS_CD4_TCELL_VS_LUPUS_BCELL_DN, GSE10325_LUPUS_CD4_TCELL_VS_LUPUS_BCELL_DN
TAATGTG_MIR323, TAATGTG_MIR323
GSE3982_DC_VS_BASOPHIL_DN, GSE3982_DC_VS_BASOPHIL_DN
GSE17721_PAM3CSK4_VS_CPG_6H_BMDC_DN, GSE17721_PAM3CSK4_VS_CPG_6H_BMDC_DN
GSE17721_PAM3CSK4_VS_GADIQUIMOD_4H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_4H_BMDC_DN
EPPERT_HSC_R, EPPERT_HSC_R
GSE34156_UNTREATED_VS_6H_NOD2_LIGAND_TREATED_MONOCYTE_DN, GSE34156_UNTREATED_VS_6H_NOD2_LIGAND_TREATED_MONOCYTE_DN
GSE10325_CD4_TCELL_VS_BCELL_DN, GSE10325_CD4_TCELL_VS_BCELL_DN
GSE9006_HEALTHY_VS_TYPE_2_DIABETES_PBMG_AT_DX_DN, GSE9006_HEALTHY_VS_TYPE_2_DIABETES_PBMG_AT_DX_DN
GSE29164_DAY3_VS_DAY7_UNTREATED_MELANOMA_UP, GSE29164_DAY3_VS_DAY7_UNTREATED_MELANOMA_UP
GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP, GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP
GSE41176_UNSTIM_VS_ANTL_IGM_STIM_BCELL_6H_DN, GSE41176_UNSTIM_VS_ANTL_IGM_STIM_BCELL_6H_DN
GSE3982_MAST_CELL_VS_BCELL_DN, GSE3982_MAST_CELL_VS_BCELL_DN
GSE17721_LPS_VS_PAM3CSK4_4H_BMDC_UP, GSE17721_LPS_VS_PAM3CSK4_4H_BMDC_UP
RASHL_RESPONSE_TO_IONIZING_RADIATION_5, RASHL_RESPONSE_TO_IONIZING_RADIATION_5
GSE10463_CD40L_AND_VA347_VS_CD40L_IN_DC_UP, GSE10463_CD40L_AND_VA347_VS_CD40L_IN_DC_UP
GSE30971_WBP7_HET_VS_KO_MACROPHAGE_DN, GSE30971_WBP7_HET_VS_KO_MACROPHAGE_DN
GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP, GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP
PID_PDGRFB_PATHWAY, PID_PDGRFB_PATHWAY
GSE17974_0H_VS_1H_IN_VITRO_ACT_CD4_TCELL_UP, GSE17974_0H_VS_1H_IN_VITRO_ACT_CD4_TCELL_UP
MODULE_292, MODULE_292
GSE3982_EOSINOPHIL_VS_TH2_UP, GSE3982_EOSINOPHIL_VS_TH2_UP
GSE13946_CTRL_VS_DSS_COLITIS_GD_TCELL_FROM_COLON_DN, GSE13946_CTRL_VS_DSS_COLITIS_GD_TCELL_FROM_COLON_DN
REACTOME_ACTIVATED_TLR4_SIGNALLING, REACTOME_ACTIVATED_TLR4_SIGNALLING
GSE22886_NAIVE_BCELL_VS_MONOCYTE_UP, GSE22886_NAIVE_BCELL_VS_MONOCYTE_UP
KEGG_ERBB_SIGNALING_PATHWAY, KEGG_ERBB_SIGNALING_PATHWAY
AMUNDSON_POOR_SURVIVAL_AFTER_GAMMA_RADIATION_8G, AMUNDSON_POOR_SURVIVAL_AFTER_GAMMA_RADIATION_8G
GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_UP, GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_UP
GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_UP, GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_UP
GO_POSITIVE_REGULATION_OF_HEMOPOIESIS, GO_POSITIVE_REGULATION_OF_HEMOPOIESIS
GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_UP, GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_UP
DACOSTA_UV_RESPONSE_VIA_ERCC3_XPCS_DN, DACOSTA_UV_RESPONSE_VIA_ERCC3_XPCS_DN
GTGCAA_MIR507, GTGCAA_MIR507
GSE3982_BCELL_VS_TH1_UP, GSE3982_BCELL_VS_TH1_UP
KEGG_CHEMOKINE_SIGNALING_PATHWAY, KEGG_CHEMOKINE_SIGNALING_PATHWAY
GSE3982_MAC_VS_NKCELL_DN, GSE3982_MAC_VS_NKCELL_DN
PID_ERBB1_INTERNALIZATION_PATHWAY, PID_ERBB1_INTERNALIZATION_PATHWAY
GSE21063_CTRL_VS_ANTL_IGM_STIM_BCELL_3H_UP, GSE21063_CTRL_VS_ANTL_IGM_STIM_BCELL_3H_UP
GO_TRANS_GOLGI_NETWORK_MEMBRANE, GO_TRANS_GOLGI_NETWORK_MEMBRANE
MORL_LARGE_PRE_BIL_LYMPHOCYTE_DN, MORL_LARGE_PRE_BIL_LYMPHOCYTE_DN
GSE28737_WT_VS_BCL6_KO_FOLLICULAR_BCELL_UP, GSE28737_WT_VS_BCL6_KO_FOLLICULAR_BCELL_UP
GO_RESPONSE_TO_REACTIVE_OXYGEN_SPECIES, GO_RESPONSE_TO_REACTIVE_OXYGEN_SPECIES
GSE37605_NOD_VS_C57BL6_IRES_GFP_TREG_UP, GSE37605_NOD_VS_C57BL6_IRES_GFP_TREG_UP
GSE19941_UNSTIM_VS_LPS_STIM_IL10_KO_MACROPHAGE_UP, GSE19941_UNSTIM_VS_LPS_STIM_IL10_KO_MACROPHAGE_UP
GSE5542_UNTREATED_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP, GSE5542_UNTREATED_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP
NUTT_GBM_VS_AO_GLIOMA_UP, NUTT_GBM_VS_AO_GLIOMA_UP
GO_POSITIVE_REGULATION_OF_LEUKOCYTE_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_LEUKOCYTE_DIFFERENTIATION
PID_FAK_PATHWAY, PID_FAK_PATHWAY
PID_TRAIL_PATHWAY, PID_TRAIL_PATHWAY
GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP
GSE1460_DP_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN, GSE1460_DP_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN
GO_HISTONE_H3_K9_MODIFICATION, GO_HISTONE_H3_K9_MODIFICATION
GSE37605_NOD_VS_C57BL6_IRES_GFP_TREG_DN, GSE37605_NOD_VS_C57BL6_IRES_GFP_TREG_DN
MODULE_411, MODULE_411
GSE22886_TCELL_VS_BCELL_NAIVE_DN, GSE22886_TCELL_VS_BCELL_NAIVE_DN
BIOCARTA_MET_PATHWAY, BIOCARTA_MET_PATHWAY
GO_REGULATION_OF_T_CELL_DIFFERENTIATION, GO_REGULATION_OF_T_CELL_DIFFERENTIATION
PID_FCR1_PATHWAY, PID_FCR1_PATHWAY
GO_POSITIVE_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION
GO_CYTOPLASMIC_SIDE_OF_MEMBRANE, GO_CYTOPLASMIC_SIDE_OF_MEMBRANE
GO_REGULATION_OF_CELL_MATRIX_ADHESION, GO_REGULATION_OF_CELL_MATRIX_ADHESION
PID_ERBB1_RECEPTOR_PROXIMAL_PATHWAY, PID_ERBB1_RECEPTOR_PROXIMAL_PATHWAY
PID_NETRIN_PATHWAY, PID_NETRIN_PATHWAY
GO_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING, GO_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING
TTGGAGA_MIR5155P_MIR519E, TTGGAGA_MIR5155P_MIR519E
PID_EPHB_FWD_PATHWAY, PID_EPHB_FWD_PATHWAY
GSE37605_FOXP3_FUSION_GFP_VS_IRES_GFP_TREG_C57BL6_DN, GSE37605_FOXP3_FUSION_GFP_VS_IRES_GFP_TREG_C57BL6_DN
GCM_PTFRD, GCM_PTFRD
PID_P75_NTR_PATHWAY, PID_P75_NTR_PATHWAY
PID_IL2_P13K_PATHWAY, PID_IL2_P13K_PATHWAY
GO_INSULIN LIKE_GROWTH_FACTOR_RECEPTOR_BINDING, GO_INSULIN LIKE_GROWTH_FACTOR_RECEPTOR_BINDING
GO_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION, GO_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION
PID_LYSOPHOSPHOLIPID_PATHWAY, PID_LYSOPHOSPHOLIPID_PATHWAY
GAZDA_DIAMOND_BLACKFAN_ANEMIA_MYELOID_UP, GAZDA_DIAMOND_BLACKFAN_ANEMIA_MYELOID_UP
GSE16266_LPS_VS_HEATSHOCK_AND_LPS_STIM_MEF_UP, GSE16266_LPS_VS_HEATSHOCK_AND_LPS_STIM_MEF_UP
BIOCARTA_FCR1_PATHWAY, BIOCARTA_FCR1_PATHWAY
PID_AR_NONGENOMIC_PATHWAY, PID_AR_NONGENOMIC_PATHWAY
ATGTCAC_MIR489, ATGTCAC_MIR489
GO_PROTEIN_TYROSINE_KINASE_ACTIVITY, GO_PROTEIN_TYROSINE_KINASE_ACTIVITY
REACTOME_NEF_MEDIATES_DOWN_MODULATION_OF_CELL_SURFACE_RECEPTORS_BY_RECRUITING_THEM_TO_CLATHRIN_ADAPTERS, REACTOME_NEF_MEDIATES_DOWN_MODULATION_OF_CELL_SURFACE_RECEPTORS_BY_RECRUITING_THEM_TO_CLATHRIN_ADAPTERS
BIOCARTA_TFF_PATHWAY, BIOCARTA_TFF_PATHWAY
GO_PEPTIDYL_TYROSINE_MODIFICATION, GO_PEPTIDYL_TYROSINE_MODIFICATION
GO_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION, GO_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION
GO_POSITIVE_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION
KEGG_PRIMARY_IMMUNODEFICIENCY, KEGG_PRIMARY_IMMUNODEFICIENCY
GO_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE, GO_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE
REACTOME_REGULATION_OF_SIGNALING_BY_CBL, REACTOME_REGULATION_OF_SIGNALING_BY_CBL
GO_POSITIVE_REGULATION_OF_B_CELL_ACTIVATION, GO_POSITIVE_REGULATION_OF_B_CELL_ACTIVATION
GO_POSITIVE_REGULATION_OF_MESENCHYMAL_CELL_PROLIFERATION, GO_POSITIVE_REGULATION_OF_MESENCHYMAL_CELL_PROLIFERATION
GO_REGULATION_OF_GOLGI_ORGANIZATION, GO_REGULATION_OF_GOLGI_ORGANIZATION
GO_IMMUNOLOGICAL_SYNAPSE, GO_IMMUNOLOGICAL_SYNAPSE
MODULE_511, MODULE_511
GO_NEGATIVE_REGULATION_OF_INTERLEUKIN_8_PRODUCTION, GO_NEGATIVE_REGULATION_OF_INTERLEUKIN_8_PRODUCTION
GO_CELLULAR_RESPONSE_TO_FLUID_SHEAR_STRESS, GO_CELLULAR_RESPONSE_TO_FLUID_SHEAR_STRESS
KEGG_DORSO_VENTRAL_AXIS_FORMATION, KEGG_DORSO_VENTRAL_AXIS_FORMATION
GO_MITOGEN_ACTIVATED_PROTEIN_KINASE_KINASE_KINASE_BINDING, GO_MITOGEN_ACTIVATED_PROTEIN_KINASE_KINASE_KINASE_BINDING
GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGM_BCELL_DN, GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGM_BCELL_DN
GSE3982_BASOPHIL_VS_NKCELL_DN, GSE3982_BASOPHIL_VS_NKCELL_DN
ST_GA12_PATHWAY, ST_GA12_PATHWAY
GO_REGULATION_OF_EPITHELIAL_CELL_MIGRATION, GO_REGULATION_OF_EPITHELIAL_CELL_MIGRATION
GO_PHAGOCYTIC_CUP, GO_PHAGOCYTIC_CUP
FINETTL_BREAST_CANCER_KINOME_GREEN, FINETTL_BREAST_CANCER_KINOME_GREEN
GO_REGULATION_OF_INTERLEUKIN_17_PRODUCTION, GO_REGULATION_OF_INTERLEUKIN_17_PRODUCTION
ST_ERK1_ERK2_MAPK_PATHWAY, ST_ERK1_ERK2_MAPK_PATHWAY
BIOCARTA_IGFIR_PATHWAY, BIOCARTA_IGFIR_PATHWAY
BIOCARTA_BAD_PATHWAY, BIOCARTA_BAD_PATHWAY
GO_ERBB_SIGNALING_PATHWAY, GO_ERBB_SIGNALING_PATHWAY
PID_EPO_PATHWAY, PID_EPO_PATHWAY
GO_POSITIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GO_POSITIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
PID_PTP1B_PATHWAY, PID_PTP1B_PATHWAY
GO_EPIDERMAL_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GO_EPIDERMAL_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
GO_BARBED_END_ACTIN_FILAMENT_CAPPING, GO_BARBED_END_ACTIN_FILAMENT_CAPPING
GO_STAT_CASCADE, GO_STAT_CASCADE
GO_PEPTIDYL_TYROSINE_AUTOPHOSPHORYLATION, GO_PEPTIDYL_TYROSINE_AUTOPHOSPHORYLATION
GO_MUSCLE_CELL_MIGRATION, GO_MUSCLE_CELL_MIGRATION
GO_CELLULAR_RESPONSE_TO_GROWTH_HORMONE_STIMULUS, GO_CELLULAR_RESPONSE_TO_GROWTH_HORMONE_STIMULUS
GO_THYROID_GLAND_DEVELOPMENT, GO_THYROID_GLAND_DEVELOPMENT
GO_PHOSPHATIDYLINOSITOL_3_KINASE_BINDING, GO_PHOSPHATIDYLINOSITOL_3_KINASE_BINDING
GO_POSITIVE_REGULATION_OF_INTERLEUKIN_17_PRODUCTION, GO_POSITIVE_REGULATION_OF_INTERLEUKIN_17_PRODUCTION
GO_POSITIVE_REGULATION_OF_T_HELPER_CELL_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_T_HELPER_CELL_DIFFERENTIATION
BUDHU_LIVER_CANCER_METASTASIS_UP, BUDHU_LIVER_CANCER_METASTASIS_UP
BIOCARTA_HER2_PATHWAY, BIOCARTA_HER2_PATHWAY
PID_EPHA2_FWD_PATHWAY, PID_EPHA2_FWD_PATHWAY
GO_ENDOTHELIAL_CELL_MIGRATION, GO_ENDOTHELIAL_CELL_MIGRATION
GO_POSITIVE_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE, GO_POSITIVE_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE
GO_CANONICAL_WNT_SIGNALING_PATHWAY, GO_CANONICAL_WNT_SIGNALING_PATHWAY
GO_IONOTROPIC_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY, GO_IONOTROPIC_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY