logo

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T162 WP 010894925.1 NC 002488:2371566-2371868   MEKGTPHYKLPVVKALVKAGKVRATATAYQGARELGIKDLTGMCTVVLALTSADFYKSMTTYADHTVFQDVYHAKTASGD    80
T6100 NP 417494.1 NC 000913:c3168544-3168248    MEKRTPHTRLSQVKKLVNAGQVRTTRSALLNADELGL.DFDGMCNVIIGLSESDFYKSMTTYSDHTIWQDVYRPRLVTG.    78
T1045 NP 417494.1 NC 000913:c3168544-3168248    MEKRTPHTRLSQVKKLVNAGQVRTTRSALLNADELGL.DFDGMCNVIIGLSESDFYKSMTTYSDHTIWQDVYRPRLVTG.    78
T314 WP 010954960.1 NC 002947:c4750751-4750455  MEKRMPHCPLERVKALAAARRIRPTGAALKGAKALGM.DFPGMLEVITSLKRTDFYKSMTSHIDHRVWQDVYRPLTAIG.    78
consensus                                       !!!**!!**!**!! !**!***!*!**!** !**!!* !**!!**!***!***!!!!!!!**!!***!!!!***  *!
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logo

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T162 WP 010894925.1 NC 002488:2371566-2371868   EVYLKLTVIDDVLIVSFKEL   100
T6100 NP 417494.1 NC 000913:c3168544-3168248    QVYLKITVIHDVLIVSFKEK    98
T1045 NP 417494.1 NC 000913:c3168544-3168248    QVYLKITVIHDVLIVSFKEK    98
T314 WP 010954960.1 NC 002947:c4750751-4750455  YVYLKLSVVDDVLIVSFKEL    98
consensus                                       *!!!!!**!**!!!!!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved