

_DNA_REPAIR, REACTOME_DNA_REPAIR

GO_GENE_SILENCING_BY_RNA, GO_GENE_SILENCING_BY_RNA
GO_RNA_POLYMERASE_COMPLEX, GO_RNA_POLYMERASE_COMPLEX
GO_DNA_DIRECTED_RNA_POLYMERASE_II_HOLOENZYME, GO_DNA_DIRECTED_RNA_POLYMERASE_II_HOLOENZYME
GO_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GO_SNRNA_METABOLIC_PROCESS, GO_SNRNA_METABOLIC_PROCESS
REACTOME_TELOMERE_MAINTENANCE, REACTOME_TELOMERE_MAINTENANCE
GSE29618_PRE_VS_DAY7_FLU_VACCINE_BCELL_DN, GSE29618_PRE_VS_DAY7_FLU_VACCINE_BCELL_DN
REACTOME_NUCLEOTIDE_EXCISION_REPAIR, REACTOME_NUCLEOTIDE_EXCISION_REPAIR
GO_DNA_TEMPLATED_TRANSCRIPTION_INITIATION, GO_DNA_TEMPLATED_TRANSCRIPTION_INITIATION
GO_PML_BODY, GO_PML_BODY
GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE, GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE
GO_CARBOHYDRATE_DERIVATIVE_CATABOLIC_PROCESS, GO_CARBOHYDRATE_DERIVATIVE_CATABOLIC_PROCESS
GO_POSITIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC, GO_POSITIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC
GO_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GO_PROTEIN_N_TERMINUS_BINDING, GO_PROTEIN_N_TERMINUS_BINDING
GO_TRANSLESION_SYNTHESIS, GO_TRANSLESION_SYNTHESIS
AMUNDSON_GENOTOXIC_SIGNATURE, AMUNDSON_GENOTOXIC_SIGNATURE
GO_BASE_EXCISION_REPAIR, GO_BASE_EXCISION_REPAIR
GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
GO_IMMUNOGLOBULIN_PRODUCTION, GO_IMMUNOGLOBULIN_PRODUCTION
GO_PROTEIN_MONOUBIQUITINATION, GO_PROTEIN_MONOUBIQUITINATION
MACAEVA_PBMC_RESPONSE_TO_IR, MACAEVA_PBMC_RESPONSE_TO_IR
KEGG_BASAL_TRANSCRIPTION_FACTORS, KEGG_BASAL_TRANSCRIPTION_FACTORS
GO_SOMATIC_RECOMBINATION_OF_IMMUNOGLOBULIN_GENE_SEGMENTS, GO_SOMATIC_RECOMBINATION_OF_IMMUNOGLOBULIN_GENE_SEGMENTS
GO_ORGANOPHOSPHATE_CATABOLIC_PROCESS, GO_ORGANOPHOSPHATE_CATABOLIC_PROCESS
GO_RESPONSE_TO_X_RAY, GO_RESPONSE_TO_X_RAY
GO_POSITIVE_REGULATION_OF_ERBB_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_ERBB_SIGNALING_PATHWAY
GO_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GO_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
GO_MULTICELLULAR_ORGANISM_GROWTH, GO_MULTICELLULAR_ORGANISM_GROWTH
GSE24574_BCL6_LOW_TFH_VS_TCONV_CD4_TCELL_UP, GSE24574_BCL6_LOW_TFH_VS_TCONV_CD4_TCELL_UP
AMUNDSON_DNA_DAMAGE_RESPONSE_TP53, AMUNDSON_DNA_DAMAGE_RESPONSE_TP53
chr2q21, chr2q21
GO_RESPONSE_TO_LITHIUM_ION, GO_RESPONSE_TO_LITHIUM_ION
GO_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_RESULTING_IN_TRANSCRIPTION, GO_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_RESULTING_IN_TRANSCRIPTION
GO_RESPONSE_TO_FIBROBLAST_GROWTH_FACTOR, GO_RESPONSE_TO_FIBROBLAST_GROWTH_FACTOR
GO_REGULATION_OF_HISTONE_PHOSPHORYLATION, GO_REGULATION_OF_HISTONE_PHOSPHORYLATION
GO_NEGATIVE_REGULATION_OF_HISTONE_ACETYLTATION, GO_NEGATIVE_REGULATION_OF_HISTONE_ACETYLTATION
REACTOME_RESOLUTION_OF_AP_SITES_VIA_THE_SINGLE_NUCLEOTIDE_REPLACEMENT_PATHWAY, REACTOME_RESOLUTION_OF_AP_SITES_VIA_THE_SINGLE_NUCLEOTIDE_REPLACEMENT_PATHWAY
chr19q12, chr19q12