```
GSE3400 UNTREATED VS IFNB TREATED MEF UP, GSE3400 UNTREATED VS IFNB TREATED MEF UP
  GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_UP
  GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_DN, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_DN
  GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN
  GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_UP
  GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_UP, GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_UP
  GSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_UP, GSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_UP
  GSE411_UNSTIM_VS_100MIN_IL6_STIM_MACROPHAGE_UP, GSE411_UNSTIM_VS_100MIN_IL6_STIM_MACROPHAGE_UP
  GSE24726_WT_VS_E2_2_KO_PDC_DN, GSE24726_WT_VS_E2_2_KO_PDC_DN
  REACTOME_AUTOPHAGY, REACTOME_AUTOPHAGY
  GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN, GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN
  GSE5960_TH1_VS_ANERGIC_TH1_DN, GSE5960_TH1_VS_ANERGIC_TH1_DN
  GSE17721_12H_VS_24H_POLYIC_BMDC_UP, GSE17721_12H_VS_24H_POLYIC_BMDC_UP
  GSE11961_MARGINAL_ZONE_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_DN, GSE11961_MARGINAL_ZONE_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_DN
  GSE339 EX_VIVO_VS_IN_CULTURE_CD4POS_DC_DN, GSE339 EX_VIVO_VS_IN_CULTURE_CD4POS_DC_DN
  GSE27434_WT_VS_DNMT1_KO_TREG_DN, GSE27434_WT_VS_DNMT1_KO_TREG_DN
  WHITFIELD_CELL_CYCLE_M_G1, WHITFIELD_CELL_CYCLE_M_G1
  CAMP_UP.V1_DN, CAMP_UP.V1_DN
  GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_UP
  GSE5542_UNTREATED_VS_IFNA_TREATED_EPITHELIAL_CELLS_6H_UP, GSE5542_UNTREATED_VS_IFNA_TREATED_EPITHELIAL_CELLS_6H_UP
  GSE15735_2H_VS_12H_HDAC_INHIBITOR_TREATED_CD4_TCELL_UP, GSE15735_2H_VS_12H_HDAC_INHIBITOR_TREATED_CD4_TCELL_UP
  GSE339_CD8POS_VS_CD4CD8DN_DC_DN, GSE339_CD8POS_VS_CD4CD8DN_DC_DN
  GSE11864_CSF1_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN, GSE11864_CSF1_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN
  GSE42021_CD24HI_TREG_VS_CD24HI_TCONV_THYMUS_DN, GSE42021_CD24HI_TREG_VS_CD24HI_TCONV_THYMUS_DN
 GSE11864_CSF1_VS_CSF1_PAM3CYS_IN_MAC_DN, GSE11864_CSF1_VS_CSF1_PAM3CYS_IN_MAC_DN
  AIZARANI_LIVER_C39_EPCAM_POS_BILE_DUCT_CELLS_4, AIZARANI_LIVER_C39_EPCAM_POS_BILE_DUCT_CELLS_4
  GSE11864_CSF1_IFNG_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN, GSE11864_CSF1_IFNG_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN
  LANDIS_ERBB2_BREAST_TUMORS_324_UP, LANDIS_ERBB2_BREAST_TUMORS_324_UP
  GSE9006 HEALTHY VS TYPE 1 DIABETES PBMC 4MONTH POST DX DN, GSE9006 HEALTHY VS TYPE 1 DIABETES PBMC 4MONTH POST DX DN
  MIR3191 5P, MIR3191 5P
  GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_DN, GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_DN
  MIR9851 5P, MIR9851 5P
  REACTOME_GLYCOSAMINOGLYCAN_METABOLISM, REACTOME_GLYCOSAMINOGLYCAN_METABOLISM
  ASTON_MAJOR_DEPRESSIVE_DISORDER_DN, ASTON_MAJOR_DEPRESSIVE_DISORDER_DN
  GSE17721_CTRL_VS_LPS_1H_BMDC_DN, GSE17721_CTRL_VS_LPS_1H_BMDC_DN
  GSE27670_CTRL_VS_BLIMP1_TRANSDUCED_GC_BCELL_DN, GSE27670_CTRL_VS_BLIMP1_TRANSDUCED_GC_BCELL_DN
  GOBP_AMINOGLYCAN_METABOLIC_PROCESS, GOBP_AMINOGLYCAN_METABOLIC_PROCESS
  HALLMARK CHOLESTEROL HOMEOSTASIS, HALLMARK CHOLESTEROL HOMEOSTASIS
  GOBP_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS, GOBP_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS
  GOBP AMINOGLYCAN BIOSYNTHETIC PROCESS, GOBP AMINOGLYCAN BIOSYNTHETIC PROCESS
  BMI1_DN.V1_UP, BMI1_DN.V1_UP
  CYTAGCAAY UNKNOWN, CYTAGCAAY UNKNOWN
  BMI1_DN_MEL18_DN.V1_UP, BMI1_DN_MEL18_DN.V1_UP
  BRUECKNER_TARGETS_OF_MIRLET7A3_DN, BRUECKNER_TARGETS_OF_MIRLET7A3_DN
  LIM MAMMARY LUMINAL MATURE UP, LIM MAMMARY LUMINAL MATURE UP
  BLANCO_MELO_COVID19_SARS_COV_2_INFECTION_A594_CELLS_UP, BLANCO_MELO_COVID19_SARS_COV_2_INFECTION_A594_CELLS_UP
  GSE9006_1MONTH_VS_4MONTH_POST_TYPE_1_DIABETES_DX_PBMC_DN, GSE9006_1MONTH_VS_4MONTH_POST_TYPE_1_DIABETES_DX_PBMC_DN
 GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNG_KO_DN, GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNG_KO_DN
  LIM MAMMARY LUMINAL MATURE DN, LIM MAMMARY LUMINAL MATURE DN
  GOMF_EXOGENOUS_PROTEIN_BINDING, GOMF_EXOGENOUS_PROTEIN_BINDING
 GOBP_POSITIVE_REGULATION_OF_ENDOCYTOSIS, GOBP_POSITIVE_REGULATION_OF_ENDOCYTOSIS
 REACTOME CHONDROITIN_SULFATE_DERMATAN_SULFATE_METABOLISM, REACTOME_CHONDROITIN_SULFATE_DERMATAN_SULFATE_METABOLISM
MIR573, MIR573
MIR3120_5P, MIR3120_5P
 MIR3616_5P, MIR3616_5P
 PTEN_DN.V1_DN, PTEN_DN.V1_DN
 GOBP_PHOSPHATIDYLCHOLINE_METABOLIC_PROCESS, GOBP_PHOSPHATIDYLCHOLINE_METABOLIC_PROCESS
 GOBP_LIPID_STORAGE, GOBP_LIPID_STORAGE
 GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BLACK_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BLACK_UP
 YORDY_RECIPROCAL_REGULATION_BY_ETS1_AND_SP100_DN, YORDY_RECIPROCAL_REGULATION_BY_ETS1_AND_SP100_DN
 AMIT_EGF_RESPONSE_120_MCF10A, AMIT_EGF_RESPONSE_120_MCF10A
GOBP_POSITIVE_REGULATION_OF_TOR_SIGNALING, GOBP_POSITIVE_REGULATION_OF_TOR_SIGNALING
GOBP_PROTEIN_QUALITY_CONTROL_FOR_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEINS, GOBP_PROTEIN_QUALITY_CONTROL_FOR_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEINS
 GOBP_DICARBOXYLIC_ACID_TRANSPORT, GOBP_DICARBOXYLIC_ACID_TRANSPORT
MIR6879_3P, MIR6879_3P
MIR3675_5P, MIR3675_5P
GOBP_FATTY_ACID_TRANSPORT, GOBP_FATTY_ACID_TRANSPORT
||||_{I}| GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OT_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OT_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONORS_WITH_INCORPORATION_OT_MOLECULAR_OXYGEN_DONORS_WITH_INCORPORATION_OT_MOLECULAR_OXYGEN_DONORS_WITH_INCORPORATION_OT_MOLECULAR_OXYGEN_DONORS_WITH_INCORPORATION_OT_MOLECULAR_OXYGEN_DONORS_WITH_INCORPORATION_OT_MOLECULAR_OXYGEN_DONORS_WITH_INCORPORATION_OT_MOLECULAR_OXYGEN_DONORS_WITH_INCORPORATION_OX_MOLECULAR_OXYGEN_DONORS_WITH_INCORPORATION_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLECULAR_OX_MOLE
// HP_ABNORMAL_BLOOD_MONOVALENT_INORGANIC_CATION_CONCENTRATION, HP_ABNORMAL_BLOOD_MONOVALENT_INORGANIC_CATION_CONCENTRATION
 // MODULE_92, MODULE_92
/// NIKOLSKY_BREAST_CANCER_7Q21_Q22_AMPLICON, NIKOLSKY_BREAST_CANCER_7Q21_Q22_AMPLICON
/// SMID_BREAST_CANCER_ERBB2_UP, SMID_BREAST_CANCER_ERBB2_UP
// GOBP_NEGATIVE_REGULATION_OF_LIPID_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_LIPID_CATABOLIC_PROCESS
 / LANDIS_ERBB2_BREAST_TUMORS_65_UP, LANDIS_ERBB2_BREAST_TUMORS_65_UP
/ HANN_RESISTANCE_TO_BCL2_INHIBITOR_DN, HANN_RESISTANCE_TO_BCL2_INHIBITOR_DN
 ─ GOBP_CARBOHYDRATE_TRANSMEMBRANE_TRANSPORT, GOBP_CARBOHYDRATE_TRANSMEMBRANE_TRANSPORT
  GOMF_HSP70_PROTEIN_BINDING, GOMF_HSP70_PROTEIN_BINDING
  GOBP_POSITIVE_REGULATION_OF_LIPID_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_LIPID_TRANSPORT
  ZHONG_PFC_MAJOR_TYPES_OPC, ZHONG_PFC_MAJOR_TYPES_OPC
 `GAUSSMANN_MLL_AF4_FUSION_TARGETS_D_UP, GAUSSMANN_MLL_AF4_FUSION_TARGETS_D_UP
 OBP_REGULATION_OF_ERAD_PATHWAY, GOBP_REGULATION_OF_ERAD_PATHWAY
\\\\ GOBP_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GOBP_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS
MIR3085_5P, MIR3085_5P
MN MIR509 3P, MIR509 3P
 \\ KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE, KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE
HP_DEGENERATION_OF_THE_LATERAL_CORTICOSPINAL_TRACTS, HP_DEGENERATION_OF_THE_LATERAL_CORTICOSPINAL_TRACTS
 SHIPP_DLBCL_CURED_VS_FATAL_UP, SHIPP_DLBCL_CURED_VS_FATAL_UP
HP_ABNORMAL_BLOOD_SODIUM_CONCENTRATION, HP_ABNORMAL_BLOOD_SODIUM_CONCENTRATION
GOBP_HEAT_GENERATION, GOBP_HEAT_GENERATION
  GOBP_MONOCARBOXYLIC_ACID_TRANSPORT, GOBP_MONOCARBOXYLIC_ACID_TRANSPORT
 HP_FRONTOTEMPORAL_CEREBRAL_ATROPHY, HP_FRONTOTEMPORAL_CEREBRAL_ATROPHY
 GOBP_REGULATION_OF_PROTEIN_DEUBIQUITINATION, GOBP_REGULATION_OF_PROTEIN_DEUBIQUITINATION
 GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC, GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC
 GOBP_NEGATIVE_REGULATION_OF_SMOOTHENED_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_SMOOTHENED_SIGNALING_PATHWAY
 GOBP_EXOCRINE_SYSTEM_DEVELOPMENT, GOBP_EXOCRINE_SYSTEM_DEVELOPMENT
 GOBP_POSITIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY
 GOBP_REGULATION_OF_LIPID_TRANSPORT, GOBP_REGULATION_OF_LIPID_TRANSPORT
 HP_SENILE_PLAQUES, HP_SENILE_PLAQUES
HP_LACK_OF_INSIGHT, HP_LACK_OF_INSIGHT
GSE9946_MATURE_STIMULATORY_VS_LISTERIA_INF_MATURE_DC_DN, GSE9946_MATURE_STIMULATORY_VS_LISTERIA_INF_MATURE_DC_DN
 GOBP_REGULATION_OF_EXOSOMAL_SECRETION, GOBP_REGULATION_OF_EXOSOMAL_SECRETION
 GOBP_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC, GOBP_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC
 HP_LOW_LEVELS_OF_VITAMIN_D, HP_LOW_LEVELS_OF_VITAMIN_D
 HP_EEG_WITH_GENERALIZED_SLOW_ACTIVITY, HP_EEG_WITH_GENERALIZED_SLOW_ACTIVITY
  GOBP_LIPID_TRANSLOCATION, GOBP_LIPID_TRANSLOCATION
 OHGUCHI_LIVER_HNF4A_TARGETS_DN, OHGUCHI_LIVER_HNF4A_TARGETS_DN
  GOBP_LENS_DEVELOPMENT_IN_CAMERA_TYPE_EYE, GOBP_LENS_DEVELOPMENT_IN_CAMERA_TYPE_EYE
  GOCC_GOLGI_TRANS_CISTERNA, GOCC_GOLGI_TRANS_CISTERNA
  HP_ABNORMALITY_OF_SMALL_INTESTINAL_VILLUS_MORPHOLOGY, HP_ABNORMALITY_OF_SMALL_INTESTINAL_VILLUS_MORPHOLOGY
  GOBP SULFATION, GOBP SULFATION
```

4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP

GOBP_CELL_ADHESION_MOLECULE_PRODUCTION, GOBP_CELL_ADHESION_MOLECULE_PRODUCTION GOMF_LYSOPHOSPHOLIPASE_ACTIVITY, GOMF_LYSOPHOSPHOLIPASE_ACTIVITY GOMF_FATTY_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_FATTY_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY HP_INTRINSIC_HAND_MUSCLE_ATROPHY, HP_INTRINSIC_HAND_MUSCLE_ATROPHY GOCC_CUTICULAR_PLATE, GOCC_CUTICULAR_PLATE GOBP_NEGATIVE_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCESS GOBP_POSITIVE_REGULATION_OF_MONOCYTE_CHEMOTAXIS, GOBP_POSITIVE_REGULATION_OF_MONOCYTE_CHEMOTAXIS GOBP_FEVER_GENERATION, GOBP_FEVER_GENERATION HP_ACANTHOCYTOSIS, HP_ACANTHOCYTOSIS GOBP_POSITIVE_REGULATION_OF_ERAD_PATHWAY, GOBP_POSITIVE_REGULATION_OF_ERAD_PATHWAY GOBP_NEGATIVE_REGULATION_OF_PLATELET_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_PLATELET_ACTIVATION HP_COLLECTIONISM, HP_COLLECTIONISM VERRECCHIA_RESPONSE_TO_TGFB1_C3, VERRECCHIA_RESPONSE_TO_TGFB1_C3 WP_CYTOKINES_AND_INFLAMMATORY_RESPONSE, WP_CYTOKINES_AND_INFLAMMATORY_RESPONSE HP_WEAKNESS_OF_THE_INTRINSIC_HAND_MUSCLES, HP_WEAKNESS_OF_THE_INTRINSIC_HAND_MUSCLES GOMF_MANNOSE_BINDING, GOMF_MANNOSE_BINDING GOBP_POSITIVE_REGULATION_OF_HEAT_GENERATION, GOBP_POSITIVE_REGULATION_OF_HEAT_GENERATION HP_CUTANEOUS_ABSCESS, HP_CUTANEOUS_ABSCESS REACTOME_PREGNENOLONE_BIOSYNTHESIS, REACTOME_PREGNENOLONE_BIOSYNTHESIS HP_ELEVATED_URINARY_CARBOXYLIC_ACID, HP_ELEVATED_URINARY_CARBOXYLIC_ACID DESCARTES_FETAL_EYE_LENS_FIBRE_CELLS, DESCARTES_FETAL_EYE_LENS_FIBRE_CELLS RAY_ALZHEIMERS_DISEASE, RAY_ALZHEIMERS_DISEASE GOBP_C21_STEROID_HORMONE_BIOSYNTHETIC_PROCESS, GOBP_C21_STEROID_HORMONE_BIOSYNTHETIC_PROCESS GOMF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_BINDING, GOMF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_BINDING GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_SURFACE, GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_SURFACE HAY_BONE_MARROW_CD34_POS_ERP, HAY_BONE_MARROW_CD34_POS_ERP HP_CORONAL_CLEFT_VERTEBRAE, HP_CORONAL_CLEFT_VERTEBRAE GOBP_REGULATION_OF_MONOCYTE_CHEMOTAXIS, GOBP_REGULATION_OF_MONOCYTE_CHEMOTAXIS GOBP_REGULATION_OF_AMINO_ACID_TRANSPORT, GOBP_REGULATION_OF_AMINO_ACID_TRANSPORT MEBARKI_HCC_PROGENITOR_WNT_UP_CTNNB1_DEPENDENT_BLOCKED_BY_FZD8CRD, MEBARKI_HCC_PROGENITOR_WNT_UP_CTNNB1_DEPENDENT_BLOCKED_BY_FZD8CRD GOBP_EPITHELIAL_MESENCHYMAL_CELL_SIGNALING, GOBP_EPITHELIAL_MESENCHYMAL_CELL_SIGNALING LI_CISPLATIN_RESISTANCE_UP, LI_CISPLATIN_RESISTANCE_UP GOMF_CCR_CHEMOKINE_RECEPTOR_BINDING, GOMF_CCR_CHEMOKINE_RECEPTOR_BINDING GNF2_MAPT, GNF2_MAPT BIOCARTA_INFLAM_PATHWAY, BIOCARTA_INFLAM_PATHWAY GOBP_CONVERGENT_EXTENSION, GOBP_CONVERGENT_EXTENSION GOBP_NEGATIVE_REGULATION_OF_OXIDOREDUCTASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_OXIDOREDUCTASE_ACTIVITY GOBP_POSITIVE_REGULATION_OF_CELL_ADHESION_MOLECULE_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_CELL_ADHESION_MOLECULE_PRODUCTION GOBP_POSITIVE_REGULATION_OF_HUMORAL_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_HUMORAL_IMMUNE_RESPONSE GOBP_OVULATION, GOBP_OVULATION