```
3601 GGGGCACAGC GGCCCTGGGC TGCCTGGTCA AGGACTACTT CCCCGAACCG
     GTGACGGTGT
                          CH2 Link GlDelCH2
     ----------------------------------------------------
     ---------
3661 CGTGGAACTC AGGCGCCCTG ACCAGCGGCG TGCACACCTT CCCGGCTGTC
     CTACAGTCCT
                          CH2 Link GlDelCH2
     ----------------------------------------------------
     ---------
3721 CAGGACTCTA CTCCCTCAGC AGCGTGGTGA CCGTGCCCTC CAGCAGCTTG
     GGCACCCAGA
                          CH2 Link GlDelCH2
     ----------------------------------------------------
     ---------
3781 CCTACATCTG CAACGTGAAT CACAAGCCCA GCAACACCAA GGTGGACAAG
     AAAGTTGAGC
                          CH2 Link GlDelCH2
     ----------------------------------------------------
     ---------
3841 CCAAATCTTG TGACAAAACT CACACATGCC CACCGTGCCC CATCGAGAAA
     ACCATCTCCA
                          CH2 Link GlDelCH2
     ----------------------------------------------------
     ---------
                                        Bsp1407I
                                        ------
3901 AAGCCAAAGG GCAGCCCCGA GAACCACAGG TGTACACCCT GCCCCCATCC
     CGGGATGAGC
                          CH2 Link GlDelCH2
     ----------------------------------------------------
     ---------
3961 TGACCAAGAA CCAGGTCAGC CTGACCTGCC TGGTCAAAGG CTTCTATCCC
     AGCGACATCG
                          CH2 Link GlDelCH2
     ----------------------------------------------------
     ---------
```

FIG. 6I