```
 Score = 1448 (373.6 bits), Expect = 3.5e-103, Sum P(3) = 3.5e-103
 Identities = 242/280 (86%), Positives = 242/280 (86%), Strand = Minus / Plus

elegans:   3392 CATTTCATAGTGTCACACGCGCACCCATGAGTTGTCGGCACACCACTCCCCACTACCCCT 3333
                ||||  ||||||| |  |||||  |||     ||||||| |||      | | |||||||
briggsae: 10570 CATTCCATAGTGGCCGACGCGAGCCCGGTTTTTGTCGGTACATGCGCGCACCCTACCCCC 10629

                                    ......>>>>.>>>>....<<<<<<<<.............
elegans:   3332 ACCCTCTCCTCCATTCAGTATCGCTTCTTCGGCTTATTAGCTAAGATCAAAGTGTAGTAT 3273
                |  ||| | |   | | ||||||||||||||||||||||||||||||||||||||||||
briggsae: 10630 CGCGCCTCGTTCTCACCGCATCGCTTCTTCGGCTTATTAGCTAAGATCAAAGTGTAGTAT 10689
                                    ----------------------------------------

                .......AAAAAA..>>>>>>aaaaaa>>>>>>>....<<<<<<<....<<<<<<.....
elegans:   3272 CTGTTCTTATCGTATTAACCTACGGTATACACTCGAATGAGTGTAATAAAGGTTATATGA 3213
                |||||||||||||||||||||||||| ||||||||||||||||||||||||||||||||
briggsae: 10690 CTGTTCTTATCGTATTAACCTACGGTATGCACTCGAATGAGTGTAATAAAGGTTATATGA 10749
                ----------------------------*--------------------------------

                ............>>>>>>..>>>>>>....<<<<<<<<<<<..>>>>>>.>>>>>>..
elegans:   3212 TTTTTGGAACCTAGGGAAGACTCGGGGCTTGCTCCGACTTCCCAAGGGTCGTCCTGGCGT 3153
                |||||||||||||| ||||||||||||||||||||||| |||||||||||| |||||
briggsae: 10750 TTTTTGGAACCTAGGAAAGACTCGGGGCTTGCTCCGACTTTCCAAGGGTCGTCCCGGCGT 10809
                ---------------*------------------------*-------------*-----

                .........<<<<<<..<<<<<<...
elegans:   3152 TGCACTGCTGCCGGGCTCGGCCCAGTCCCCGAGGGGACAA 3113
                |||||||||||||||||||||||||| | ||||||||
briggsae: 10810 TGCACTGCTGCCGGGCTCGGCCCAGTCCCTGTGGGGACAA 10849
                -------------------------
```

Figure 3