N_METHYLATION, GOBP_PROTEIN_METHYLATION

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HP ABNORMALITY OF THE FRONTAL HAIRLINE, HP ABNORMALITY OF THE FRONTAL HAIRLINE
 ATGCAGT_MIR217, ATGCAGT_MIR217
 GOCC SITE OF DOUBLE STRAND BREAK, GOCC SITE OF DOUBLE STRAND BREAK
 GOMF N METHYLTRANSFERASE ACTIVITY, GOMF N METHYLTRANSFERASE ACTIVITY
 REACTOME PROTEIN UBIQUITINATION, REACTOME PROTEIN UBIQUITINATION
 GOMF_NUCLEAR_HORMONE_RECEPTOR_BINDING, GOMF_NUCLEAR_HORMONE_RECEPTOR_BINDING
 MIR1913, MIR1913
 GOBP POSITIVE REGULATION OF RESPONSE TO DNA DAMAGE STIMULUS, GOBP POSITIVE REGULATION OF RESPONSE TO DNA DAMAGE STIMULUS
 MIR6129, MIR6129
 MIR6133, MIR6133
 MIR6130, MIR6130
- HP BLEPHAROPHIMOSIS, HP BLEPHAROPHIMOSIS
GOBP_BLASTOCYST_DEVELOPMENT, GOBP_BLASTOCYST_DEVELOPMENT
 GOMF_LYSINE_N_METHYLTRANSFERASE_ACTIVITY, GOMF_LYSINE_N_METHYLTRANSFERASE_ACTIVITY
 REACTOME PKMTS METHYLATE HISTONE LYSINES, REACTOME PKMTS METHYLATE HISTONE LYSINES
 HP_ACUTE_LYMPHOBLASTIC_LEUKEMIA, HP_ACUTE_LYMPHOBLASTIC_LEUKEMIA
 HP ABNORMAL NASOLACRIMAL SYSTEM MORPHOLOGY, HP ABNORMAL NASOLACRIMAL SYSTEM MORPHOLOGY
 GOBP_PEPTIDYL_ARGININE_MODIFICATION, GOBP_PEPTIDYL_ARGININE_MODIFICATION
 GOBP SENSORY PERCEPTION OF PAIN, GOBP SENSORY PERCEPTION OF PAIN
GOBP_HISTONE_ARGININE_METHYLATION, GOBP_HISTONE_ARGININE_METHYLATION
GOBP_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMOLOGOUS_END_JOINING, GOBP_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMO.
 WP NO METABOLISM IN CYSTIC FIBROSIS, WP NO METABOLISM IN CYSTIC FIBROSIS
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