



B

GTAGGTTAA GTGTGTGTT
 17651 AAGGCTCGAG ACTTCCTGCC CCACCCAGTG GGTAGGCCAA GTGTGTGCT
 TTAGTA ▶
 17701 TCAGCAAACC GGACCAGCAG GGCCAGGGCC GGATGTGGG ACCCTCTTC
 AGTATAG TAAAGTTGGT TTTT AGAAAT A ▶ BstUI
 17751 TCTAGCACAG TAAAGCTGCC CTCCAGAA ACgGGTATCT CCgCgTGGTG
 MspI
 17801 CTTTGCgGTC gCCgTCgTTG TGGCCgTCCg GGGTGGGGTG TGAGGAGGGG
 BstUI
 17851 ACgAAGGAGG GAAGGAAGGG CAAGGCgGGG GGGGCTCTGC gAGAGCgCgC
 BstUI
 17901 CCAGCCCCgC CTTcGGCCCC CACAGTCCCT GCACCCAGGT TTCCATTGCg
 BstUI
 17951 CgGCTCTCCT CAGCTCCTC CCgCCgCCCA GTCTGGATCC TGGGGGAGGC
 MspI BstUI
 18001 gCTGAAgTcG GGGCCCgCCC TGTGGCCCCg CCCgGCCcGc gCTTGCTAGC
 18051 gCCCAAAGCC AGCgAAGCAC gGGCCCAACC gGGCCATGTC gGGGGAGCCT
 A
 18101 GAGCTCATTG AGCTGCGGGA GCTGGCACCC GCTGGGCGCG CTGGGAAGGG
 CTCAAATAAC TCAACRCCCT CAACCATAA
 18151 CCGCACCCGG CTGGAGCGTG CCAACGGCT GCGCATCGCG CGGGGCACCG