

methyltransferase activity, GO:0008168

- tRNA processing, GO:0008033
- tRNA binding, GO:0000049
- negative regulation of gene expression, GO:0045814
- regulation of megakaryocyte differentiation, GO:0045652
- stem cell population maintenance, GO:0019827
- chromatin organization, GO:0006325
- p53 binding, GO:0002039
- tRNA modification, GO:0006400
- histone-lysine N-methyltransferase activity, GO:0018024
- 7-methylguanosine mRNA capping, GO:0006370
- germ cell development, GO:0007281
- histone methyltransferase complex, GO:0035097
- tRNA methylation, GO:0030488
- RNA polymerase II core promoter sequence-specific DNA binding, GO:0000979
- histone methyltransferase activity (H3-K4 specific), GO:0042800
- methyl-CpG binding, GO:0008327
- histone H3-K4 methylation, GO:0051568
- histone H3-K4 trimethylation, GO:0080182
- protein-lysine N-methyltransferase activity, GO:0016279
- histone methylation, GO:0016571
- response to vitamin A, GO:0033189
- ubiquinone biosynthetic process, GO:0006744
- rRNA methylation, GO:0031167
- oxidoreductase activity, GO:0016706
- homeostasis of number of cells within a tissue, GO:0048873
- S-adenosylmethionine-dependent methyltransferase activity, GO:0008757
- promoter-specific chromatin binding, GO:1990841
- protein-arginine N-methyltransferase activity, GO:0016274
- positive regulation of histone H3-K4 methylation, GO:0051571
- RNA methylation, GO:0001510
- rRNA modification, GO:0000154
- negative regulation of G1/S transition of mitotic cell cycle, GO:2000134
- Set1C/COMPASS complex, GO:0048188
- retinoic acid receptor binding, GO:0042974
- lysine-acetylated histone binding, GO:0070577
- hair follicle maturation, GO:0048820
- RNA methyltransferase activity, GO:0008173
- tRNA wobble uridine modification, GO:0002098
- peptidyl-lysine methylation, GO:0018022
- regulation of protein binding, GO:0043393
- one-carbon metabolic process, GO:0006730
- peptidyl-arginine N-methylation, GO:0035246
- ferrous iron binding, GO:0008198
- fertilization, GO:0009566
- histone H4-K20 trimethylation, GO:0034773
- histone demethylase activity (H3-K36 specific), GO:0051864
- histone H3-K36 demethylation, GO:0070544