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GO_NEGATIVE_REGULATION_OF_MITOTIC_CELL_CYCLE, GO_NEGATIVE_REGULATION_OF_MITOTIC_CELL_CYCLE
GO DNA INTEGRITY CHECKPOINT, GO DNA INTEGRITY CHECKPOINT
GO_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS, GO_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS
WHITFIELD_CELL_CYCLE_S, WHITFIELD_CELL_CYCLE_S
GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_PHASE_TRANSITION, GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_PHASE_TRANSITION
GSE1925 3H VS 24H IFNG STIM IFNG PRIMED MACROPHAGE DN, GSE1925 3H VS 24H IFNG STIM IFNG PRIMED MACROPHAGE DN
GO_PROTEIN_UBIQUITINATION_INVOLVED_IN_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_PROTEIN_UBIQUITINATION_INVOLVED_IN_UBIQUITIN_DEPENDENT_PROTEIN_CATA
GO_GENE_SILENCING_BY_RNA, GO_GENE_SILENCING_BY_RNA
GO_NUCLEASE_ACTIVITY, GO_NUCLEASE_ACTIVITY
GO_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION, GO_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION
WHITFIELD_CELL_CYCLE_G1_S, WHITFIELD_CELL_CYCLE_G1_S
GO DOUBLE STRAND BREAK REPAIR, GO DOUBLE STRAND BREAK REPAIR
HALLMARK_P53_PATHWAY, HALLMARK_P53_PATHWAY
SMIRNOV_RESPONSE_TO_IR_6HR_UP, SMIRNOV_RESPONSE_TO_IR_6HR_UP
GO_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS, GO_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS
HALLMARK_APOPTOSIS, HALLMARK_APOPTOSIS
GSE3982_MAST_CELL_VS_TH1_DN, GSE3982_MAST_CELL_VS_TH1_DN
GO_PROTEIN_C_TERMINUS_BINDING, GO_PROTEIN_C_TERMINUS_BINDING
GO_PROTEIN_DNA_COMPLEX, GO_PROTEIN_DNA_COMPLEX
GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT, GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT
PID_P53_DOWNSTREAM_PATHWAY, PID_P53_DOWNSTREAM_PATHWAY
GO_ACTIVATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY, GO_ACTIVATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY
GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION, GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION
GO_RESPONSE_TO_IONIZING_RADIATION, GO_RESPONSE_TO_IONIZING_RADIATION
GO_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
CGTSACG_PAX3_B, CGTSACG_PAX3_B
GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMC_AT_DX_DN, GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMC_AT_DX_DN
MORF_DDX11, MORF_DDX11
KOKKINAKIS_METHIONINE_DEPRIVATION_48HR_UP, KOKKINAKIS_METHIONINE_DEPRIVATION_48HR_UP
GSE26351_WNT_VS_BMP_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_UP, GSE26351_WNT_VS_BMP_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_UP
GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE, GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE
GO_NEGATIVE_REGULATION_OF_CELLULAR_CATABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_CELLULAR_CATABOLIC_PROCESS
GO_REGULATION_OF_CELL_CYCLE_ARREST, GO_REGULATION_OF_CELL_CYCLE_ARREST
GSE11961_FOLLICULAR_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_DN, GSE11961_FOLLICULAR_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_DN
MACAEVA_PBMC_RESPONSE_TO_IR, MACAEVA_PBMC_RESPONSE_TO_IR
GSE1460_CORD_VS_ADULT_BLOOD_NAIVE_CD4_TCELL_UP, GSE1460_CORD_VS_ADULT_BLOOD_NAIVE_CD4_TCELL_UP
PID_AR_PATHWAY, PID_AR_PATHWAY
GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
GSE22611_MUTANT_NOD2_TRANSDUCED_VS_CTRL_HEK293T_STIMULATED_WITH_MDP_6H_DN, GSE22611_MUTANT_NOD2_TRANSDUCED_VS_CTRL_HEK293T_STIMULATED_WITH_MDP_6H_DN
GAZDA DIAMOND BLACKFAN ANEMIA PROGENITOR DN, GAZDA DIAMOND BLACKFAN ANEMIA PROGENITOR DN
KOKKINAKIS_METHIONINE_DEPRIVATION_96HR_UP, KOKKINAKIS_METHIONINE_DEPRIVATION_96HR_UP
GSE3039 ALPHAALPHA CD8 TCELL VS B1 BCELL DN, GSE3039 ALPHAALPHA CD8 TCELL VS B1 BCELL DN
REACTOME_ACTIVATED_TLR4_SIGNALLING, REACTOME_ACTIVATED_TLR4_SIGNALLING
REACTOME_MEIOSIS, REACTOME_MEIOSIS
DACOSTA_UV_RESPONSE_VIA_ERCC3_COMMON_UP, DACOSTA_UV_RESPONSE_VIA_ERCC3_COMMON_UP
GO_EXONUCLEASE_ACTIVITY, GO_EXONUCLEASE_ACTIVITY
TAANNYSGCG_UNKNOWN, TAANNYSGCG_UNKNOWN
GSE11961_MEMORY_BCELL_DAY40_VS_GERMINAL_CENTER_BCELL_DAY40_UP, GSE11961_MEMORY_BCELL_DAY40_VS_GERMINAL_CENTER_BCELL_DAY40_UP
FIRESTEIN PROLIFERATION, FIRESTEIN PROLIFERATION
GO REGULATION OF CYCLIN DEPENDENT PROTEIN KINASE ACTIVITY, GO REGULATION OF CYCLIN DEPENDENT PROTEIN KINASE ACTIVITY
GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS
JACKSON_DNMT1_TARGETS_UP, JACKSON_DNMT1_TARGETS_UP
GO_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE, GO_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE
GO_EXECUTION_PHASE_OF_APOPTOSIS, GO_EXECUTION_PHASE_OF_APOPTOSIS
GSE21927_UNTREATED_VS_GMCSF_IL6_TREATED_BONE_MARROW_UP, GSE21927_UNTREATED_VS_GMCSF_IL6_TREATED_BONE_MARROW_UP
KEGG SMALL CELL LUNG CANCER, KEGG SMALL CELL LUNG CANCER
GO_3_5_EXONUCLEASE_ACTIVITY, GO_3_5_EXONUCLEASE_ACTIVITY
GO_POSITIVE_REGULATION_OF_CELL_CYCLE_ARREST, GO_POSITIVE_REGULATION_OF_CELL_CYCLE_ARREST
CCAWYNNGAAR_UNKNOWN, CCAWYNNGAAR_UNKNOWN
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC
GO_JNK_CASCADE, GO_JNK_CASCADE
FRIDMAN_SENESCENCE_UP, FRIDMAN_SENESCENCE_UP
GO_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS
GO_POSTTRANSCRIPTIONAL_GENE_SILENCING, GO_POSTTRANSCRIPTIONAL_GENE_SILENCING
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_52H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_52H_UP
GO_RECEPTOR_SIGNALING_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY, GO_RECEPTOR_SIGNALING_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY
GO_CELLULAR_RESPONSE_TO_IONIZING_RADIATION, GO_CELLULAR_RESPONSE_TO_IONIZING_RADIATION
GO_REGULATION_OF_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_REGULATION_OF_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_
GO_REGULATION_OF_JNK_CASCADE, GO_REGULATION_OF_JNK_CASCADE
CHIARADONNA_NEOPLASTIC_TRANSFORMATION_KRAS_CDC25_UP, CHIARADONNA_NEOPLASTIC_TRANSFORMATION_KRAS_CDC25_UP
REACTOME_MEIOTIC_SYNAPSIS, REACTOME_MEIOTIC_SYNAPSIS
GO_SITE_OF_DOUBLE_STRAND_BREAK, GO_SITE_OF_DOUBLE_STRAND_BREAK
GHANDHI_DIRECT_IRRADIATION_UP, GHANDHI_DIRECT_IRRADIATION_UP
GSE13411_IGM_MEMORY_BCELL_VS_PLASMA_CELL_DN, GSE13411_IGM_MEMORY_BCELL_VS_PLASMA_CELL_DN
GO_NEGATIVE_REGULATION_OF_ERBB_SIGNALING_PATHWAY, GO_NEGATIVE_REGULATION_OF_ERBB_SIGNALING_PATHWAY
E2F_01, E2F_01
WEIGEL_OXIDATIVE_STRESS_BY_HNE_AND_TBH, WEIGEL_OXIDATIVE_STRESS_BY_HNE_AND_TBH
GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION, GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION
REACTOME SIGNALING BY SCF KIT, REACTOME SIGNALING BY SCF KIT
REACTOME_PI3K_EVENTS_IN_ERBB2_SIGNALING, REACTOME_PI3K_EVENTS_IN_ERBB2_SIGNALING
GO_ACTIVATING_TRANSCRIPTION_FACTOR_BINDING, GO_ACTIVATING_TRANSCRIPTION_FACTOR_BINDING
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CLE CHECKPOINT, GO CELL CYCLE CHECKPOINT

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REACTOME_GAB1_SIGNALOSOME, REACTOME_GAB1_SIGNALOSOME
GO_CELLULAR_COMPONENT_DISASSEMBLY_INVOLVED_IN_EXECUTION_PHASE_OF_APOPTOSIS, GO_CELLULAR_COMPONENT_DISASSEMBLY_INVOLVED_IN_EXECUTION_PHASE_OF_APOPTOSIS
FRIDMAN IMMORTALIZATION DN, FRIDMAN IMMORTALIZATION DN
GO_REGULATION_OF_FIBROBLAST_PROLIFERATION, GO_REGULATION_OF_FIBROBLAST_PROLIFERATION
BONCI TARGETS OF MIR15A AND MIR16 1, BONCI TARGETS OF MIR15A AND MIR16 1
KRIGE_AMINO_ACID_DEPRIVATION, KRIGE_AMINO_ACID_DEPRIVATION
MORF_CNTN1, MORF_CNTN1
GO_DIGESTIVE_SYSTEM_DEVELOPMENT, GO_DIGESTIVE_SYSTEM_DEVELOPMENT
PID_EPO_PATHWAY, PID_EPO_PATHWAY
GO_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS, GO_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS
WARTERS_RESPONSE_TO_IR_SKIN, WARTERS_RESPONSE_TO_IR_SKIN
GO_ACTIVATION_OF_MAPK_ACTIVITY, GO_ACTIVATION_OF_MAPK_ACTIVITY
GO_SOMATIC_DIVERSIFICATION_OF_IMMUNOGLOBULINS, GO_SOMATIC_DIVERSIFICATION_OF_IMMUNOGLOBULINS
SCIAN_INVERSED_TARGETS_OF_TP53_AND_TP73_UP, SCIAN_INVERSED_TARGETS_OF_TP53_AND_TP73_UP
REACTOME_PI3K_EVENTS_IN_ERBB4_SIGNALING, REACTOME_PI3K_EVENTS_IN_ERBB4_SIGNALING
REACTOME_TRAF6_MEDIATED_INDUCTION_OF_TAK1_COMPLEX, REACTOME_TRAF6_MEDIATED_INDUCTION_OF_TAK1_COMPLEX
GO_DNA_POLYMERASE_BINDING, GO_DNA_POLYMERASE_BINDING
GO_SKELETAL_SYSTEM_MORPHOGENESIS, GO_SKELETAL_SYSTEM_MORPHOGENESIS
GO REGULATION OF MEIOTIC CELL CYCLE, GO REGULATION OF MEIOTIC CELL CYCLE
GO_RESPONSE_TO_KETONE, GO_RESPONSE_TO_KETONE
REACTOME SIGNALING BY ERBB4, REACTOME SIGNALING BY ERBB4
GO_NEGATIVE_REGULATION_OF_CELL_CELL_ADHESION, GO_NEGATIVE_REGULATION_OF_CELL_CELL_ADHESION
REACTOME_PIP3_ACTIVATES_AKT_SIGNALING, REACTOME_PIP3_ACTIVATES_AKT_SIGNALING
GO_REGULATION_OF_DNA_DAMAGE_CHECKPOINT, GO_REGULATION_OF_DNA_DAMAGE_CHECKPOINT
GO_RESPONSE_TO_VITAMIN, GO_RESPONSE_TO_VITAMIN
SESTO_RESPONSE_TO_UV_C3, SESTO_RESPONSE_TO_UV_C3
GO_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION, GO_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION
MATZUK_SPERMATOZOA, MATZUK_SPERMATOZOA
REACTOME_IRAK2_MEDIATED_ACTIVATION_OF_TAK1_COMPLEX_UPON_TLR7_8_OR_9_STIMULATION, REACTOME_IRAK2_MEDIATED_ACTIVATION_OF_TAK1_COMPLEX_UPON_TLR7_8_OR_9_STIMULATION_OF_TAK1_OR_9_STIMULATION_OF_TAK1_OR_9_STIMULATION_OF_TAK1_OR_9_STIMULATION_OF_TAK1_OR_9_STIMULATION_OF_TAK1_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATION_OR_9_STIMULATI
DING_LUNG_CANCER_MUTATED_SIGNIFICANTLY, DING_LUNG_CANCER_MUTATED_SIGNIFICANTLY
SHIN_B_CELL_LYMPHOMA_CLUSTER_8, SHIN_B_CELL_LYMPHOMA_CLUSTER_8
MORF_PTPN9, MORF_PTPN9
GO REGULATION OF MULTICELLULAR ORGANISMAL METABOLIC PROCESS, GO REGULATION OF MULTICELLULAR ORGANISMAL METABOLIC PROCESS
DING_LUNG_CANCER_MUTATED_FREQUENTLY, DING_LUNG_CANCER_MUTATED_FREQUENTLY
GSE28449_WT_VS_LRF_KO_GERMINAL_CENTER_BCELL_DN, GSE28449_WT_VS_LRF_KO_GERMINAL_CENTER_BCELL_DN
GSE6259_FLT3L_INDUCED_DEC205_POS_DC_VS_BCELL_UP, GSE6259_FLT3L_INDUCED_DEC205_POS_DC_VS_BCELL_UP
GO_REGULATION_OF_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GO_REGULATION_OF_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE
GSE2585_CD80_HIGH_VS_LOW_AIRE_KO_MTEC_DN, GSE2585_CD80_HIGH_VS_LOW_AIRE_KO_MTEC_DN
NUNODA_RESPONSE_TO_DASATINIB_IMATINIB_DN, NUNODA_RESPONSE_TO_DASATINIB_IMATINIB_DN
BIOCARTA_VDR_PATHWAY, BIOCARTA_VDR_PATHWAY
MATZUK_MEIOTIC_AND_DNA_REPAIR, MATZUK_MEIOTIC_AND_DNA_REPAIR
GO_MAMMARY_GLAND_EPITHELIUM_DEVELOPMENT, GO_MAMMARY_GLAND_EPITHELIUM_DEVELOPMENT
GO_SKELETAL_MUSCLE_CELL_DIFFERENTIATION, GO_SKELETAL_MUSCLE_CELL_DIFFERENTIATION
ROYLANCE_BREAST_CANCER_16Q_COPY_NUMBER_UP, ROYLANCE_BREAST_CANCER_16Q_COPY_NUMBER_UP
SATO_SILENCED_BY_METHYLATION_IN_PANCREATIC_CANCER_2, SATO_SILENCED_BY_METHYLATION_IN_PANCREATIC_CANCER_2
GO_SYNAPTONEMAL_COMPLEX, GO_SYNAPTONEMAL_COMPLEX
BIOCARTA_TID_PATHWAY, BIOCARTA_TID_PATHWAY
NOJIMA_SFRP2_TARGETS_DN, NOJIMA_SFRP2_TARGETS_DN
REACTOME_ACTIVATION_OF_IRF3_IRF7_MEDIATED_BY_TBK1_IKK_EPSILON, REACTOME_ACTIVATION_OF_IRF3_IRF7_MEDIATED_BY_TBK1_IKK_EPSILON
GO_POSITIVE_REGULATION_OF_MULTICELLULAR_ORGANISMAL_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_MULTICELLULAR_ORGANISMAL_METABOLIC_PROCESS
GO_MEGAKARYOCYTE_DIFFERENTIATION, GO_MEGAKARYOCYTE_DIFFERENTIATION
MASSARWEH_RESPONSE_TO_ESTRADIOL, MASSARWEH_RESPONSE_TO_ESTRADIOL
GO_POSITIVE_REGULATION_OF_PROTEIN_ACETYLATION, GO_POSITIVE_REGULATION_OF_PROTEIN_ACETYLATION
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GO_RNA_POLYMERASE_II_ACTIVATING_TRANSCRIPTION_FACTOR_BINDING, GO_RNA_POLYMERASE_II_ACTIVATING_TRANSCRIPTION_FACTOR_BINDING

GO_NEGATIVE_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE, GO_NEGATIVE_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE

GO_REGULATION_OF_EXTRACELLULAR_MATRIX_ORGANIZATION, GO_REGULATION_OF_EXTRACELLULAR_MATRIX_ORGANIZATION

REACTOME_AKT_PHOSPHORYLATES_TARGETS_IN_THE_CYTOSOL, REACTOME_AKT_PHOSPHORYLATES_TARGETS_IN_THE_CYTOSOL

GO POSITIVE REGULATION OF DNA DEPENDENT DNA REPLICATION, GO POSITIVE REGULATION OF DNA DEPENDENT DNA REPLICATION

GO_POSITIVE_REGULATION_OF_FAT_CELL_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_FAT_CELL_DIFFERENTIATION

GO_MITOGEN_ACTIVATED_PROTEIN_KINASE_BINDING, GO_MITOGEN_ACTIVATED_PROTEIN_KINASE_BINDING

GO_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION, GO_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION

GO_HOMOLOGOUS_CHROMOSOME_SEGREGATION, GO_HOMOLOGOUS_CHROMOSOME_SEGREGATION

ZHU_SKIL_TARGETS_UP, ZHU_SKIL_TARGETS_UP

GCAAGGA_MIR502, GCAAGGA_MIR502

GO_SYNAPSIS, GO_SYNAPSIS

MURATA_VIRULENCE_OF_H_PILORI, MURATA_VIRULENCE_OF_H_PILORI GO_MYELOID_CELL_DEVELOPMENT, GO_MYELOID_CELL_DEVELOPMENT

SIMBULAN_PARP1_TARGETS_UP, SIMBULAN_PARP1_TARGETS_UP

STREICHER_LSM1_TARGETS_DN, STREICHER_LSM1_TARGETS_DN

PID_CIRCADIAN_PATHWAY, PID_CIRCADIAN_PATHWAY

BIOCARTA_IL6_PATHWAY, BIOCARTA_IL6_PATHWAY

GO_CARDIAC_ATRIUM_DEVELOPMENT, GO_CARDIAC_ATRIUM_DEVELOPMENT

GO_VENTRICULAR_SEPTUM_DEVELOPMENT, GO_VENTRICULAR_SEPTUM_DEVELOPMENT

GO_MAMMARY_GLAND_DUCT_MORPHOGENESIS, GO_MAMMARY_GLAND_DUCT_MORPHOGENESIS

TURJANSKI_MAPK8_AND_MAPK9_TARGETS, TURJANSKI_MAPK8_AND_MAPK9_TARGETS