```
# Program: water
# Rundate: Sat 8 Apr 2023 10:22:33
# Commandline: water
    -auto
#
    -stdout
    -asequence emboss_water-I20230408-102231-0207-96177647-p2m.asequence
#
    -bsequence emboss_water-I20230408-102231-0207-96177647-p2m.bsequence
    -datafile EBLOSUM62
#
#
    -gapopen 10.0
#
    -gapextend 0.5
    -aformat3 pair
#
#
    -sprotein1
#
    -sprotein2
# Align_format: pair
# Report file: stdout
#
# Aligned sequences: 2
# 1: YP 009724390.1
# 2: YP 009047204.1
# Matrix: EBLOSUM62
# Gap penalty: 10.0
# Extend_penalty: 0.5
#
# Length: 1440
# Identity:
             433/1440 (30.1%)
# Similarity:
              655/1440 (45.5%)
# Gaps:
              276/1440 (19.2%)
# Score: 1568.5
#
#
YP 009724390.
                3 VFLVLLPL-----PPAYTNSFTR
                                                                  34
                                    |.|.|:::... |....
YP 009047204.
                5 VFLLMFLLTPTESYVDVGPDSVKSACIEVDIQQTFFDKTWPRPIDVSKAD
                                                                  54
YP 009724390.
               35 GVYYPDKVFRSSVLHSTQDLFLPF---FSNVTWFHAIHVSGTNGTKRF--
                                                                  79
                  [:.[].....[::..:.[.]] [: ...:..:.[.].::[]...[.]
YP 009047204.
               55 GIIYPQGRTYSNITITYQGLF-PYQGDHGDMYVYSAGHATGTTPQKLFVA
                                                                 103
YP_009724390.
               80 -- DNPVLPF------GWIF
                                                                 106
                    . . . | . . |
                                    :.|....|...|...|
YP_009047204.
               104 NYSQDVKQFANGFVVRIGAAANSTGTVIISPSTSATIRKIYPAFMLGSSV
                                                                 153
YP_009724390.
               107 GTTLDSK-----TQSLLIVNNATNVVIKVCEFQFCNDPFLGVY-----
                                                                 144
                  YP 009047204.
               154 GNFSDGKMGRFFNHTLVLLPDGCGTLLRA--FYCILEPRSGNHCPAGNSY
                                                                 201
               145 ----YH-----KNNKSWMESEFRVYSSANNCTFEY---VSQPFLM
                                                                 177
YP 009724390.
                               -..|::...:.|:.|.:..|||||.| :::..::
YP 009047204.
               202 TSFATYHTPATDCSDGNYNRNASLNSFKEYFNLRNCTFMYTYNITEDEIL
                                                                 251
               178 DLEGKQGNFKNLREFVFKNIDGYFKIYSKHTPINLVRDLPQGFSALEPLV
YP 009724390.
                                                                 227
                  :..|....:.:.|..:.|..
               252 EWFGITQTAQGVHLFSSRYVDLY------
YP 009047204.
                                                                 274
              228 DLPIGINITRFQTL-----LALHRSYLTPGDSSSGWTAGAAAYYVG
                                                                 268
YP 009724390.
                                ..:..|..:.....
                      |.|:.:|.||
               275 ----GGNMFQFATLPVYDTIKYYSIIPHSIRSIQSDRKAW----AAFYVY
YP 009047204.
                                                                 316
YP 009724390.
              269 YLQPRTFLLKYNENGTITDAVDCALDPLSETKCTLKSFTVEKGIYQTSNF
                                                                 318
                  YP 009047204.
               317 KLQPLTFLLDFSVDGYIRRAIDCGFNDLSQLHCSYESFDVESGVYSVSSF
                                                                 366
```

08/04/2023, 14:52 YP 009724390.	https://www.ebi.ac.uk/Tools/services/rest/emboss_water/result/emboss_water-I2023	30408-102231-0207-961 368
YP_009047204.	: :. :	414
YP_009724390.	369 YNSASFSTFKCYGVSPTKLNDLCFTNVYADSFVIRGDEVRQIAPGQTGKI	418
YP_009047204.	.: .:. . : : :::: .	464
YP_009724390.	419 ADYNYKLPDDFTGC-VIAWNSNNLDSKVGG-NYNYLYRLFRKSNLKPFER	466
YP_009047204.	::: :: : .: : ::: : 465 SQFNYKQSFSNPTCLILATVPHNLTTITKPLKYSYINKCSRLLS	508
YP_009724390.	467 DISTEIYQAGSTPCNGVEGFNCYFPLQSYGFQPTNG	502
YP_009047204.	:. : : : :. . : :: 509 DDRTEVPQLVNANQYSPCVSIVPSTVWEDGDYYRKQLSPLEGGGWLVASG	558
YP_009724390.	503 VGYQPYRVVVLSFELLHAPATVCGPKKSTNLVKNKCVNFNF	543
YP_009047204.	:.:. .: : 559 STVAMTEQLQMGFGITVQYGTDTNSVCPKLEFANDTKIASQLGNCVEYSL	608
YP_009724390.	544 NGLTGTGVLTESNKKFLPFQQFGRDIADTTDAVRDPQTLEILDITPCSFG	593
YP_009047204.	609 YGVSGRGVFQNCTAVGVRQQRFVYDAYQ	636
YP_009724390.	594 GVSVITPGTNTSNQVAVLYQDVNCTEVPVAIHA	626
YP_009047204.	. :: 637NLVGYYSDDGNYYCLRACVSVPVSVIYDKETKTHATLF	674
YP_009724390.	627DQLTPTWRVYSTGSNVFQTRAGCLIGAEHVNNSY	660
YP_009047204.	675 GSVACEHISSTMSQYSRSTRSMLKRRDSTYGPLQTPVGCVLGL VNSSL	722
YP_009724390.	661ECDIPIGAGICASYQT-QTNSPRRARSVASQSIIAYTMSLGAENSVA	706
YP_009047204.	: .: : : .: : : 723 FVEDCKLPLGQSLCALPDTPSTLTPRSVRSVPGEMRLASIA	763
YP_009724390.	707 YSNNSIAIPTNFTISVTTEILPVSMTKTSVDCTMYICGD :: : :: : . .:.::. .: :	745
YP_009047204.	764 FNHPIQVDQLNSSYFKLSIPTNFSFGVTQEYIQTTIQKVTVDCKQYVCNG	813
YP_009724390.	746 STECSNLLLQYGSFCTQLNRALTGIAVEQDKNTQEVFAQVKQIYKTPPIK	795
YP_009047204.	: .: . ::: : . :. .:.::: . .	863
YP_009724390.	796 DFGG-FNFSQILPDPSKPSKRSFIEDLLFNKVTLADAGFIKQYGDCL	841
YP_009047204.	864 GFGGDFNLTLLEPVSISTGSRSARSAIEDLLFDKVTIADPGYMQGYDDCM	913
YP_009724390.	842GDIAARDLICAQKFNGLTVLPPLLTDEMIAQYTSALLAGTITSGWTFG	889
YP_009047204.	914 QQĠPASARDĿĬĊAQYVAĠYKVĿPPĿMDVNMEAAYTSSĿĿGSIAGVĠWTAĠ	963
YP_009724390.	890 AGAALQIPFAMQMAYRFNGIGVTQNVLYENQKLIANQFNSAIGKIQDSLS: .: . : . . .	939
YP_009047204.	964 LSSFAAİPFAQSIFYRLNĞVĞITQQVLSENQKLİANKFNQALĞAMQTGFT	1013
YP_009724390.	940 STASALGKLQDVVNQNAQALNTLVKQLSSNFGAISSVLNDILSRLDKVEA : : . :. :. .	989
YP_009047204.	1014 TŤNEÁFQKVQĎAVŇNŇÁQÁĽSKĽASEĽŠNTFĠÁÍŠASIGĎÍIQŘĽĎVLÉQ	1063
YP_009724390.	990 EVQIDRLITGRLQSLQTYVTQQLIRAAEIRASANLAATKMSECVLGQSKR :. . :	1039
YP_009047204.	1064 DAQİDRLİNGRLTTLNAFVAQQLVRSESAALSAQLAKDKVNECVKAQSKR	1113
YP_009724390.	1040 VDFCGKGYHLMSFPQSAPHGVVFLHVTYVPAQEKNFTTAPAICHDGK : : : : : : : : . : : : : : : : :	1086
YP_009047204.	1114 SGFCGQGTHIVSFVVNAPNGLYFMHVGYYPSNHIEVVSAYGLCDAANPTN	1163
YP_009724390.	1087 AHFPREGVFV-SNGTHWFVTQRNFYEPQIITTDNT-FVSGNCDVVI	1130

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YP_009047204.	. : : . . : . :. : : 1164 CIAPVNGYFIKTNNTRIVDEWSYTGSSFYAPEPITSLNTKYVAPQ	1208
YP_009724390.	1131 GIVNNTVYDPLQPELDSFKEELDKYFKNHTSPDVDLGDISGINA	1174
YP_009047204.	:: . . :: :: .:::. . 1209 -VTYQNISTNLPPPLLGNSTGIDFQDELDEFFKNVSTSIPNFGSLTQINT	1257
YP_009724390.	1175 SVVNIQKEIDRLNEVAKNLNESLIDLQELGKYEQYIKWPWYIWLGFIAGL	1224
YP_009047204.	::::: : :	1307
YP_009724390.	1225 IAIVMVTIMLCCMTSCCSCLKGCCSCGSCC-KFDEDDSEP 1263	
YP_009047204.	: :.::. . . ::: . . 1308 VALALCVFFILCCTGCGTNCMGKLKCNRCCDRYEEYDLEP 1347	
#		
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