

GOBP_HISTONE_METHYLATION, GOBP_PROTEIN_METHYLATION

- GOBP_HISTONE_METHYLATION, GOBP_HISTONE_METHYLATION
- GOBP_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE, GOBP_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE
- GOBP_SPLICEOSOMAL_SNRNP_ASSEMBLY, GOBP_SPLICEOSOMAL_SNRNP_ASSEMBLY
- GOBP_PROTEIN_ACETYLATION, GOBP_PROTEIN_ACETYLATION
- GOBP_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GOBP_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
- GOBP_PEPTIDYL_LYSINE_ACETYLATION, GOBP_PEPTIDYL_LYSINE_ACETYLATION
- GOBP_PEPTIDYL_LYSINE_METHYLATION, GOBP_PEPTIDYL_LYSINE_METHYLATION
- GOBP_HISTONE_H4_ACETYLATION, GOBP_HISTONE_H4_ACETYLATION
- GOBP_REGULATION_OF_DNA_RECOMBINATION, GOBP_REGULATION_OF_DNA_RECOMBINATION
- GOBP_HISTONE_UBIQUITINATION, GOBP_HISTONE_UBIQUITINATION
- GOBP_HISTONE_H4_K5_ACETYLATION, GOBP_HISTONE_H4_K5_ACETYLATION
- GOBP_NON_RECOMBINATIONAL_REPAIR, GOBP_NON_RECOMBINATIONAL_REPAIR
- GOBP_HISTONE_H3_K4_METHYLATION, GOBP_HISTONE_H3_K4_METHYLATION
- GOBP_MITOCHONDRIAL_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT, GOBP_MITOCHONDRIAL_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
- GOBP_ENDODERM_DEVELOPMENT, GOBP_ENDODERM_DEVELOPMENT
- GOBP_RRNA_TRANSCRIPTION, GOBP_RRNA_TRANSCRIPTION
- GOBP_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMOLOGOUS_END_JOINING, GOBP_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMOLOGOUS_END_JOINING
- GOBP_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION, GOBP_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION
- GOBP_POSITIVE_REGULATION_OF_HISTONE_METHYLATION, GOBP_POSITIVE_REGULATION_OF_HISTONE_METHYLATION
- GOBP_REGULATION_OF_BILE_ACID_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_BILE_ACID_BIOSYNTHETIC_PROCESS
- GOBP_HISTONE_H4_K16_ACETYLATION, GOBP_HISTONE_H4_K16_ACETYLATION
- GOBP_LYMPHOCYTE_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_LYMPHOCYTE_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE
- GOBP_POSITIVE_REGULATION_OF_B_CELL_MEDIATED_IMMUNITY, GOBP_POSITIVE_REGULATION_OF_B_CELL_MEDIATED_IMMUNITY
- GOBP_SULFUR_AMINO_ACID_BIOSYNTHETIC_PROCESS, GOBP_SULFUR_AMINO_ACID_BIOSYNTHETIC_PROCESS
- GOBP_ASPARTATE_FAMILY_AMINO_ACID_BIOSYNTHETIC_PROCESS, GOBP_ASPARTATE_FAMILY_AMINO_ACID_BIOSYNTHETIC_PROCESS
- GOBP_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC, GOBP_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC
- GOBP_BLASTOCYST_GROWTH, GOBP_BLASTOCYST_GROWTH
- GOBP_HISTONE_H3_K9_TRIMETHYLATION, GOBP_HISTONE_H3_K9_TRIMETHYLATION
- GOBP_DNA_MODIFICATION, GOBP_DNA_MODIFICATION
- GOBP_PYRIMIDINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS, GOBP_PYRIMIDINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS
- GOBP_REGULATION_OF_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY
- GOBP_SNO_S_RNA_3_END_PROCESSING, GOBP_SNO_S_RNA_3_END_PROCESSING
- GOBP_REGULATION_OF_BILE_ACID_METABOLIC_PROCESS, GOBP_REGULATION_OF_BILE_ACID_METABOLIC_PROCESS
- GOBP_REGULATION_OF_RRNA_PROCESSING, GOBP_REGULATION_OF_RRNA_PROCESSING