Figure S1. cDNA Sequences of Piriform Silk Proteins.

A. *A. trifasciata* Piriform:

```
      A  P  L  P  A  P  A  P  R  P  R  P  A  P  A  P  R  P  A  P
3     gctccattgcctgctccggctccgagacccagaccagcacccgctccgcgaccagcgcca  62
      V  Y  A  P  A  P  V  V  S  Q  I  Q  A  F  T  S  N  R  G  S
63    gtttatgctccagctccagttgtttcgcaaattcaggcatttacatcgaatcgaggttcc  122
      T  Q  Q  N  S  F  A  Q  Q  S  S  V  A  Q  Q  S  S  V  A  Q
123   actcaacagaattcctttgcacagcaatcatctgttgcacagcagtcatctgttgcacag  182
      Q  S  S  V  A  Q  Q  S  S  V  A  Q  Q  S  S  T  A  Q  Q  S
183   caatcatctgttgcacagcaatcatctgttgcacagcaatcatctactgcacagcaatca  242
      S  V  A  Q  Q  S  A  V  A  Q  S  Q  Q  S  S  Y  A  A  A  S
243   tccgttgctcaacagtcagcagttgcccaatcgcaacaaagttcctacgccgctgcgtct  302
      S  G  G  S  S  A  S  Q  S  Q  A  T  V  S  S  A  P  V  Y  F
303   tctggtggttcgagtgcgtcgcagtcacaagctactgtctcaagcgccccagtatatttc  362
      N  S  Q  I  L  T  N  N  L  A  S  S  L  Q  S  L  S  G  L  N
363   aactcgcaaattttgacaaataacctggcttcatctctgcaatcactgagtggccttaat  422
      Y  V  S  S  G  Q  L  S  S  S  D  V  A  S  I  V  A  E  A  V
423   tacgtatcaagtggtcagttgagttcctcggacgtcgcttccattgtcgcggaagctgta  482
      S  Q  S  L  G  V  S  Q  G  S  V  Q  N  I  I  S  Q  R  L  N
483   tcccagtccctcggagtgtcacaagggtcagttcaaaatattataagccaaagattgaac  542
      G  I  G  P  G  A  S  P  S  S  V  S  A  A  I  A  N  A  V  S
543   ggcataggccctggagcttccccatcatcagtctcagcagcgatagcaaatgcagtatct  602
      S  A  V  Q  G  S  A  S  A  A  P  G  Q  E  Q  S  I  A  Q  S
603   tcggcagttcaaggatcagcctctgcagctcccggacaggaacaatcgattgcgcaaagt  662
      M  S  S  A  I  S  S  A  F  Q  Q  I  I  S  Q  R  T  A  V  A
663   atgtcttcagctatttcctcagctttccaacaaattatttcgcagagaacagcagtggct  722
      P  A  P  S  T  L  P  S  P  L  P  A  P  R  P  R  P  A  P  A
723   cccgctcctagcaccctaccatctcctcttcctgctccgagaccgagaccagcgccagct  782
      L  Q  P  G  P  V  Y  A  P  A  L  A  Y  A  P  A  P  V  Y  A
783   ctacaaccagggccagtttatgcaccagcacttgcgtatgcaccagcgccagtttatgcc  842
      P  A  P  V  V  S  Q  F  H  S  S  A  S  T  E  A  T  A  E  Q
843   ccagctccagttgtttcgcagtttcattcatctgcttctactgaagccacagctgaacag  902
      N  S  F  V  Q  T  S  L  A  Q  S  Q  P  G  S  Y  S  S  Q  S
903   aattcctttgtacagacgtcacttgcccaatcgcaaccaggctcctattcgtcacaatca  962
      N  A  L  S  S  N  S  G  Q  Q  A  S  V  A  R  S  Q  S  S  G
963   aatgctctgtcttcaaattcaggtcaacaggcaagtgtagctcgttctcagtctagcgga 1022
      S  S  F  S  S  G  S  S  S  D  F  G  S  S  S  S  S  G  A  F
1023  tcttccttttctagcggatcgtcttcagatttcgggtcttcttcgtcttctggtgcgttt 1082
      S  T  A  S  S  P  V  S  Y  A  P  A  T  I  S  S  S  S  S  V
1083  tcaactgccagcagccccgtgtcctatgctccagcaacaatcagcagctcttcgtctgtt 1142
      A  A  S  S  F  S  S  G  S  S  S  G  F  S  S  S  A  S  G  S
1143  gcggcctcatcgttttcgagtggatcttcctcaggttttagctcttctgcttctggatca 1202
      A  L  A  A  S  A  A  Q  Q  L  L  S  P  A  A  N  Q  R  I  A
1203  gcattggccgcttccgctgcccagcagcttttgtcacctgcagcgaatcagaggattgca 1262
      A  L  S  N  T  L  K  S  A  V  S  G  G  R  V  N  Y  G  A  L
1263  gcactgtccaatactttgaagtctgctgtgtcaggaggacgcgtcaattacggggctctt 1322
      S  N  S  L  A  S  A  A  S  Q  I  Q  S  S  S  R  L  S  K  N
1323  tcgaattctcttgcgagcgctgcaagtcaaatacaaagttcctccagattatcaaaaaac 1382
      E  V  L  V  E  V  L  L  E  T  L  A  A  L  L  D  S  L  S  I
1383  gaagtcctagtcgaggttctcttggaaactttggccgctttgttggattccctttcaatt 1442
      S  G  S  S  S  S  Q  F  A  Q  A  V  L  Q  A  F  A  *
1443  tctggatcatccagctctcaattcgcacaagcagtgctacaagcatttgcctagatttcg 1502
1503  ctcctttactctggaaaaccacagcaaaaataatatgaaattttaattgtatctttattt 1562
1563  gttatatattgtatctgattttgggtaatggtaataaaaatttaatgaaagcataaaaaa 1622
1623  aaaaaaa  1629
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B. *N. cruentata* Piriform:

```
      A  P  R  P  L  P  A  P  A  P  R  P  L  P  A  P  A  P  R  P
2     gctcctcgtcctctaccagctcctgctcctcgtcctctaccagctcctgctcctcgtccc   61
      L  P  A  P  L  P  A  P  L  P  R  P  R  P  A  P  I  V  S  Q
62    ctaccagcacctttgccagctcctcttccaagaccacgcccagcacccatagtttctcaa  121
      V  Q  Q  A  S  A  L  Q  A  Q  S  Q  Q  S  A  F  A  Q  S  Q
122   gtgcaacaggcatccgctctacaggcacagtcacaacagtctgcttttgctcaatcccaa  181
      Q  S  S  I  A  Q  S  Q  Q  A  S  V  A  Q  S  Q  R  A  S  V
182   caatcgtccattgcacaatctcagcaagcctctgtcgcccaatcccaacgagcctccgtc  241
      S  Q  S  Q  Q  S  S  N  A  F  S  S  A  A  S  F  G  A  S  S
242   tctcaatcccagcaatctagtaatgcgttttcttctgcagcatcatttggagcttctagc  301
      V  A  S  S  A  S  T  Y  F  N  S  G  I  V  Q  S  S  I  A  S
302   gtagcatccagtgcttcgacttactttaattcgggaatagtacaaagtagcatcgcgtcg  361
      S  L  Q  S  S  S  A  L  S  S  I  A  Y  G  Q  T  T  A  S  I
362   tcgttgcagtcttccagtgctctcagttccattgcctacggccagacaaccgcttctatc  421
      N  D  I  A  S  A  V  A  G  S  I  A  N  S  I  G  L  S  Q  Q
422   aacgatatagcatcggcagtcgctggcagcattgcaaattcaatcggactctcgcaacaa  481
      T  V  Q  S  I  I  S  Q  Q  L  A  S  A  G  S  G  A  S  A  Q
482   accgttcaaagtattattagtcaacaactagccagtgcaggatccggagcatctgctcaa  541
      T  L  A  S  L  I  S  S  A  V  S  S  L  V  Q  Q  S  G  S  V
542   acattggcttcattgatatccagcgcagtttcctccttggttcaacaatctggatcggta  601
      S  A  G  Q  E  Q  S  I  S  Q  A  L  S  S  S  I  S  S  S  L
602   tcagccggacaagaacagagtatttcgcaagcactttccagttctatctcgagttctttg  661
      N  Q  L  V  A  A  R  P  L  P  A  P  A  P  R  P  L  P  A  P
662   aatcaattggtagctgcaagacctctaccagctcctgctcctcgtcccctaccagcacct  721
      L  P  A  P  L  P  R  P  R  P  A  P  I  V  S  Q  V  Q  Q  A
722   ttgcctgctcctcttccaagaccacgcccagcacccatagtttctcaagtgcagcaggca  781
      S  A  L  Q  A  Q  S  Q  Q  S  A  F  A  Q  S  Q  Q  S  S  I
782   tccgctctacaggcacagtcacaacagtctgcttttgctcaatcccaacagtcgtccatt  841
      A  Q  S  Q  Q  A  S  V  A  Q  S  Q  Q  S  S  I  S  Q  S  Q
842   gcacagtctcagcaagcctctgtcgcccaatcccaacaatcctccatctcccaatcccaa  901
      Q  A  S  V  S  Q  S  Q  Q  S  S  N  A  F  S  S  A  A  S  S
902   caagcctccgtctctcaatcccagcaatctagtaatgcgttttcttctgcagcatcttct  961
      G  A  S  S  V  A  S  S  A  S  T  Y  F  N  S  G  I  V  Q  S
962   ggagcttctagcgtagcatccagtgcttcgacttacttcaattcgggcatagtacaaagc 1021
      S  I  A  S  S  L  Q  S  S  S  A  L  S  S  I  A  Y  G  Q  T
1022  agcatcgcgtcgtcgttgcagtcttccagtgctctcagttccattgcctacggccagaca 1081
      T  A  S  I  S  D  I  A  S  A  V  A  G  S  I  A  N  S  I  G
1082  accgcttctatcagcgatatagcatcggcagtcgctggcagcattgcaaattcaatcgga 1141
      L  S  Q  Q  T  V  Q  S  V  I  S  Q  Q  L  A  S  A  G  S  G
1142  ctctcgcaacaaaccgttcaaagtgttattagtcaacaactagccagtgcaggatccgga 1201
      A  S  A  Q  T  L  A  S  L  I  S  S  A  V  S  S  L  V  Q  Q
1202  gcatctgctcaaacattggcttcattgatatccagcgcagtttcctccttggttcaacaa 1261
      S  G  S  V  S  A  G  Q  E  Q  S  I  S  Q  A  L  S  S  S  I
1262  tctggatcggtatcagccggacaagaacagagtatttcgcaagcactttccagttctatc 1321
      S  S  S  L  N  Q  L  V  A  A  R  P  L  P  A  P  A  P  R  P
1322  tcgagttctttgaatcagttggtagccgcaagacctctaccagcccctgctcctcgtccc 1381
      L  P  A  P  L  P  A  P  L  S  R  P  R  P  V  P  V  Q  R  P
1382  ctaccagcacctttaccagctcctctttcaagacctcgaccagttccagtccaaagacct 1441
      Q  P  V  F  S  P  S  P  A  P  A  Y  A  P  A  P  F  T  Q  Q
1442  caacccgtattttcacccagtcccgctcctgcctatgctcctgccccattcactcagcag 1501
      S  T  F  A  Q  S  Q  Q  A  S  L  A  Q  S  Q  Q  Q  A  S  I
1502  tcgacttttgctcagtctcaacaagcttctcttgctcaatctcaacaacaagcatctatc 1561
      A  R  S  Q  Q  A  S  L  A  Q  S  Q  Q  S  A  F  A  Q  S  Q
1562  gctcgatcccaacaagcgtctctagcgcaatctcaacaatcggcttttgctcaatcccaa 1621
      Q  V  A  T  A  Q  S  Q  Q  S  S  G  G  F  S  T  S  S  T  G
1622  caagtagctacagcacaatctcagcaatcttctggtggattctccacatcatctactgga 1681
```

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      A  S  Q  I  S  S  S  A  I  S  T  S  S  G  S  A  L  A  N  S
1682  gcttctcaaatcagttcttcagccataagtacttcttcgggatctgcattggctaattcc 1741
      A  Q  Q  L  T  S  P  A  A  S  Q  R  I  S  Q  L  S  N  S  L
1742  gcacaacaactcacatcacctgcagcctctcaaagaatatctcagttatccaattcccta 1801
      A  S  A  V  S  G  G  Q  V  N  Y  A  A  L  S  N  S  I  A  S
1802  gcatctgcagtttctggtggacaggtcaactatgcagccttatctaattctattgctagt 1861
      A  A  S  Q  I  G  G  G  S  G  L  S  K  T  E  V  L  I  E  T
1862  gctgcaagtcaaattggaggtggatctggattatccaaaacggaagttctaattgaaact 1921
      L  L  E  T  L  A  A  L  L  E  S  L  S  L  P  G  S  A  S  G
1922  ctcttagaaaccctggctgctttattggaatctctttctcttcctggatcagctagtggt 1981
      G  S  Q  F  A  Q  A  M  L  A  A  L  A  *
1982  ggaagtcaattcgctcaagcaatgcttgcagctcttgcataaaatgtggttaataacaaa 2041
2042  taattttgttggaatgcttatgaatatttttaggggaatatatgatatacactttaaatg 2101
2102  aataatattgtataacttttgttaatgggaaataaaatttattagcaagcatcaaaaaa 2161
2162  aaaaaaaaaaa  2172
```

C. *N. clavipes* piriform:

```
      A  P  R  P  L  P  A  P  L  P  A  P  R  P  I  P  A  P  L  P
1     gctcctcgtccactcccagctcctttgcccgctcctcgtccaatcccagctcctttaccc   60
      A  P  R  P  I  P  A  P  L  P  R  P  V  P  I  V  S  Q  V  Q
61    gctcctcgtccaatcccagctcctcttccccgcccagtacccatagtttctcaagtacag  120
      Q  A  S  I  Q  Q  A  Q  S  S  S  A  Q  S  R  Q  S  S  V  A
121   caggcatccattcaacaggcacagtcgtcttctgctcagtcccgacaatcgtccgttgct  180
      Q  Q  A  S  I  S  Q  S  Q  Q  A  S  V  S  Q  S  Q  Q  A  S
181   cagcaagcctccatctctcaatcccagcaggcttcggtatcccaatcccagcaggcttcg  240
      V  S  Q  S  Q  Q  A  S  V  S  Q  S  Q  Q  S  S  N  A  Y  S
241   gtatcccaatcccagcaggcttcagtatcccaatctcagcagtcctctaatgcatattct  300
      A  A  S  N  A  A  S  S  V  S  Q  A  S  S  D  S  S  Y  F  N
301   gctgcgtctaatgccgcatccagcgtatctcaagcatcaagcgattcaagttactttaac  360
      S  Q  V  V  Q  S  A  L  S  S  S  L  Q  S  S  S  A  L  S  S
361   tcgcaagtagtacagagcgcactctcatcgtccttgcaatcttctagtgctctcagttca  420
      I  A  Y  G  Q  T  S  A  N  I  N  D  V  A  A  A  V  A  R  S
421   attgcttacggtcagacctcagccaatataaatgatgtagcagctgcggtcgctagaagc  480
      V  S  Q  S  L  G  V  S  Q  Q  A  A  Q  S  V  I  S  Q  Q  L
481   gtttcacaatcccttggagtttcccaacaagccgcacaaagtgtcatcagccaacaacta  540
      A  S  A  G  S  G  A  S  A  Q  T  L  A  Q  L  I  S  S  A  V
541   gcaagtgcaggatcgggagcatcggctcaaacattggcgcaattgatatccagcgcagtg  600
      S  S  L  V  Q  Q  S  G  T  V  S  A  G  Q  E  Q  S  I  S  Q
601   tcttccttggttcagcaatctggaacggtgtctgccggacaagaacagagtatttcacaa  660
      S  L  S  S  S  I  L  S  S  L  S  Q  V  V  A  Q  R  P  L  P
661   tccctttctagttcaatcttgtcttctctcagtcaagtagttgcccagaggccactacca  720
      V  P  A  P  R  P  L  P  A  P  L  P  A  P  R  P  I  P  A  P
721   gtacctgctcctcgtccactcccagcacctttgcccgctcctcgtccaatcccagctcct  780
      L  P  R  P  V  P  I  V  S  Q  V  Q  Q  A  S  I  Q  Q  A  Q
781   cttccccgcccagtacccatagtttctcaagtacaacaagcatccattcaacaggcacag  840
      S  S  S  A  Q  S  R  Q  S  S  V  A  Q  Q  A  S  V  S  Q  S
841   tcgtcttctgctcagtcccgacaatcgtccgttgctcagcaagcctccgtctctcaatcc  900
      Q  Q  A  S  V  S  Q  S  Q  Q  A  S  V  S  Q  S  Q  Q  S  S
901   cagcaggcttctgtttcccaatcccagcaggcttcggtctcccaatctcagcagtcctct  960
      N  A  Y  S  A  A  S  N  A  A  S  S  V  S  Q  V  S  S  A  S
961   aatgcatattctgctgcgtctaatgccgcatccagcgtatctcaagtatcaagcgcttca 1020
      S  Y  F  N  S  Q  V  V  Q  S  A  L  S  S  S  L  Q  S  S  S
1021  agttactttaactcgcaagtagtacagagcgcactctcatcgtccttgcaatcttctagt 1080
      A  L  S  S  I  A  Y  G  Q  T  S  A  N  I  N  D  V  A  A  A
1081  gctctcagttcaattgcttacggtcagacctcagccaatataaacgatgtagcagcagcg 1140
      V  A  R  S  V  S  Q  S  L  G  V  S  Q  Q  A  A  Q  S  V  I
1141  gtcgctagaagcgtttcacaatcccttggagtttcccaacaagccgcacaaagtgtcatc 1200
      S  Q  Q  L  S  S  A  G  S  G  A  S  A  Q  T  L  A  Q  L  I
1201  agccaacaactatcaagtgcaggatcgggagcatcggctcaaacattggcgcaattgata 1260
      S  S  A  V  S  S  L  V  Q  Q  S  G  T  V  S  A  G  Q  E  Q
1261  tccagcgcagtgtcttccttggttcagcaatctggaacggtgtctgccggacaagaacag 1320
      S  I  S  Q  A  L  S  S  S  I  L  S  S  L  S  Q  V  V  S  Q
1321  agtatttcacaagctctttccagttcaatcttgtcttctctcagtcaagtagtttcccag 1380
      R  P  L  P  A  P  R  P  A  P  A  P  R  P  L  P  E  P  L  P
1381  aggcctctaccagcaccacgaccagcacctgctcctcgtccactcccagaacctctgccc 1440
      A  P  R  P  I  P  A  P  L  P  R  P  V  P  I  V  S  Q  V  Q
1441  gctcctcgtccaatcccagctcctcttccccgcccagtacccatagtttctcaagtacag 1500
      Q  A  S  I  Q  Q  A  Q  S  S  S  A  Q  S  R  Q  S  A  V  A
1501  caggcatccattcaacaggcacagtcgtcttctgctcagtcccgacaatcggccgttgct 1560
      Q  Q  A  S  V  S  Q  S  Q  Q  A  S  V  S  Q  S  Q  Q  A  S
1561  cagcaagcctccgtctctcaatcccagcaggcttcggtctcccaatcccagcaggcttcg 1620
      V  S  Q  S  Q  Q  A  S  L  S  Q  T  Q  Q  A  S  V  S  Q  S
1621  gtatcccaatcccagcaggcttcgctatcccaaacccagcaggcttcggtatcccaatct 1680
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      Q  Q  S  S  N  A  Y  S  A  A  S  N  A  A  S  S  V  S  Q  A
1681  cagcagtcttctaatgcatattctgctgcgtctaatgccgcatccagcgtatctcaagca 1740
      S  S  A  S  S  Y  F  N  S  Q  V  V  Q  S  T  L  S  S  S  L
1741  tcaagcgcttcaagttactttaactcgcaagtagtacagagtacactctcatcatccttg 1800
      Q  S  S  S  A  L  S  S  I  A  Y  G  Q  T  S  A  N  I  N  D
1801  caatcttctagtgctctcagttcaattgcttacggtcagacctcagccaatataaacgat 1860
      V  A  A  A  V  A  R  S  V  S  Q  S  L  G  V  S  Q  Q  A  A
1861  gtagcagcagcggtcgctagaagcgtttcacaatcccttggagtttcccaacaagccgca 1920
      Q  S  V  I  S  Q  Q  L  A  S  A  G  A  G  A  S  A  Q  T  L
1921  caaagtgtcatcagccaacaactagcaagtgcaggagcaggagcatcggctcaaacattg 1980
      A  Q  L  I  S  S  A  V  S  S  L  V  Q  Q  S  G  T  V  S  A
1981  gcacaattgatatccagcgcagtgtcttccttggttcaacaatctggaacggtatctgcc 2040
      G  Q  E  Q  S  I  S  Q  A  L  S  S  S  I  L  S  S  L  S  Q
2041  ggacaagaacagagtatttcacaagcactttctagttcaatcttgtcttctctcagtcaa 2100
      V  V  A  Q  R  P  L  P  V  P  A  P  R  P  L  P  A  P  L  P
2101  gtagttgcccagaggccactaccagtacctgctcctcgtccactcccagcacctttgccc 2160
      A  P  R  P  I  P  .  .  .  .  .  .  .  .  .  .  .  .  .  .
2161  gctcctcgtccaatcccag .  .  .  .  .  .  .  .  .  .  .  .  .  .  2179
      V  I  L  S  S  L  S  Q  V  V  A  Q  R  P  L  P  A  P  R  P
3001  gtcatcttgtcttctcttagtcaagtagttgcccagagacctctaccagctcctcgtcca 3060
      L  P  V  P  L  P  R  P  Q  P  A  P  R  P  L  P  A  P  L  P
3061  ttgccagtacccctccaagaccccagcctgctcctcgtccattaccagcacctcttcca 3120
      R  P  Q  P  A  P  I  Q  R  P  Q  P  V  F  T  P  S  P  S  P
3121  agaccccaacctgctcctatccaaagacctcaacccgtatttactccaagtccttctccg 3180
      V  F  A  P  S  A  F  T  Q  Q  S  T  I  A  Q  S  Q  Q  V  S
3181  gtatttgctccttcagcattcactcaacaatcgactattgcgcaatcccaacaagtctct 3240
      Q  A  Q  S  S  L  A  Q  S  Q  Q  S  S  L  A  Q  S  Q  Q  S
3241  caagcacaatcctctttagcgcagtctcaacaatcctctttagcgcagtctcaacaatcc 3300
      S  L  A  Q  S  Q  Q  S  S  L  A  Q  S  Q  Q  S  S  L  A  Q
3301  tccttagcgcagtctcaacaatcctctttagcgcagtctcaacaatcctctttagcgcag 3360
      S  Q  Q  S  S  L  A  Q  S  Q  Q  S  S  L  A  Q  S  Q  Q  S
3361  tctcaacaatcctctttagcgcagtctcaacaatcctctttagcgcagtctcaacaatcc 3420
      S  L  A  Q  S  Q  Q  S  S  L  A  Q  S  Q  Q  S  S  F  A  Q
3421  tctttagcgcagtctcaacaatcctctttagcgcagtctcaacaatcttcttttgcacaa 3480
      S  Q  Q  V  A  A  A  Q  T  Q  Q  S  S  G  G  Y  S  A  T  S
3481  tcccaacaagtagcagcagcacaaacacagcagtcttcaggtggatactccgcaacttcc 3540
      M  G  A  S  Q  I  S  S  S  A  I  S  S  S  S  G  S  A  L  A
3541  atgggtgcgtcgcaaatctcttcttcggccataagttcttcttcaggatctgcattggct 3600
      S  S  A  Q  Q  L  T  S  A  A  A  S  Q  R  I  S  Q  L  S  N
3601  agttcagcacagcaactcacatcagcagcagcatctcaaagaatatctcagttatccaat 3660
      S  L  A  S  A  V  S  N  G  R  V  N  Y  A  A  L  S  N  S  I
3661  tcccttgcatctgctgtttcaaatggacgcgttaactatgcggccttatcgaactctatt 3720
      A  S  A  A  S  Q  I  G  S  G  S  R  L  S  K  T  E  V  L  I
3721  gccagtgctgcaagtcagattggaagtggatctagattgtccaaaactgaagttctaatt 3780
      E  T  L  L  E  T  L  A  A  L  L  E  S  I  S  I  P  G  S  A
3781  gaaactctcttggaaaccttggctgctttattggaatcaatttccatccccggatcagcg 3840
      S  G  G  S  Q  F  A  Q  A  M  L  A  A  L  A  *
3841  agtggtggaagtcaatttgcccaagctatgcttgccgcacttgcataggaatttggttta 3900
3901  taacaattatttgttaaaatcgtggtgaaagtttttgggaattatataacagaatattc 3960
3961  aattaactcttta                                               3974
```

Figure S2. Alignment of Spidroin C-terminal amino acid sequences.