```
                                                        ATGCAATACACTTACCAGCACATTCAGGATTTGGTACCGGTCCGACTCCACAAAAC    57
..........................................................................................................    0
ATGCAATACACTTACCAGCACATTCAGGATTTGGTACCGGGTCCGACTCCACAAAACTTCTACGGGAAAATAATTTTCATCAAGAAAAAAATCAACCAAATTGTTGTTCTTATCAAAGAT    120
ATGCAATACACTTACCAGCACATTCAGGATTTGGTACCGGGTCCGACTCCACAAAACTTCTACGGGAAAATAATTTTCATCAAGAAAAAAATCAACCAAATTGTTGTTCTTATCAAAGAT    120
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TTCTACGGGAAAATAATTTTCATCAAGAAAAAAATCAACCAAATTGTTGTTCTTATCAAACTACGCCAAGTGGTACGAGTTCATCGGTGCAAAATTCAGTCAATTCTCAATTCAAAAGAA    177
..........................................................................................................    0
GAAACGCAATCTATCTATTTACGAGTAATACCAAAAGAAGACCAGGAGCTTGAATTTCAGCTACGCCAAGTGGTACGAGTTCATCGGTGCAAAATTCAGTCAATTCTCAATTCAAAAGAA    240
GAAACGCAATCTATCTATTTACGAGTAATACCAAAAGAAGACCAGGAGCTTGAATTTCAGCTACGCCAAGTGGTACGAGTTCATCGGTGCAAAATTCAGTCAATTCTCAATTCAAAAGAA    240
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GGAATTGCTCAAATCGGATTATTTGGTTGTCATCTTATAGCTTGGTCTCAATCCGGAAAAGTGGATAATCCTGTTATTATCAGCTCAAGATCATGGACTAAAAGCGATGAAGACAGTGAA    297
..........................................................................................................    0
GGAATTGCTCAAATCGGATTATTTGGTTGTCATCTTATAGCTTGGTCTCAATCCGGAAAAGTGGATAATCCTGTTATTATCAGCTCAAGATCATGGACTAAAAGCGATGAAGACAGTGAA    360
GGAATTGCTCAAATCGGATTATTTGGTTGTCATCTTATAGCTTGGTCTCAATCCGGAAAAGTGGATAATCCTGTTATTATCAGCTCAAGATCATGGACTAAAAGCGATGAAGACAGTGAA    360
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AGGCTCCAGACGCTCCGCAAACTAGGAAAATCAAGAAGAAAGTCAGGTCGGAAAACATCAGTTGATACTATGGCAAACAAATTAATTGAGCGCCGAGAAGCGATGTTTGCTGATACATTT    417
..........................................................................................................    0
AGGCTCCAGACGCTCCGCAAACTAGGAAAATCAAGAAGAAAGTCAGGTCGGAAAACATCAGTTGATACTATGGCAAACAAATTAATTGAGCGCCGAGAAGCGATGTTTGCTGATACATTT    480
AGGCTCCAGACGCTCCGCAAACTAGGAAAATCAAGAAGAAAGTCAGGTCGGAAAACATCAGTTGATACTATGGCAAACAAATTAATTGAGCGCCGAGAAGCGATGTTTGCTGATACATTT    480
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ATCAAGAGCCTTTTCAACAAAATTGCACTTTCAAGAAAGGAACATCTTTCAAGAAATGCTCGAGAATTGTTCTATCACCGGCCTGGAGACATTGTTGAAACTCAAAATTTACTAGAAATT    537
..........................................................................................................    0
ATCAAGAGCCTTTTCAACAAAATTGCACTTTCAAGAAAGGAACATCTTTCAAGAAATGCTCGAGAATTGTTCTATCACCGGCCTGGAGACATTGTTGAAACTCAAAATTTACTAGAAATT    600
ATCAAGAGCCTTTTCAACAAAATTGCACTTTCAAGAAAGGAACATCTTTCAAGAAATGCTCGAGAATTGTTCTATCACCGGCCTGGAGACATTGTTGAAACTCAAAATTTACTAGAAATT    600
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GATGATTCTTGGTTCAACGATGAGAATTCGGAACAATTTGTGCAATATGTATTGAACTGTACAACTTGTCATGTCGAATACAATCACGTTGAGTATGCACAAAACAATATCCCAACAAAT    657
..........................................................................................................    0
GATGATTCTTGGTTCAACGATGAGAATTCGGAACAATTTGTGCAATATGTATTGAACTGTACAACTTGTCATGTCGAATACAATCACGTTGAGTATGCACAAAACAATATCCCAACAAAT    720
GATGATTCTTGGTTCAACGATGAGAATTCGGAACAATTTGTGCAATATGTATTGAACTGTACAACTTGTCATGTCGAATACAATCACGTTGAGTATGCACAAAACAATATCCCAACAAAT    720
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TGTCGGTTCTGTCAAGAAGCAATGGAAAGTTTCCACGCTGCATTTCGGATTCGTATTTCTATTGAAACTTATGGAGTATTTCTTACAATCCCTTTGGAATTGATTAAAACGGAGCTGGAC    777
.....................ATGGAAAGTTTCCACGCTGCATTTCGGATTCGTATTTCTATTGAAACTTATGGAGTATTTCTTACAATCCCTTTGGAATTGATTAAAACGGAGCTGGAC    99
TGTCGGTTCTGTCAAGAAGCAATGGAAAGTTTCCACGCTGCATTTCGGATTCGTATTTCTATTGAAACTTATGGAGTATTTCTTACAATCCCTTTGGAATTGATTAAAACGGAGCTGGAC    840
TGTCGGTTCTGTCAAGAAGCAATGGAAAGTTTCCACGCTGCATTTCGGATTCGTATTTCTATTGAAACTTATGGAGTATTTCTTACAATCCCTTTGGAATTGATTAAAACGGAGCTGGAC    840
*********************!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

ATTTGTGAAGATTGGGATTCTGAAAGCAATATCGTAGAAGAGGAAGAGAAAGTTACTAGATTCAAAAAAAATATTCAGGAAAAGGTTCGCGATGCGTCAATTGTGCATATAAAAGGTATT    897
ATTTGTGAAGATTGGGATTCTGAAAGCAATATCGTAGAAGAGGAAGAGAAAGTTACTAGATTCAAAAAAAATATTCAGGAAAAGGTTCGCGATGCGTCAATTGTGCATATAAAAGGTATT    219
ATTTGTGAAGATTGGGATTCTGAAAGCAATATCGTAGAAGAGGAAGAGAAAGTTACTAGATTCAAAAAAAATATTCAGGAAAAGGTTCGCGATGCGTCAATTGTGCATATAAAAGGTATT    960
ATTTGTGAAGATTGGGATTCTGAAAGCAATATCGTAGAAGAGGAAGAGAAAGTTACTAGATTCAAAAAAAATATTCAGGAAAAGGTTCGCGATGCGTCAATTGTGCATATAAAAGG.GTT    959
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TCTTCGCTTCTACTAATAATCATGTTAAATATAAATTCAATTTCACTTGTGAATAATATTACTGAGAATAAACGGATTTTATTGCAAAAGTCTAATGTTCCGTCTAGTTTACAATTAAAA    1017
TCTTCGCTTCTACTAATAATCATGTTAAATATAAATTCAATTTCACTTGTGAATAATATTACTGAGAATAAACGGATTTTATTGCAAAAGTCTAATGTTCCGTCTAGTTTACAATTAAAA     339
TCTTCGCTTCTACTAATAATCATGTTAAATATAAATTCAATTTCACTTGTGAATAATATTACTGAGAATAAACGGATTTTATTGCAAAAGTCTAATGTTCCGTCTAGTTTACAATTAAAA    1080
GGTAGTGTTCCGTTCAC..TTGAATTGAAGTCAATTGATATTCATCTCTCGGAAGTTGTTTTCTACAATATTAATATTTGA........................................    1038
**!*****!!!***!*!***!****!!*!!***!!*!***!!!**!!***!*!***!*!!***!*!!!!***!!!!*!*******************************************
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ATCCTAATCACTCCATTCGTCATCGTCGTCGTCAGATTCTTCGGAATCACTTGGATTTATTCCGGATCTTCTTGCATCGAGGAAGTTCTGAATCTGATCGACAATTGTTCCAGACGTCGT    1137
ATCCTAATCACTCCATTCGTCATCGTCGTCGTCAGATTCTTCGGAATCACTTGGATTTATTCCGGATCTTCTTGCATCGAGGAAGTTCTGAATCTGATCGACAATTGTTCCAGACGTCGT     459
ATCCTAATCACTCCATTCGTCATCGTCGTCGTCAGATTCTTCGGAATCACTTGGATTTATTCCGGATCTTCTTGCATCGAGGAAGTTCTGAATCTGATCGACAATTGTTCCAGACGTCGT    1200
.......................................................................................................................    1038
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TGA    1140
TGA     462
TGA    1203
...    1038
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☒ non conserved
☒ ≥ 50% conserved