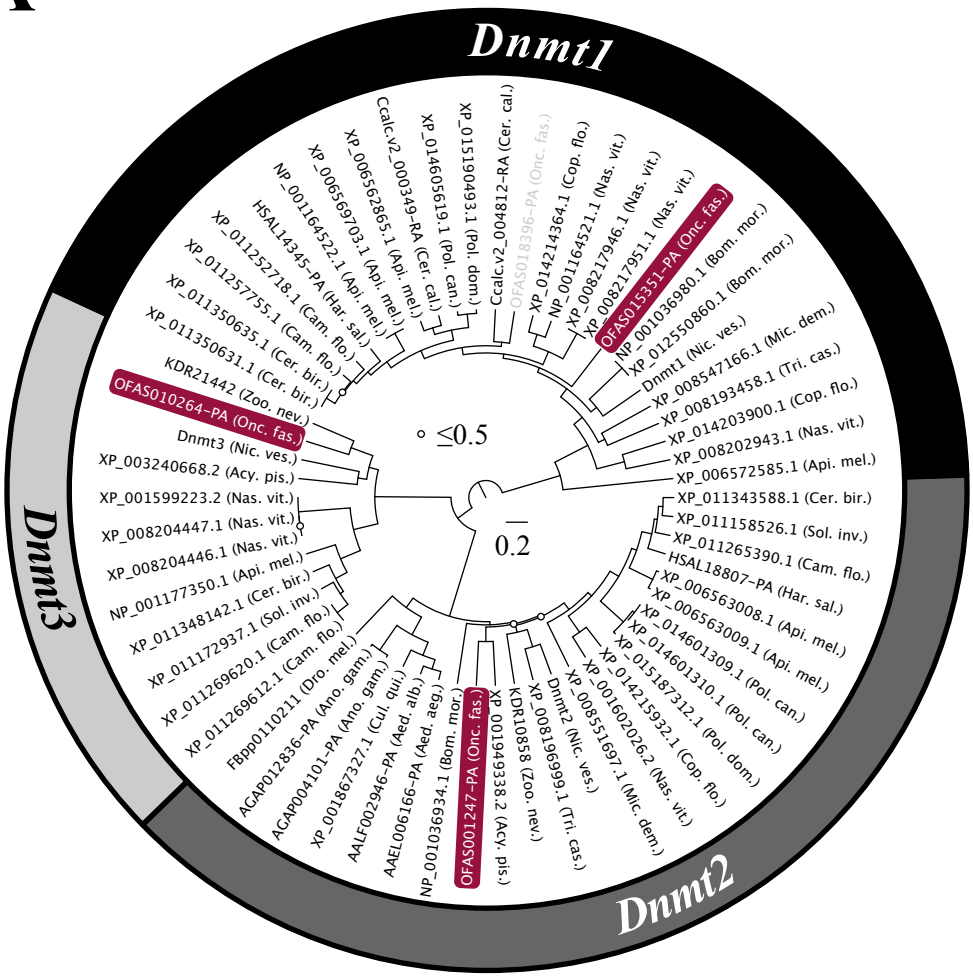
A


Dnmt1
Dnmt3
Dnmt2
≤0.5
0.2
OFAS018396-PA (Onc. fas.)
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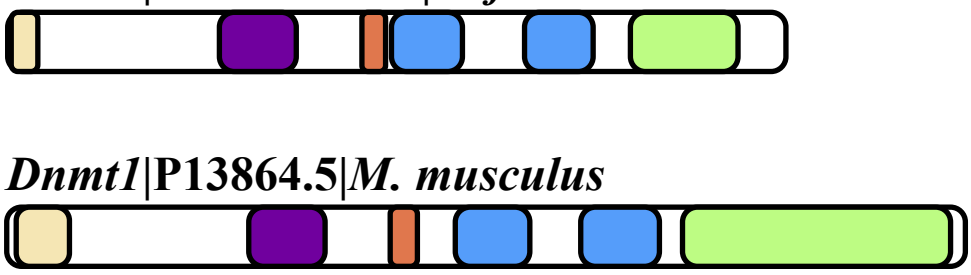
OFAS010264-PA (Onc. fas.)
OFAS001247-PA (Onc. fas.)
XP_015190693.1 (Del. dom.)
Ccalc.v2_004812-RA (Cer. cal.)
XP_014214364.1 (Cop. flo.)
NP_001164521.1 (Nas. vit.)
XP_008217946.1 (Nas. vit.)
XP_008217951.1 (Nas. vit.)
NP_001036980.1 (Bom. mor.)
XP_012550860.1 (Bom. mor.)
XP_011343588.1 (Cer. bir.)
XP_011158526.1 (Sol. inv.)
XP_011265390.1 (Cam. flo.)
HSAL18807-PA (Har. sal.)
XP_006563008.1 (Api. mel.)
XP_006563009.1 (Api. mel.)
XP_014601309.1 (Pol. can.)
XP_014601310.1 (Pol. can.)
XP_015187312.1 (Pol. dom.)
XP_014215932.1 (Cop. flo.)
XP_008551697.1 (Mic. dem.)
XP_011602026.2 (Nas. vit.)
Dnmt2 (Nic. ves.)
XP_008196999.1 (Tri. cas.)
KDR10858 (Zoo. nev.)
XP_001949338.2 (Acy. pis.)
NP_001036934.1 (Bom. mor.)
AAEL006166-PA (Aed. aeg.)
AALF002946-PA (Aed. alb.)
XP_001867327.1 (Cul. qui.)
AGAP004101-PA (Ano. gam.)
AGAP012836-PA (Ano. gam.)
FBpp0110211 (Dro. mel.)
XP_011269612.1 (Cam. flo.)
XP_011269620.1 (Cam. flo.)
XP_011172937.1 (Sol. inv.)
XP_011348142.1 (Cer. bir.)
NP_001177350.1 (Api. mel.)
XP_008204446.1 (Nas. vit.)
XP_008204447.1 (Nas. vit.)
XP_001599223.2 (Nas. vit.)
XP_003240668.2 (Acy. pis.)
Dnmt3 (Nic. ves.)
KDR21442 (Zoo. nev.)
XP_011350631.1 (Cer. bir.)
XP_011350635.1 (Cer. bir.)
XP_011257755.1 (Cam. flo.)
XP_011257718.1 (Cam. flo.)
HSAL14345-PA (Har. sal.)
NP_001164522.1 (Api. mel.)
XP_006569703.1 (Api. mel.)
XP_006562865.1 (Api. mel.)
Ccalc.v2_000349-RA (Cer. cal.)
XP_014605619.1 (Pol. can.)
XP_015190693.1 (Del. dom.)

B


Dnmt1|OFAS015351|*O. fasciatus*
Dnmt1|P13864.5|*M. musculus*
BAH domain
C-5 cytosine-specific DNA methylase
Cytosine specific DNA methyltransferase replication foci domain
DMAP1-binding domain
CXXC zinc finger domain
100 aa