| | GSE37532_WT_VS_PPARG_KO_VISCERAL GO_SISTER_CHROMATID_SEGREGATION | I, GSE7460_CD8_TCELL_VS_TREG_ACT_DN L_ADIPOSE_TISSUE_TREG_UP, GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TREG_UP N, GO_SISTER_CHROMATID_SEGREGATION |
|--|--|---|
| | GSE19825_CD24LOW_VS_IL2RA_HIGH_DA GSE1460_DP_THYMOCYTE_VS_NAIVE_CI GSE29164_UNTREATED_VS_CD8_TCELL_A | GION, GO_CHROMOSOME_CENTROMERIC_REGION AY3_EFF_CD8_TCELL_DN, GSE19825_CD24LOW_VS_IL2RA_HIGH_DAY3_EFF_CD8_TCELL_DN D4_TCELL_CORD_BLOOD_UP, GSE1460_DP_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_UP AND_IL12_TREATED_MELANOMA_DAY3_DN, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_DN CD4_TCELL_EROM_LN_UP_CSE37522_TREC_VS_TCONV_RPARC_VO_CD4_TCELL_EROM_LN_UP |
| | GO_CONDENSED_CHROMOSOME, GO_C | ATED_CD4_TCELL_PPARG1_AND_FOXP3_TRASDUCED_DN, GSE37534_UNTREATED_VS_GW1929_TREATED_CD4_TCELL_PPARG1_AND_FOXP3_TRASDUCED_DN |
| | GSE25085_FETAL_LIVER_VS_ADULT_BM_ GSE9878_CTRL_VS_EBF_TRANSDUCED_P | TED_STAT3_KO_CD4_TCELL_UP, GSE21670_IL6_VS_TGFB_AND_IL6_TREATED_STAT3_KO_CD4_TCELL_UP _SP4_THYMIC_IMPLANT_DN, GSE25085_FETAL_LIVER_VS_ADULT_BM_SP4_THYMIC_IMPLANT_DN PAX5_KO_PRO_BCELL_DN, GSE9878_CTRL_VS_EBF_TRANSDUCED_PAX5_KO_PRO_BCELL_DN A_CD4_TCELL_UP, GSE39820_CTRL_VS_TGFBETA3_IL6_IL23A_CD4_TCELL_UP |
| | GSE3920_UNTREATED_VS_IFNB_TREATE GSE37532_TREG_VS_TCONV_CD4_TCELL | ELL_24H_UP, GSE7764_IL15_TREATED_VS_CTRL_NK_CELL_24H_UP ED_ENDOTHELIAL_CELL_DN, GSE3920_UNTREATED_VS_IFNB_TREATED_ENDOTHELIAL_CELL_DN L_FROM_VISCERAL_ADIPOSE_TISSUE_DN, GSE37532_TREG_VS_TCONV_CD4_TCELL_FROM_VISCERAL_ADIPOSE_TISSUE_DN MACROPHAGE_DN, GSE33162_HDAC3_KO_VS_HDAC3_KO_MACROPHAGE_DN |
| | GSE7852_LN_VS_THYMUS_TCONV_DN, G GSE40274_CTRL_VS_FOXP3_AND_HELIOS | _ACT_CD4_TCELL_48H_DN, GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_48H_DN GSE7852_LN_VS_THYMUS_TCONV_DN S_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN TCELL_UP, GSE9650_EFFECTOR_VS_MEMORY_CD8_TCELL_UP |
| | GSE16450_IMMATURE_VS_MATURE_NEU GSE22229_RENAL_TRANSPLANT_VS_HEA | JRON_CELL_LINE_DN, GSE16450_IMMATURE_VS_MATURE_NEURON_CELL_LINE_DN ALTHY_PBMC_UP, GSE22229_RENAL_TRANSPLANT_VS_HEALTHY_PBMC_UP INE_PBMC_DN, GSE13485_DAY1_VS_DAY7_YF17D_VACCINE_PBMC_DN |
| | GO_CELL_CYCLE_CHECKPOINT, GO_CEL GSE27786_NEUTROPHIL_VS_MONO_MAG | LL_CYCLE_CHECKPOINT C_DN, GSE27786_NEUTROPHIL_VS_MONO_MAC_DN D8_SINGLE_POSITIVE_THYMOCYTE_UP, GSE22601_CD4_SINGLE_POSITIVE_VS_CD8_SINGLE_POSITIVE_THYMOCYTE_UP |
| | GSE11386_NAIVE_VS_MEMORY_BCELL_U GSE19941_LPS_VS_LPS_AND_IL10_STIM_I | A_CD4_TCELL_UP, GSE39820_CTRL_VS_TGFBETA1_IL6_IL23A_CD4_TCELL_UP UP, GSE11386_NAIVE_VS_MEMORY_BCELL_UP IL10_KO_MACROPHAGE_UP, GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_UP M_CD8_TCELL_DN, GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_CD8_TCELL_DN |
| | PYEON_CANCER_HEAD_AND_NECK_VS GSE25146_UNSTIM_VS_HELIOBACTER_PY | _UP, GSE2128_C57BL6_VS_NOD_THYMOCYTE_UP S_CERVICAL_UP, PYEON_CANCER_HEAD_AND_NECK_VS_CERVICAL_UP YLORI_LPS_STIM_AGS_CELL_DN, GSE25146_UNSTIM_VS_HELIOBACTER_PYLORI_LPS_STIM_AGS_CELL_DN REATED_STAT3_KO_CD4_TCELL_UP, GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_STAT3_KO_CD4_TCELL_UP |
| | GSE6269_FLU_VS_STREP_PNEUMO_INF_E GSE25146_UNSTIM_VS_HELIOBACTER_PY GSE16451_CTRL_VS_WEST_EQUINE_ENC | PBMC_UP, GSE6269_FLU_VS_STREP_PNEUMO_INF_PBMC_UP YLORI_LPS_STIM_AGS_CELL_UP, GSE25146_UNSTIM_VS_HELIOBACTER_PYLORI_LPS_STIM_AGS_CELL_UP C_VIRUS_IMMATURE_NEURON_CELL_LINE_DN, GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_IMMATURE_NEURON_CELL_LINE_DN REATED_CD4_TCELL_UP, GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_UP |
| | GSE6269_HEALTHY_VS_STAPH_AUREUS_ GSE33425_CD161_HIGH_VS_INT_CD8_TCI GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_N | S_INF_PBMC_UP, GSE6269_HEALTHY_VS_STAPH_AUREUS_INF_PBMC_UP ELL_DN, GSE33425_CD161_HIGH_VS_INT_CD8_TCELL_DN MONOCYTE_12H_UP, GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_12H_UP IL6_STIM_MACROPHAGE_45MIN_UP, GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_UP |
| | GO_SPINDLE_POLE, GO_SPINDLE_POLE WHITFIELD_CELL_CYCLE_G2, WHITFIEL GSE18893_TCONV_VS_TREG_24H_CULTU | |
| | GO_NEGATIVE_REGULATION_OF_MITO GSE39110_UNTREATED_VS_IL2_TREATED GSE6259_33D1_POS_DC_VS_CD4_TCELL_U | OTIC_CELL_CYCLE, GO_NEGATIVE_REGULATION_OF_MITOTIC_CELL_CYCLE D_CD8_TCELL_DAY3_POST_IMMUNIZATION_DN, GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY3_POST_IMMUNIZATION_DN UP, GSE6259_33D1_POS_DC_VS_CD4_TCELL_UP _UP, GSE17721_CTRL_VS_POLYIC_24H_BMDC_UP |
| | GSE24634_TEFF_VS_TCONV_DAY10_IN_C GSE7596_AKT_TRANSD_VS_CTRL_CD4_TG GSE3720_UNSTIM_VS_PMA_STIM_VD2_G | CULTURE_UP, GSE24634_TEFF_VS_TCONV_DAY10_IN_CULTURE_UP CONV_WITH_TGFB_UP, GSE7596_AKT_TRANSD_VS_CTRL_CD4_TCONV_WITH_TGFB_UP GAMMADELTA_TCELL_DN, GSE3720_UNSTIM_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_DN VDOSE_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_UP, GSE32986_GMCSF_AND_CURDLAN_LOWDOSE_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_UP |
| | GO_SISTER_CHROMATID_COHESION, GO REACTOME_MITOTIC_PROMETAPHASE, GO_CONDENSED_CHROMOSOME_CENT | O_SISTER_CHROMATID_COHESION |
| | EPPERT_PROGENITOR, EPPERT_PROGEN GSE17186_CD21LOW_VS_CD21HIGH_TRA GO_MITOTIC_SISTER_CHROMATID_SEGI | |
| | GSE8621_LPS_PRIMED_UNSTIM_VS_LPS_I GO_DNA_INTEGRITY_CHECKPOINT, GO_ HOXA9_DN.V1_DN, HOXA9_DN.V1_DN | PRIMED_AND_LPS_STIM_MACROPHAGE_DN, GSE8621_LPS_PRIMED_UNSTIM_VS_LPS_PRIMED_AND_LPS_STIM_MACROPHAGE_DN |
| | GSE37532_WT_VS_PPARG_KO_LN_TCONG GSE6259_FLT3L_INDUCED_DEC205_POS_1 | CROPHAGE_45MIN_UP, GSE5589_WT_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_UP V_DN, GSE37532_WT_VS_PPARG_KO_LN_TCONV_DN DC_VS_BCELL_DN, GSE6259_FLT3L_INDUCED_DEC205_POS_DC_VS_BCELL_DN ANCE, REACTOME_CHROMOSOME_MAINTENANCE |
| | GSE14415_ACT_TCONV_VS_ACT_NATUR GSE13485_CTRL_VS_DAY7_YF17D_VACCI | RAL_TREG_DN, GSE14415_ACT_TCONV_VS_ACT_NATURAL_TREG_DN INE_PBMC_DN, GSE13485_CTRL_VS_DAY7_YF17D_VACCINE_PBMC_DN 1_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_UP, GSE10240_CTRL_VS_IL17_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_UP |
| | GSE9239_CTRL_VS_TNF_INHIBITOR_TREA GSE14415_INDUCED_TREG_VS_FAILED_II ERB2_UP.V1_DN, ERB2_UP.V1_DN | ATED_DC_DN, GSE9239_CTRL_VS_TNF_INHIBITOR_TREATED_DC_DN INDUCED_TREG_DN, GSE14415_INDUCED_TREG_VS_FAILED_INDUCED_TREG_DN _CD8_TCELL_DN, GSE17812_WT_VS_THPOK_KO_MEMORY_CD8_TCELL_DN |
| | GO_NEGATIVE_REGULATION_OF_PROTE LEE_EARLY_T_LYMPHOCYTE_UP, LEE_E GSE20727_DNFB_ALLERGEN_VS_ROS_INIT | EIN_COMPLEX_DISASSEMBLY, GO_NEGATIVE_REGULATION_OF_PROTEIN_COMPLEX_DISASSEMBLY |
| | VEGF_A_UP.V1_DN, VEGF_A_UP.V1_DN GSE19941_UNSTIM_VS_LPS_STIM_IL10_KG GSE37301_HEMATOPOIETIC_STEM_CELL | |
| | GO_CHROMATIN_ASSEMBLY_OR_DISAS GSE3982_DC_VS_TH1_DN, GSE3982_DC_V GSE32901_NAIVE_VS_TH17_NEG_CD4_TC | SSEMBLY, GO_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY |
| | GSE15930_STIM_VS_STIM_AND_TRICHOS GO_DNA_SYNTHESIS_INVOLVED_IN_DN | STATINA_24H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_24H_CD8_T_CELL_DN NA_REPAIR, GO_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR _BCELL_UP, GSE13547_2H_VS_12_H_ANTI_IGM_STIM_BCELL_UP |
| | GSE29614_DAY3_VS_DAY7_TIV_FLU_VAC GSE17301_ACD3_ACD28_VS_ACD3_ACD2 GSE12003_MIR223_KO_VS_WT_BM_PROG | CCINE_PBMC_DN, GSE29614_DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMC_DN 28_AND_IFNA5_STIM_CD8_TCELL_UP, GSE17301_ACD3_ACD28_VS_ACD3_ACD28_AND_IFNA5_STIM_CD8_TCELL_UP 3ENITOR_4D_CULTURE_UP, GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_4D_CULTURE_UP D_ACT_CD4_TCELL_48H_DN, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_DN |
| | GO_NUCLEASE_ACTIVITY, GO_NUCLEASE GSE3982_MAST_CELL_VS_TH1_DN, GSE39 GSE20366_EX_VIVO_VS_HOMEOSTATIC_0 | SE_ACTIVITY |
| | GSE15330_LYMPHOID_MULTIPOTENT_VS | S_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_UP, GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_UP L_UP, GSE39820_CTRL_VS_IL1B_IL6_CD4_TCELL_UP |
| | GSE27786_LIN_NEG_VS_BCELL_UP, GSE27 GSE10240_CTRL_VS_IL17_AND_IL22_STIM GSE15735_2H_VS_12H_HDAC_INHIBITOR | |
| | GSE7509_UNSTIM_VS_IFNA_STIM_IMMA' GSE40274_CTRL_VS_EOS_TRANSDUCED_ GSE25088_WT_VS_STAT6_KO_MACROPH. | TURE_DC_UP, GSE7509_UNSTIM_VS_IFNA_STIM_IMMATURE_DC_UP _ACTIVATED_CD4_TCELL_UP, GSE40274_CTRL_VS_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP IAGE_ROSIGLITAZONE_AND_IL4_STIM_UP, GSE25088_WT_VS_STAT6_KO_MACROPHAGE_ROSIGLITAZONE_AND_IL4_STIM_UP TIONAL_BCELL_CORD_BLOOD_DN, GSE17186_NAIVE_VS_CD21LOW_TRANSITIONAL_BCELL_CORD_BLOOD_DN |
| | GO_RESPONSE_TO_IONIZING_RADIATIO GSE33292_DN3_THYMOCYTE_VS_TCELL_ REACTOME_MEIOSIS, REACTOME_MEIO | ON, GO_RESPONSE_TO_IONIZING_RADIATION _LYMPHOMA_FROM_TCF1_KO_DN, GSE33292_DN3_THYMOCYTE_VS_TCELL_LYMPHOMA_FROM_TCF1_KO_DN OSIS |
| | GO_REGULATION_OF_MICROTUBULE_P PIGF_UP.V1_UP, PIGF_UP.V1_UP GO_RECOMBINATIONAL_REPAIR, GO_R | S, REACTOME_MITOTIC_G2_G2_M_PHASES POLYMERIZATION_OR_DEPOLYMERIZATION, GO_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION RECOMBINATIONAL_REPAIR CMV_INFECTION_DN, GSE45365_WT_VS_IFNAR_KO_BCELL_MCMV_INFECTION_DN |
| | GSE17721_POLYIC_VS_CPG_0.5H_BMDC_I GSE13485_DAY3_VS_DAY7_YF17D_VACCI | DN, GSE17721_POLYIC_VS_CPG_0.5H_BMDC_DN INE_PBMC_DN, GSE13485_DAY3_VS_DAY7_YF17D_VACCINE_PBMC_DN FF_CD8_TCELL_DN, GSE10239_KLRG1INT_VS_KLRG1HIGH_EFF_CD8_TCELL_DN |
| | GSE23502_BM_VS_COLON_TUMOR_MYEIGSE17186_CD21LOW_VS_CD21HIGH_TRA | TION_FORK LOID_DERIVED_SUPPRESSOR_CELL_DN, GSE23502_BM_VS_COLON_TUMOR_MYELOID_DERIVED_SUPPRESSOR_CELL_DN ANSITIONAL_BCELL_DN, GSE17186_CD21LOW_VS_CD21HIGH_TRANSITIONAL_BCELL_DN ATID_SEGREGATION, GO_REGULATION_OF_SISTER_CHROMATID_SEGREGATION |
| | GO_EXONUCLEASE_ACTIVITY, GO_EXON GSE23502_WT_VS_HDC_KO_MYELOID_DI GSE13411_PLASMA_CELL_VS_MEMORY_I | NUCLEASE_ACTIVITY DERIVED_SUPPRESSOR_CELL_COLON_TUMOR_UP, GSE23502_WT_VS_HDC_KO_MYELOID_DERIVED_SUPPRESSOR_CELL_COLON_TUMOR_UP BCELL_UP, GSE13411_PLASMA_CELL_VS_MEMORY_BCELL_UP , GSE7831_1H_VS_4H_CPG_STIM_PDC_DN |
| | AMUNDSON_GAMMA_RADIATION_REST GSE39556_UNTREATED_VS_3H_POLYIC_I REACTOME_DEPOSITION_OF_NEW_CEN | , GSE7831_1H_VS_4H_CPG_STIM_PDC_DN GPONSE, AMUNDSON_GAMMA_RADIATION_RESPONSE INJ_MOUSE_NK_CELL_UP, GSE39556_UNTREATED_VS_3H_POLYIC_INJ_MOUSE_NK_CELL_UP IPA_CONTAINING_NUCLEOSOMES_AT_THE_CENTROMERE, REACTOME_DEPOSITION_OF_NEW_CENPA_CONTAINING_NUCLEOSOMES_AT_THE_CENTROMERE _METABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_DNA_METABOLIC_PROCESS |
| | GSE27786_CD8_TCELL_VS_NEUTROPHIL_ PYEON_HPV_POSITIVE_TUMORS_UP, PY WILCOX_RESPONSE_TO_PROGESTERON | _DN, GSE27786_CD8_TCELL_VS_NEUTROPHIL_DN TEON_HPV_POSITIVE_TUMORS_UP IE_UP, WILCOX_RESPONSE_TO_PROGESTERONE_UP |
| | GO_DNA_PACKAGING, GO_DNA_PACKA | ULAR_CATABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_CELLULAR_CATABOLIC_PROCESS AGING ION_CD11B_DC_IFNAR_KO_DN, GSE45365_HEALTHY_VS_MCMV_INFECTION_CD11B_DC_IFNAR_KO_DN |
| | GO_SPINDLE_MICROTUBULE, GO_SPIND GSE17721_POLYIC_VS_GARDIQUIMOD_21 | H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_2H_BMDC_UP |
| | GO_NON_RECOMBINATIONAL_REPAIR, NAKAYAMA_SOFT_TISSUE_TUMORS_PC GSE15330_LYMPHOID_MULTIPOTENT_VS | NT_PBMC_DN, GSE34205_HEALTHY_VS_RSV_INF_INFANT_PBMC_DN , GO_NON_RECOMBINATIONAL_REPAIR CA2_UP, NAKAYAMA_SOFT_TISSUE_TUMORS_PCA2_UP S_GRANULOCYTE_MONOCYTE_PROGENITOR_UP, GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_UP |
| | AAGWWRNYGGC_UNKNOWN, AAGWW MOLENAAR_TARGETS_OF_CCND1_AND GO_MICROTUBULE_ORGANIZING_CENT | D_CDK4_DN, MOLENAAR_TARGETS_OF_CCND1_AND_CDK4_DN TER_ORGANIZATION, GO_MICROTUBULE_ORGANIZING_CENTER_ORGANIZATION |
| | REACTOME_LOSS_OF_NLP_FROM_MITO GEORGES_CELL_CYCLE_MIR192_TARGET | OOD_VS_THYMIC_STROMAL_CELL_DN, GSE1460_NAIVE_CD4_TCELL_ADULT_BLOOD_VS_THYMIC_STROMAL_CELL_DN OTIC_CENTROSOMES, REACTOME_LOSS_OF_NLP_FROM_MITOTIC_CENTROSOMES TS, GEORGES_CELL_CYCLE_MIR192_TARGETS |
| | GO_POSTREPLICATION_REPAIR, GO_POSTREPLICATION_RE | CHERT_MITOSIS_LIN9_TARGETS |
| | GO_MICROTUBULE_ORGANIZING_CENT | EGION_VS_WHOLE_MEDULLA_THYMUS_UP, GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_UP TER_PART, GO_MICROTUBULE_ORGANIZING_CENTER_PART OS_CD4_EFF_TCELL_D6_LCMV_UP, GSE43863_NAIVE_VS_LY6C_INT_CXCR5POS_CD4_EFF_TCELL_D6_LCMV_UP ROTUBULE_BINDING |
| | GSE23505_IL6_IL1_VS_IL6_IL1_IL23_TREAT | Έ_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_9H_UP, GSE2405_HEAT_KILLED_LYSATE_VS_LIVE_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_9H_UP TED_CD4_TCELL_UP, GSE23505_IL6_IL1_VS_IL6_IL1_IL23_TREATED_CD4_TCELL_UP ΓURE_2_UP, MULLIGHAN_NPM1_MUTATED_SIGNATURE_2_UP 3_5_EXONUCLEASE_ACTIVITY |
| | SMIRNOV_RESPONSE_TO_IR_6HR_DN, SI GSE36078_UNTREATED_VS_AD5_T425A_H | MODELING, GO_ATP_DEPENDENT_CHROMATIN_REMODELING MIRNOV_RESPONSE_TO_IR_6HR_DN HEXON_INF_MOUSE_LUNG_DC_DN, GSE36078_UNTREATED_VS_AD5_T425A_HEXON_INF_MOUSE_LUNG_DC_DN SOME, GO_CONDENSED_NUCLEAR_CHROMOSOME |
| | GO_SPINDLE_ASSEMBLY, GO_SPINDLE_A GSE12839_CTRL_VS_IL12_TREATED_PBMG GSE8921_UNSTIM_VS_TLR1_2_STIM_MON | C_UP, GSE12839_CTRL_VS_IL12_TREATED_PBMC_UP NOCYTE_6H_DN, GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_6H_DN |
| | GSE3982_DC_VS_MAC_LPS_STIM_DN, GSEGSE41978_WT_VS_ID2_KO_AND_BIM_KO_GSE45365_HEALTHY_VS_MCMV_INFECTE | CKLRG1_LOW_EFFECTOR_CD8_TCELL_UP, GSE41978_WT_VS_ID2_KO_AND_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_UP CION_CD11B_DC_DN, GSE45365_HEALTHY_VS_MCMV_INFECTION_CD11B_DC_DN |
| | GO_DNA_REPLICATION_INDEPENDENT GSE1925_CTRL_VS_IFNG_PRIMED_MACR GO_MITOTIC_SPINDLE, GO_MITOTIC_SP | C_VS_BLOOD_UP, PETROVA_ENDOTHELIUM_LYMPHATIC_VS_BLOOD_UP I_NUCLEOSOME_ORGANIZATION, GO_DNA_REPLICATION_INDEPENDENT_NUCLEOSOME_ORGANIZATION ROPHAGE_UP, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_UP PINDLE 10_STIM_IL10_KO_MACROPHAGE_DN, GSE9509_10MIN_VS_30MIN_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_DN |
| | RIGGI_EWING_SARCOMA_PROGENITOR ZHANG_RESPONSE_TO_IKK_INHIBITOR_ GO_MICROTUBULE_ASSOCIATED_COME | IO_STING_IETO_RO_MACROTTIAGE_DIV, GSE9309_TOMIN_VS_30MIN_EFS_AND_IETO_STING_IETO_RO_MACROTTIAGE_DIV R_DN, RIGGI_EWING_SARCOMA_PROGENITOR_DN R_AND_TNF_DN, ZHANG_RESPONSE_TO_IKK_INHIBITOR_AND_TNF_DN PLEX, GO_MICROTUBULE_ASSOCIATED_COMPLEX IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_UP |
| | GSE2405_0H_VS_24H_A_PHAGOCYTOPHI GSE369_SOCS3_KO_VS_IFNG_KO_LIVER_ GOTZMANN_EPITHELIAL_TO_MESENCH | ILUM_STIM_NEUTROPHIL_DN, GSE2405_0H_VS_24H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_DN DN, GSE369_SOCS3_KO_VS_IFNG_KO_LIVER_DN HYMAL_TRANSITION_UP, GOTZMANN_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_UP REGION_VS_WHOLE_CORTEX_THYMUS_UP, GSE18281_PERIMEDULLARY_CORTICAL_REGION_VS_WHOLE_CORTEX_THYMUS_UP |
| | // MODULE_331, MODULE_331 // SHETH_LIVER_CANCER_VS_TXNIP_LOSS // IWANAGA_E2F1_TARGETS_INDUCED_BY | S_PAM2, SHETH_LIVER_CANCER_VS_TXNIP_LOSS_PAM2 Y_SERUM, IWANAGA_E2F1_TARGETS_INDUCED_BY_SERUM IT_DNA_REPLICATION, GO_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION |
| 'ARGETS_36HR_DN, ZHANG_TLX_TARGETS_36HR_DN | PID_AURORA_A_PATHWAY, PID_AUROR MODULE_397, MODULE_397 GSE40274_FOXP3_VS_FOXP3_AND_HELIC GO_MEIOTIC_CELL_CYCLE, GO_MEIOTIC | OS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN |
| | GSE41978_WT_VS_ID2_KO_AND_BIM_KO_GSE27786_ERYTHROBLAST_VS_NEUTROB | R, GO_INTERSTRAND_CROSS_LINK_REPAIR D_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN, GSE41978_WT_VS_ID2_KO_AND_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN PHIL_UP, GSE27786_ERYTHROBLAST_VS_NEUTROPHIL_UP ION, GO_REGULATION_OF_NUCLEAR_DIVISION |
| | GO_NUCLEAR_REPLICATION_FORK, GO_GSE27786_CD4_TCELL_VS_MONO_MAC_I | ED_ENDOTHELIAL_CELL_DN, GSE3920_UNTREATED_VS_IFNA_TREATED_ENDOTHELIAL_CELL_DN |
| | GSE11961_UNSTIM_VS_ANTI_IGM_AND_ODONNELL_TFRC_TARGETS_DN,ODON | |
| | GO_CELLULAR_RESPONSE_TO_RADIATI GO_RESPONSE_TO_UV, GO_RESPONSE_T DE_YY1_TARGETS_DN, DE_YY1_TARGETS | |
| | GSE9960_GRAM_POS_VS_GRAM_NEG_ANGO_DEOXYRIBONUCLEASE_ACTIVITY, C | |
| | GO_CENTROMERE_COMPLEX_ASSEMBLE GSE17721_POLYIC_VS_GARDIQUIMOD_61 PID_PLK1_PATHWAY, PID_PLK1_PATHW | |
| | GRAHAM_CML_QUIESCENT_VS_NORMA GO_CENTROSOME_CYCLE, GO_CENTRO RRCCGTTA_UNKNOWN, RRCCGTTA_UN | |
| | GTGACTT_MIR224, GTGACTT_MIR224 GSE28783_ANTI_MIR33_VS_CTRL_ATHER GSE6875_WT_VS_FOXP3_KO_TREG_DN, G | ROSCLEROSIS_MACROPHAGE_DN, GSE28783_ANTI_MIR33_VS_CTRL_ATHEROSCLEROSIS_MACROPHAGE_DN |
| | GACTGTT_MIR212_MIR132, GACTGTT_MICORDENONSI_YAP_CONSERVED_SIGNA | IR212_MIR132 ATURE, CORDENONSI_YAP_CONSERVED_SIGNATURE DN, GSE17721_CTRL_VS_POLYIC_2H_BMDC_DN |
| | GO_ENDODEOXYRIBONUCLEASE_ACTIV SCIBETTA_KDM5B_TARGETS_DN, SCIBET GO_REGULATION_OF_DNA_RECOMBINA | VITY, GO_ENDODEOXYRIBONUCLEASE_ACTIVITY |
| | ATTACAT_MIR3803P, ATTACAT_MIR3803 | BP IMUNE_RECEPTORS, GO_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS TILE_RESPONSE_CLUSTER_D3 |
| | REACTOME_HOMOLOGOUS_RECOMBIN KMCATNNWGGA_UNKNOWN, KMCATN | AST_DN, GSE27786_CD4_TCELL_VS_ERYTHTROBLAST_DN IATION_REPAIR_OF_REPLICATION_INDEPENDENT_DOUBLE_STRAND_BREAKS, REACTOME_HOMOLOGOUS_RECOMBINATION_REPAIR_OF_REPLICATION_INDEPENDEN NNWGGA_UNKNOWN ROGENITOR_UP, HADDAD_T_LYMPHOCYTE_AND_NK_PROGENITOR_UP |
| | GO_SOMATIC_CELL_DNA_RECOMBINAT SERVITJA_LIVER_HNF1A_TARGETS_UP, S chr15q21, chr15q21 | TION, GO_SOMATIC_CELL_DNA_RECOMBINATION |
| | GSE17721_LPS_VS_CPG_8H_BMDC_UP, GS GSE11961_FOLLICULAR_BCELL_VS_GERN | N, BYSTROEM_CORRELATED_WITH_IL5_DN SE17721_LPS_VS_CPG_8H_BMDC_UP MINAL_CENTER_BCELL_DAY7_DN, GSE11961_FOLLICULAR_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_DN , GO_EXON_EXON_JUNCTION_COMPLEX |
| | GSE45365_WT_VS_IFNAR_KO_CD11B_DC_PRC2_EZH2_UP.V1_UP, PRC2_EZH2_UP.V | OR_HIV_SPECIFIC_CD8_TCELL_DN, GSE24081_CONTROLLER_VS_PROGRESSOR_HIV_SPECIFIC_CD8_TCELL_DN _MCMV_INFECTION_DN, GSE45365_WT_VS_IFNAR_KO_CD11B_DC_MCMV_INFECTION_DN /1_UP _ASSEMBLY, GO_REGULATION_OF_CELL_JUNCTION_ASSEMBLY |
| | GSE3982_MAST_CELL_VS_CENT_MEMOR MODULE_303, MODULE_303 | CE_OF_CELL_POLARITY, GO_ESTABLISHMENT_OR_MAINTENANCE_OF_CELL_POLARITY RY_CD4_TCELL_DN, GSE3982_MAST_CELL_VS_CENT_MEMORY_CD4_TCELL_DN MPHOBLASTIC_LEUKEMIA, TAYLOR_METHYLATED_IN_ACUTE_LYMPHOBLASTIC_LEUKEMIA |
| | CEBP_C, CEBP_C GO_PROTEIN_LOCALIZATION_TO_CYTO PID_ATM_PATHWAY, PID_ATM_PATHW GO_HISTONE_LYSINE_N_METHYLTRAN | OSKELETON, GO_PROTEIN_LOCALIZATION_TO_CYTOSKELETON /AY ISFERASE_ACTIVITY, GO_HISTONE_LYSINE_N_METHYLTRANSFERASE_ACTIVITY |
| | GO_CHROMOSOME_LOCALIZATION, GO GO_ENDONUCLEASE_ACTIVITY, GO_EN GO_MITOTIC_SPINDLE_ASSEMBLY, GO_I GSE17721_LPS_VS_PAM3CSK4_0.5H_BMD0 | DONUCLEASE_ACTIVITY |
| | PID_PRL_SIGNALING_EVENTS_PATHWA GO_DNA_DEPENDENT_DNA_REPLICATI GO_PEPTIDYL_LYSINE_DIMETHYLATION | _DN, DORN_ADENOVIRUS_INFECTION_24HR_DN AY, PID_PRL_SIGNALING_EVENTS_PATHWAY ION_MAINTENANCE_OF_FIDELITY, GO_DNA_DEPENDENT_DNA_REPLICATION_MAINTENANCE_OF_FIDELITY N, GO_PEPTIDYL_LYSINE_DIMETHYLATION |
| | GO_MITOTIC_G2_M_TRANSITION_CHEC GO_DNA_LIGATION, GO_DNA_LIGATIO | T_DN, SHEPARD_CRUSH_AND_BURN_MUTANT_DN CKPOINT, GO_MITOTIC_G2_M_TRANSITION_CHECKPOINT |
| | KEGG_HOMOLOGOUS_RECOMBINATION GRAHAM_CML_QUIESCENT_VS_CML_D | N, KEGG_HOMOLOGOUS_RECOMBINATION DIVIDING_DN, GRAHAM_CML_QUIESCENT_VS_CML_DIVIDING_DN N, GO_METAPHASE_PLATE_CONGRESSION |
| | GO_CHROMOSOME_CONDENSATION, G KENNY_CTNNB1_TARGETS_UP, KENNY_ GO_NEGATIVE_REGULATION_OF_CHRO GO_MALE_MEIOSIS, GO_MALE_MEIOSIS | GO_CHROMOSOME_CONDENSATION _CTNNB1_TARGETS_UP DMOSOME_SEGREGATION, GO_NEGATIVE_REGULATION_OF_CHROMOSOME_SEGREGATION |
| | GO_SPINDLE_CHECKPOINT, GO_SPINDL PID_EPHB_FWD_PATHWAY, PID_EPHB_F GSE12963_UNINF_VS_ENV_AND_NEF_DE REACTOME_FACTORS_INVOLVED_IN_M | LE_CHECKPOINT FWD_PATHWAY EFICIENT_HIV1_INF_CD4_TCELL_DN, GSE12963_UNINF_VS_ENV_AND_NEF_DEFICIENT_HIV1_INF_CD4_TCELL_DN IEGAKARYOCYTE_DEVELOPMENT_AND_PLATELET_PRODUCTION, REACTOME_FACTORS_INVOLVED_IN_MEGAKARYOCYTE_DEVELOPMENT_AND_PLATELET_PRODUC |
| | CHIBA_RESPONSE_TO_TSA_DN, CHIBA_ | KPOINT, GO_MITOTIC_G2_DNA_DAMAGE_CHECKPOINT |
| | GSE29614_CTRL_VS_TIV_FLU_VACCINE_I GO_GAMMA_TUBULIN_COMPLEX, GO_C GO_MITOTIC_CYTOKINESIS, GO_MITOTI MODULE_222, MODULE_222 | PBMC_2007_DN, GSE29614_CTRL_VS_TIV_FLU_VACCINE_PBMC_2007_DN GAMMA_TUBULIN_COMPLEX |
| | MODULE_421, MODULE_421 WANG_METASTASIS_OF_BREAST_CANC WEST_ADRENOCORTICAL_TUMOR_MAI GSE2585_CD80_HIGH_VS_LOW_AIRE_KO | CER_ESR1_UP, WANG_METASTASIS_OF_BREAST_CANCER_ESR1_UP RKERS_UP, WEST_ADRENOCORTICAL_TUMOR_MARKERS_UP O_MTEC_DN, GSE2585_CD80_HIGH_VS_LOW_AIRE_KO_MTEC_DN |
| | GSE17721_LPS_VS_POLYIC_2H_BMDC_UP GO_REGULATION_OF_ADHERENS_JUNC | SPORT, GO_REGULATION_OF_GLUCOSE_TRANSPORT P, GSE17721_LPS_VS_POLYIC_2H_BMDC_UP CTION_ORGANIZATION, GO_REGULATION_OF_ADHERENS_JUNCTION_ORGANIZATION DHESION, GO_REGULATION_OF_CELL_MATRIX_ADHESION |
| | YAATNANRNNNCAG_UNKNOWN, YAA GO_CONDENSED_CHROMOSOME_OUTE CCAWNWWNNNGGC_UNKNOWN, CCA | ER_KINETOCHORE, GO_CONDENSED_CHROMOSOME_OUTER_KINETOCHORE |
| | GO_NUCLEAR_INCLUSION_BODY, GO_N GO_LAMELLIPODIUM_ASSEMBLY, GO_L GO_INCLUSION_BODY, GO_INCLUSION_ | NUCLEAR_INCLUSION_BODY LAMELLIPODIUM_ASSEMBLY |
| | GO_POSITIVE_REGULATION_OF_CELL_JI | TIVITY _CYCLE_G2_M_PHASE_TRANSITION, GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION UNCTION_ASSEMBLY, GO_POSITIVE_REGULATION_OF_CELL_JUNCTION_ASSEMBLY RATION, GO_INNER_CELL_MASS_CELL_PROLIFERATION |
| | GO_CONDENSED_NUCLEAR_CHROMOS GO_REGULATION_OF_ATTACHMENT_O STAMBOLSKY_TARGETS_OF_MUTATED_ | SOME_CENTROMERIC_REGION, GO_CONDENSED_NUCLEAR_CHROMOSOME_CENTROMERIC_REGION OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GO_REGULATION_OF_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE _TP53_UP, STAMBOLSKY_TARGETS_OF_MUTATED_TP53_UP |
| | GSE14350_TREG_VS_TEFF_IN_IL2RB_KO_I | |
| | _ | _DEPENDENT_DNA_REPLICATION, GO_NEGATIVE_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION 45_TREG_VS_TCONV_UP VA_PROX1_TARGETS_UP |
| | GO_BLASTOCYST_GROWTH, GO_BLASTO | MBLY_INVOLVED_IN_EXECUTION_PHASE_OF_APOPTOSIS, GO_CELLULAR_COMPONENT_DISASSEMBLY_INVOLVED_IN_EXECUTION_PHASE_OF_APOPTOSIS |
| | REACTOME_KINESINS, REACTOME_KINE CHEOK_RESPONSE_TO_MERCAPTOPURI GO_POSITIVE_REGULATION_OF_ADHER GO_3_5_DNA_HELICASE_ACTIVITY, GO_ | ESINS INE_DN, CHEOK_RESPONSE_TO_MERCAPTOPURINE_DN RENS_JUNCTION_ORGANIZATION, GO_POSITIVE_REGULATION_OF_ADHERENS_JUNCTION_ORGANIZATION _3_5_DNA_HELICASE_ACTIVITY |
| | BIOCARTA_PDGF_PATHWAY, BIOCARTA TAXCREB_01, TAXCREB_01 GO_HISTONE_PHOSPHORYLATION, GO_ GO_SITE_OF_DOUBLE_STRAND_BREAK, | A_PDGF_PATHWAY _HISTONE_PHOSPHORYLATION GO_SITE_OF_DOUBLE_STRAND_BREAK |
| | ST_FAS_SIGNALING_PATHWAY, ST_FAS_ GO_CELL_AGING, GO_CELL_AGING REACTOME_SYNTHESIS_OF_PIPS_AT_TH | HE_PLASMA_MEMBRANE, REACTOME_SYNTHESIS_OF_PIPS_AT_THE_PLASMA_MEMBRANE |
| | BILANGES_SERUM_SENSITIVE_VIA_TSC2 MATZUK_MEIOTIC_AND_DNA_REPAIR, | EGION_VS_WHOLE_CORTEX_THYMUS_DN, GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_CORTEX_THYMUS_DN 2, BILANGES_SERUM_SENSITIVE_VIA_TSC2 . MATZUK_MEIOTIC_AND_DNA_REPAIR |
| | GO_REGULATION_OF_PROTEIN_EXPORT GO_NEGATIVE_REGULATION_OF_MITO | [_RECOMBINATION L_TGFB_TREATED_DN, GSE21670_STAT3_KO_VS_WT_CD4_TCELL_TGFB_TREATED_DN T_FROM_NUCLEUS, GO_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS PTIC_NUCLEAR_DIVISION, GO_NEGATIVE_REGULATION_OF_MITOTIC_NUCLEAR_DIVISION S, GO_EXECUTION_PHASE_OF_APOPTOSIS |
| | ZHANG_ADIPOGENESIS_BY_BMP7, ZHANG_CO_CENTROSOME_DUPLICATION, GO_CENTA_GO_PATHWAY, BIOCARTA_GO_PATHWAY, BIOCARTA_GO_CENTA | NG_ADIPOGENESIS_BY_BMP7 CENTROSOME_DUPLICATION |
| | BIOCARTA_EGF_PATHWAY, BIOCARTA_chr2p16, chr2p16 BENPORATH_ES_2, BENPORATH_ES_2 | _EGF_PATHWAY |
| | BIOCARTA_IGF1_PATHWAY, BIOCARTA_GO_NEGATIVE_REGULATION_OF_CELL_GO_NEGATIVE_REGULATION_OF_NUCL | SS, GO_ONE_CARBON_METABOLIC_PROCESS _IGF1_PATHWAY _MATRIX_ADHESION, GO_NEGATIVE_REGULATION_OF_CELL_MATRIX_ADHESION LEAR_DIVISION, GO_NEGATIVE_REGULATION_OF_NUCLEAR_DIVISION JOMA_VS_DUCTAL_NORMAL_UP, TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_DUCTAL_NORMAL_UP |
| | MODULE_552, MODULE_552 | IFICANTLY, DING_LUNG_CANCER_MUTATED_SIGNIFICANTLY _TPO_PATHWAY |
| | KEGG_TGF_BETA_SIGNALING_PATHWA RAY_TUMORIGENESIS_BY_ERBB2_CDC25 GO_TETRAHYDROFOLATE_METABOLIC_ | _TRANSLOCATION AY, KEGG_TGF_BETA_SIGNALING_PATHWAY 5A_UP, RAY_TUMORIGENESIS_BY_ERBB2_CDC25A_UP _PROCESS, GO_TETRAHYDROFOLATE_METABOLIC_PROCESS LL_DN, GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_DN |
| | PID_CIRCADIAN_PATHWAY, PID_CIRCAGO_RHO_PROTEIN_SIGNAL_TRANSDUCTORS GO_NEGATIVE_REGULATION_OF_CELL_GO_INSULIN_RECEPTOR_BINDING, GO_INSULIN_RECEPTOR_BINDING, GO_IN | ADIAN_PATHWAY CTION, GO_RHO_PROTEIN_SIGNAL_TRANSDUCTION _JUNCTION_ASSEMBLY, GO_NEGATIVE_REGULATION_OF_CELL_JUNCTION_ASSEMBLY |
| | · · · · · · · · · · · · · · · · · · · | INSULIN_RECEPTOR_BINDING ON, GO_RECIPROCAL_DNA_RECOMBINATION D_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION, GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION |
| | GO_REGULATION_OF_DOUBLE_STRAND MODULE_320, MODULE_320 | |
| | GO_RECIPROCAL_DNA_RECOMBINATION GO_REGULATION_OF_DOUBLE_STRAND MODULE_320, MODULE_320 AMIT_EGF_RESPONSE_40_MCF10A, AMIT REACTOME_CDC6_ASSOCIATION_WITH REACTOME_G2_M_DNA_DAMAGE_CHECOME_CO_PROTEIN_LOCALIZATION_TO_CHRO | T_EGF_RESPONSE_40_MCF10A I_THE_ORC_ORIGIN_COMPLEX, REACTOME_CDC6_ASSOCIATION_WITH_THE_ORC_ORIGIN_COMPLEX CCKPOINT, REACTOME_G2_M_DNA_DAMAGE_CHECKPOINT OMOSOME_CENTROMERIC_REGION, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_CENTROMERIC_REGION |
| | GO_RECIPROCAL_DNA_RECOMBINATION GO_REGULATION_OF_DOUBLE_STRAND MODULE_320, MODULE_320 AMIT_EGF_RESPONSE_40_MCF10A, AMIT REACTOME_CDC6_ASSOCIATION_WITH REACTOME_G2_M_DNA_DAMAGE_CHECOME GO_PROTEIN_LOCALIZATION_TO_CHROUN REACTOME_ASSOCIATION_OF_LICENSIN BIOCARTA_PAR1_PATHWAY, BIOCARTA chr10p13, chr10p13 chrxq27, chrxq27 | T_EGF_RESPONSE_40_MCF10A I_THE_ORC_ORIGIN_COMPLEX, REACTOME_CDC6_ASSOCIATION_WITH_THE_ORC_ORIGIN_COMPLEX ICKPOINT, REACTOME_G2_M_DNA_DAMAGE_CHECKPOINT OMOSOME_CENTROMERIC_REGION, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_CENTROMERIC_REGION NG_FACTORS_WITH_THE_PRE_REPLICATIVE_COMPLEX, REACTOME_ASSOCIATION_OF_LICENSING_FACTORS_WITH_THE_PRE_REPLICATIVE_COMPLEX A_PAR1_PATHWAY |
| | GO_RECIPROCAL_DNA_RECOMBINATION GO_REGULATION_OF_DOUBLE_STRAND MODULE_320, MODULE_320 AMIT_EGF_RESPONSE_40_MCF10A, AMIT REACTOME_CDC6_ASSOCIATION_WITH REACTOME_G2_M_DNA_DAMAGE_CHECOME GO_PROTEIN_LOCALIZATION_TO_CHROUND REACTOME_ASSOCIATION_OF_LICENSIN BIOCARTA_PAR1_PATHWAY, BIOCARTA chr10p13, chr10p13 chrxq27, chrxq27 GO_CHROMOSOME_SEPARATION, GO_COMECOMECOMECOMECOMECOMECOMECOMECOMECOME | T_EGF_RESPONSE_40_MCF10A I_THE_ORC_ORIGIN_COMPLEX, REACTOME_CDC6_ASSOCIATION_WITH_THE_ORC_ORIGIN_COMPLEX ICKPOINT, REACTOME_G2_M_DNA_DAMAGE_CHECKPOINT OMOSOME_CENTROMERIC_REGION, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_CENTROMERIC_REGION NG_FACTORS_WITH_THE_PRE_REPLICATIVE_COMPLEX, REACTOME_ASSOCIATION_OF_LICENSING_FACTORS_WITH_THE_PRE_REPLICATIVE_COMPLEX A_PAR1_PATHWAY CHROMOSOME_SEPARATION MIDZONE YY COR_MAINTENANCE_OF_CELL_POLARITY, GO_REGULATION_OF_ESTABLISHMENT_OR_MAINTENANCE_OF_CELL_POLARITY |
| | GO_RECIPROCAL_DNA_RECOMBINATION GO_REGULATION_OF_DOUBLE_STRAND MODULE_320, MODULE_320 AMIT_EGF_RESPONSE_40_MCF10A, AMIT REACTOME_CDC6_ASSOCIATION_WITH REACTOME_G2_M_DNA_DAMAGE_CHE GO_PROTEIN_LOCALIZATION_TO_CHROUSE REACTOME_ASSOCIATION_OF_LICENSIN BIOCARTA_PAR1_PATHWAY, BIOCARTA chr10p13, chr10p13 chrxq27, chrxq27 GO_CHROMOSOME_SEPARATION, GO_C GO_SPINDLE_MIDZONE, GO_SPINDLE_M ST_GA13_PATHWAY, ST_GA13_PATHWA GO_REGULATION_OF_ESTABLISHMENT_ KUMAR_AUTOPHAGY_NETWORK, KUM GO_EXTRINSIC_APOPTOTIC_SIGNALING YOKOE_CANCER_TESTIS_ANTIGENS, YO GO_REGULATION_OF_CIRCADIAN_RHY | T_EGF_RESPONSE_40_MCF10A L_THE_ORC_ORIGIN_COMPLEX, REACTOME_CDC6_ASSOCIATION_WITH_THE_ORC_ORIGIN_COMPLEX ICKPOINT, REACTOME_G2_M_DNA_DAMAGE_CHECKPOINT OMOSOME_CENTROMERIC_REGION, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_CENTROMERIC_REGION NG_FACTORS_WITH_THE_PRE_REPLICATIVE_COMPLEX, REACTOME_ASSOCIATION_OF_LICENSING_FACTORS_WITH_THE_PRE_REPLICATIVE_COMPLEX A_PAR1_PATHWAY CHROMOSOME_SEPARATION MIDZONE NY COR_MAINTENANCE_OF_CELL_POLARITY, GO_REGULATION_OF_ESTABLISHMENT_OR_MAINTENANCE_OF_CELL_POLARITY IAR_AUTOPHAGY_NETWORK G_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS, GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS |