FIG. 17A

```
  1  M  E  K  K  I  T  G  Y  T  T  V  D  I  S  Q  W  H  R  K  E  20
  1  ATGGAGAAAAAAATCACTGGATATACCACCGTTGATATATCCGAATGGCATCGTAAAGAA  60

                    Cleavage site

 21  H  F  E  A  F  Q  S  V  A  Q  C  T  Y  N  Q  T  V  Q  L  D  40
 61  CATTTTGAGGCATTTCAGTCAGTTGCTCAATGTACCTATAACCAGACCGTTCAGCTGGAT  120

 41  I  T  A  F  L  K  T  V  K  K  N  K  H  K  F  Y  P  A  F  I  60
121  ATTACGGCCTTTTTAAAGACCGTAAAGAAAAATAAGCACAAGTTTTATCCGGCCTTTATT  180

 61  H  I  L  A  R  L  M  N  A  H  P  E  F  R  M  A  M  K  D  G  80
181  CACATTCTTGCCCGCCTGATGAATGCTCATGCGGAATTCCGTATGGCAATGAAAGACGGT  240

 81  E  L  V  I  W  D  S  V  H  P  C  Y  T  V  F  H  E  Q  T  E  100
241  GAGCTGGTGATATGGGATAGTGTTCACCCTTGTTACACCGTTTTCCATGAGCAAACTGAA  300

101  T  F  S  S  L  W  S  E  Y  H  D  D  F  R  Q  F  L  H  I  Y  120
301  ACGTTTTCATCGCTCTGGAGTGAATACCACGACGATTTCCGGCAGTTTCTACACATATAT  360

121  S  Q  D  V  A  C  Y  G  E  N  L  A  Y  F  P  K  G  F  I  E  140
361  TCGCAAGATGTGGCGTGTTACGGTGAAAACCTGGCCTATTTCCCTAAAGGGTTTATTGAG  420

141  N  M  F  F  V  S  A  N  P  W  V  S  F  T  S  F  D  L  N  V  160
421  AATATGTTTTTCGTCTCAGCCAATCCCTGGGTGAGTTTCACCAGTTTTGATTTAAACGTG  480

161  A  N  M  D  N  F  F  A  P  V  F  T  M  G  K  Y  Y  T  Q  G  180
481  GCCAATATGGACAACTTCTTCGCCCCCGTTTTCACCATGGGCAAATATTATACGCAAGGC  540

181  D  K  V  L  M  P  L  A  I  Q  V  H  H  A  V  C  D  G  F  H  200
541  GACAAGGTGCTGATGCCGCTGGCGATTCAGGTTCATCATGCCGTCTGTGATGGCTTCCAT  600

201  V  G  R  M  L  N  E  L  Q  Q  Y  C  D  E  W  Q  G  G  A  *  220
601  GTCGGCAGAATGCTTAATGAATTACAACAGTACTGCGATGAGTGGCAGGGCGGGGCGTAA  660
```

FIG. 17B

```
  1  M  E  K  K  I  T  G  Y  T  T  V  D  I  S  Q  W  H  R  K  E  20
  1  ATGGAGAAAAAAATCACTGGATATACCACCGTTGATATATCCCAATGGCATCGTAAAGAA  60

 21  H  F  E  A  F  Q  S  V  A  Q  *  30
 61  CATTTTGAGGCATTTCAGTCAGTTGCTCAATAA  94
                                                N-fragment

  1  M  C  T  Y  N  Q  T  V  Q  L  D  11
  1  ATGTGTACCTATAACCAGACCGTTCAGCTGGAT  33
                                                C-fragment

 12  I  T  A  F  L  K  T  V  K  K  N  K  H  K  F  Y  P  A  F  I  31
 34  ATTACGGCCTTTTTAAAGACCGTAAAGAAAAATAAGCACAAGTTTTATCCGGCCTTTATT  93

 32  H  I  L  A  R  L  M  N  A  H  P  E  F  R  M  A  M  K  D  G  51
 94  CACATTCTTGCCCGCCTGATGAATGCTCATGCGGAATTCCGTATGGCAATGAAAGACGGT  153

 52  E  L  V  I  W  D  S  V  H  P  C  Y  T  V  F  H  E  Q  T  E  71
154  GAGCTGGTGATATGGGATAGTGTTCACCCTTGTTACACCGTTTTCCATGAGCAAACTGAA  213

 72  T  F  S  S  L  W  S  E  Y  H  D  D  F  R  Q  F  L  H  I  Y  91
214  ACGTTTTCATCGCTCTGGAGTGAATACCACGACGATTTCCGGCAGTTTCTACACATATAT  273

 92  S  Q  D  V  A  C  Y  G  E  N  L  A  Y  F  P  K  G  F  I  E  111
274  TCGCAAGATGTGGCGTGTTACGGTGAAAACCTGGCCTATTTCCCTAAAGGGTTTATTGAG  333

112  N  M  F  F  V  S  A  N  P  W  V  S  F  T  S  F  D  L  N  V  131
334  AATATGTTTTTCGTCTCAGCCAATCCCTGGGTGAGTTTCACCAGTTTTGATTTAAACGTG  393

132  A  N  M  D  N  F  F  A  P  V  F  T  M  G  K  Y  Y  T  Q  G  151
454  GCCAATATGGACAACTTCTTCGCCCCCGTTTTCACCATGGGCAAATATTATACGCAAGGC  453

152  D  K  V  L  M  P  L  A  I  Q  V  H  H  A  V  C  D  G  F  H  171
514  GACAAGGTGCTGATGCCGCTGGCGATTCAGGTTCATCATGCCGTCTGTGATGGCTTCCAT  513

172  V  G  R  M  L  N  E  L  Q  Q  Y  C  D  E  W  Q  G  G  A  *  190
574  GTCGGCAGAATGCTTAATGAATTACAACAGTACTGCGATGAGTGGCAGGGCGGGGCGTAA  573
```