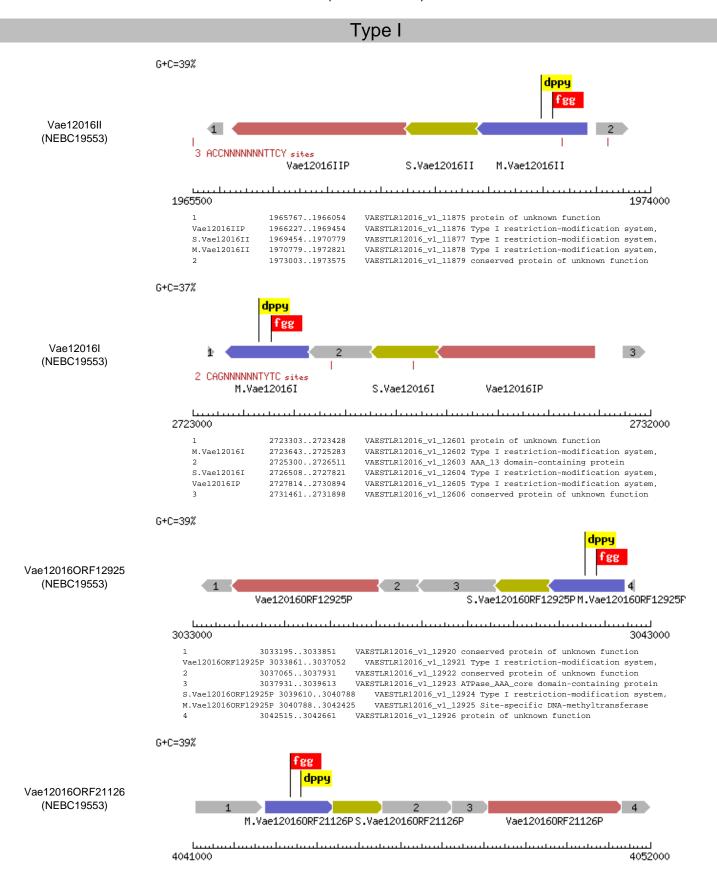
## Vibrio aestuarianus 12\_016

Created 03/12/23 (NEBC19553)



1 4041084..4042658 VAESTLR12016\_v1\_21125 TnsD domain-containing protein

M.Vael20160RF21126P 4042748..4044385 VAESTLR12016\_v1\_21126 Site-specific DNA-methyltransferase

S.Vael20160RF21126P 4044385..4045563 VAESTLR12016\_v1\_21127 Type I restriction-modification system,

2 4045560..4047242 VAESTLR12016\_v1\_21128 ATPase\_AAA\_core domain-containing protein

3 4047242..4048108 VAESTLR12016\_v1\_21129 conserved protein of unknown function

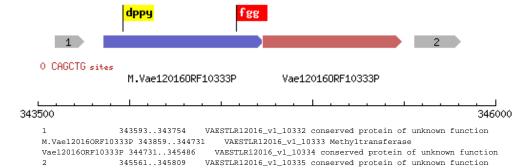
Vael20160RF21126P 4048121..4051312 VAESTLR12016\_v1\_21130 Type I restriction-modification system,

4 4051322..4051978 VAESTLR12016\_v1\_21131 conserved protein of unknown function

## Type II

### G+C=37%

## Vae12016ORF10333 (NEBC19553)



### G+C=45%

## Vae12016ORF11144 (NEBC19553)

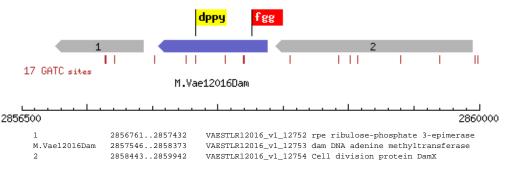


1215500 1220500

1 1215906..1216436 VAESTLR12016\_v1\_11143 DNA methyltransferase
M.Vael20160RF11144P 1216537..1217370 VAESTLR12016\_v1\_11144 Site-specific DNA-methyltransferase
2 1217367..1220312 VAESTLR12016\_v1\_11145 Putative phage protein

## G+C=44%

## Vae12016Dam (NEBC19553)



#### G+C=40%

## Vae12016ORF20737 (NEBC19553)



# 18819500 3821000

1 3819640..3819822 VAESTLR12016\_v1\_20736 hag DNA for RVC repeated sequence M.Vae12016ORF20737P 3819902..3820576 VAESTLR12016\_v1\_20737 Methyltransferase 2 3820713..3820883 VAESTLR12016\_v1\_20738 protein of unknown function

