

DDSAAADYCPPPKRLKTN--CYNNGKDRGEEDQSREOMALDVTNPKSLEDSCLSCGRKNPVSFHPLFEGGLCQTCRDRFLELFYMYDDDGYSYCTVCC 466
 IDFAASEYSTPPKRLKTN--SYG--GKDRGDEEESRERMASDVTNNGKLEDRCLSCGKKNPVSFHPLFEGGLCQSCRDRFLELFYMYDEEDGYQSYCTVCC 488
 NDSAASE-SPPPKRLKTN--SYG--GKDRGDEEESRERMASEVTNNGKLEDRCLSCGKKNPVSFHPLFEGGLCQSCRDRFLELFYMYDEEDGYQSYCTVCC 488
 EEVTVDLSLPPPKRLKPS--CVNNKKEVEEDQSREKMSVSLKNNKSNPELSCIAAGRRNPPSFHPLFEGGLCHTCRDRFLELFYMYDDDGYSYCTVCC 386
 LESSPPEFP--SAKRARLP--LNKAKPGIEEVSREQMVNEVLKMHRSIEEFCLSCGKTRVATFHPLFEGGLCLTCKDAYLENSYMYDDDGYSYCTVCC 446

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GKELLLCSNA--SCCRFCVCECLEVLVGRGSSAKAKEQEPWNCYMCQPORSYGVLRQXQDSSRLQDFFTSK--GEYDAPKIYPVPPAKRRRPIRVLSLFD 576
 GRELLLCNSNTSCCRFCVCECLEVLVGTGTAAEAKLQEPWSCYMCLPQRCHGVLRRRKDNVRLQAFFTSDT--GLEYEAPKLYPAIPAARRRPIRVLSLFD 582
 GRELLLCNSNTSCCRFCVCECLEVLVGTGTAAEAKLQEPWSCYMCLPQRCHGVLRRRKDNVRLQAFFTSDT--GLEYEAPKLYPAIPAARRRPIRVLSLFD 575
 GRELLLCNSNTSCCRFCVCECLEVLVGTGTAAEAKLQEPWSCYMCLPQRCHGVLRRRKDNVRLQAFFTSDT--GLEYEAPKLYPAIPAARRRPIRVLSLFD 581
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 GRELLLCNSNTSCCRFCVCECLEVLVGTGTAAEAKLQEPWSCYMCLPQRCHGVLRRRKDNVRLQAFFTSDT--GLEYEAPKLYPAIPAARRRPIRVLSLFD 565
 GRELLLCNSNTSCCRFCVCECLEVLVGTGTAAEAKLQEPWSCYMCLPQRCHGVLRRRKDNVRLQAFFTSDT--GLEYEAPKLYPAIPAARRRPIRVLSLFD 588
 GRELLLCNSNTSCCRFCVCECLEVLVGTGTAAEAKLQEPWSCYMCLPQRCHGVLRRRKDNVRLQAFFTSDT--GLEYEAPKLYPAIPAARRRPIRVLSLFD 588
 GRELLLCNSNTSCCRFCVCECLEVLVGTGTAAEAKLQEPWSCYMCLPQRCHGVLRRRKDNVRLQAFFTSDT--GLEYEAPKLYPAIPAARRRPIRVLSLFD 485
 GRELLLCNSNTSCCRFCVCECLEVLVGTGTAAEAKLQEPWSCYMCLPQRCHGVLRRRKDNVRLQAFFTSDT--GLEYEAPKLYPAIPAARRRPIRVLSLFD 545

DNA Methylase domain

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GIATGYLVKDLGIQVEKYIASEICEDPLAVGTVRHEGNIITYVH--DVRNITKRNIEEWGPFDLVIGGSPCNDLSLASPERKAPVGEAAWPHGAGSPQAA 674
 GIATGYLVKELGKIKVGYVASEVCEESIAVGTVKHEGNIKYV--DVRNITKRNIEEWGPFDLVIGGSPCNDLSNVNPAKGLY----- 665
 GIATGYLVKELGKIKVGYVASEVCEESIAVGTVKHEGNIKYV--DVRNITKRNIEEWGPFDLVIGGSPCNDLSNVNPAKGLY----- 658
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 GIATGYLVKELGKIKVGYVASEVCEESIAVGTVKHEGNIKYV--DVRNITKRNIEEWGPFDLVIGGSPCNDLSNVNPAKGLY----- 649
 GIATGYLVKDLGIQVEKYIASEICEDPLAVGTVRHEGNIITYVH--DVRNITKRNIEEWGPFDLVIGGSPCNDLSLASPERKAPVGEAAWPHGAGSPQAA 671
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 ----- EGTGRLFFEFYHLLNYSRPKEGDDRPFFWMFENVVAMKVGDKRDISR 713
 ----- EGTGRLFFEFYHLLNYSRPKEGDDRPFFWMFENVVAMKVGDKRDISR 906
 ----- EGTGRLFFEFYHLLNYSRKEGDDRPFFWMFENVVAMQVGDKRDISR 712
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 ----- EGTGRLFFEFYHLLNYSRPKEGDDRPFFWMFENVVAMKVGDKRDISR 719
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 ----- EGTGRLFFEFYHLLNYSRPKEGDDRPFFWMFENVVAMRVNDKRDISR 616
 ----- EGTGRLFFEFYHLLNYSRPKEGDDRPFFWMFENVVAMRVNDKRDISR 676

LECNPVMIDAIKISAHRARYFWGNLPGMN-----RIFGFPV 811
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HCTDVSNSISRGARQKLLGRSWSIPVIOHLFSPLKDYFACE 851aa, chickenDNMT3B
 HYTDVSNMGRGARQKLLGRSWSVPVIRHLFAPLKDYFACE 853aa, humanDNMT3B
 HYTDVSNMGRGARQKLLGRSWSVPVIRHLFAPLKDYFACE 1046aa, chimpanzeeDNMT3B
 HYTDVSNMGRVARQKLPGRSWSVPVIRHLFAPLKDYFACE 852aa, pigDNMT3B
 HYTDVSNMGRVARQKLLGRSWSVPVIRHLFAPLKDYFACE 844aa, cattleDNMT3B
 HYTDVSNMGRGARQKLLGRSWSVPVIRHLFAPLKDYFACE 774aa, horseDNMT3B
 HYTDVSNMGRGARQKLLGRSWSVPVIRHLFAPLKDYFACE 859aa, ratDNMT3B
 HYTDVSNMGRGARQKLLGRSWSVPVIRHLFAPLKDYFACE 859aa, mouseDNMT3B
 HYTDVSNMGRGARQKLLGRSWSVPVIRHLFAPLKDYFACE 756aa, opossumDNMT3B
 HYTDVSNMGRGARQKLLGRSWSVPVIRHLFAPLKDYFACE 816aa, zebrafishDNMT3B