

Fig. 1 (cont.)

(M) Dph-CRE06-CRE04-Csk-SH5 (SEQ ID NO:13)

GGGCCAGGGGACGGTGGCTTCTACGTGCTTGGGACGTTCCCAGCCACCGTCCCATGTTCCCGGCGGGGGGCCA
 GCTGTCCCCACCGCCAGCCCAACTCAGCACTTGGTCAGGGTATCAGCTTGGTGGGGGGGCGTGAGCCCAGCCC
 CTGGGGCGGCTCAGCCCATACAAGGCCATGGGGCTGGGCGCAAAGCATGCCGTGGGTTTCAGGGTGGGTATGGTG
 CGGGAGCAGGGAGGTGAGAGGCTCAGCTGCCCTCCAGAACTCCTCCCTGGGGACAACCCCTCCCAGCCAATAG
 CACAGCCTAGGTCCCCCTATATAAGGCCACGGCTGCTGGCCCTTCCCTTTGGGTGAGTGTACCTCCAGGATAC
 AGACAGCCCCCCTTCAGCCCAGCCCAGCCAGGTACGGCGCGCCGTCGACGGATCCCCCTTTTAGAGAATCCACA
 CCTGTCCAGTTGCTGGGTTCCTACTACCAAAAGTGAATTGCAACTATTTTAGGAGCACTTAAGCACATCCGAA
 AAATGAGTGATTCTGTCTTGGCCACACCACATCACTGATGTACCCCTTAAAGCATGTCCCTGAGTTCATCA
 CAGAAGACTGCTCCTCCTGTGCCCTCCACAAGGTTAGAACTGTCTTGTCTTAGGGAAAAAGGAGAGAGAGAG
 AGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG
 CCCACTCTACTTTACCATAAGTAGCTCCAAATCCTTCTAGAAAATCTGAAAGGCATAGCCCCATATATCAGTG
 ATATAAATAGAACCTGCAGCAGGCTCTGGTAAATGATGACTACAAGGTGGACTGGGAGGCAGCCCGGCCCTTGG
 CAGGCATCATCTCTAAATATAAAGATGAGTTTGTTCAGCCTTTGCAGAAGGAGGATCCGTCGACGGCGCGCC
 ACGCGTCAGTTTACTCACCAGGGATTTCAGAGGCAGCACTGCTGAACCCCTGAGCCCTTGGCACATCAGGTTGGC
 TGTCAGAAGTCGGCCTTTGTACATACACAGTTCCCTTGTGAGGCCAGCTGCGTGTCTTAGGAGCGGGGCCCTC
 TCTCCACAGCAGAGCTCAGCCTCTCAAGTGTATGGACAGCAGGGTGCCGTGATGGGTGGATTTAGCCATGAGT
 TGAAGGTGGCTTGGGGAGAATGAGAGTTCTAGAGATAGGGAGAAGGGTTGCCAATAGGAGAGTGGAATTCCT
 GAGCACCTCGTCACAGGCAGCCGACAGAACATGAGCCGCAGGGCCCAGGCTATTTATACCTCGCCTGTCTACTA
 TCAGGGTCCCCACAGCTCCCCCACCCTCCAGCCACACACAGCAGGTCCCTTTTGTCTTTCTGGTCCCTTCTCT
 ACTCCTCCCCCTCCCTACCTAA

(N) Dph-CRE02-CRE04-CRE06-Csk-SH5 (SEQ ID NO:14)

GACAGGTGCGGTTCCCGGAGCGCAGGCGCACACATGCACCCACCGCGGAACGCGGTGACCCTCGCCCCACCCC
 ATCCCCCTCCGGCGGGCAACTGGGTTCGGGTTCAGGAGGGGCAAACCCGCTAGGGAGACACTCCATATACGGCCCG
 GCGCGGTTACCTGGGACCGGGCCAAACCCGCTCCTTCTTTGGTCAACGCGAGGGGACCCGGGCGGGGGCCAGG
 CCGCGAACCAGGCCGAGGGAGGGGGCTCTAGTGCCCAACACCCAAATATGGCTCGAGAAGGGCAGCGACATTCC
 TGCGGGGTGGCGCGGAGGGAATGCCCCGCGGGCTATATAAAACCTGAGCAGAGGGACAAGCGGCCACCGCAGCG
 GACAGCGCCAAGTGAAGCCTCGCTTCCCCCTCCGCGGCGACCAGGGCCCCGAGCCGAGAGTAGCAGTTGTAGCTA
 CCGCCCCAGGTAGGGGCGCGCCGTACGGTCGACGGATCCCCCTTTTAGAGAATCCACACCTGTCCCAGTTGCT
 GGGTTCCACTACCAAAAGTGAATTGCAACTATTTTAGGAGCACTTAAGCACATCCGAAAAATGAGTGATTCTG
 TTCTGGCCACACCACATCACTGATGTACCCCTTAAAGCATGTCCCTGAGTTCATCACAGAAGACTGCTCCT
 CCTGTGCCCTCCACAAGGTTAGAACTGTCTTGTCTTAGGGAAAAAGGAGAGAGAGAGAGAGAGAGAGAGAGA
 GAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG
 CATAAGTAGCTCCAAATCCTTCTAGAAAATCTGAAAGGCATAGCCCCATATATCAGTGATATAAATAGAACCT
 GCAGCAGGCTCTGGTAAATGATGACTACAAGGTGGACTGGGAGGCAGCCCGGCCCTTGGCAGGCATCATCCTCT
 AAATATAAAGATGAGTTTGTTCAGCCTTTGCAGAAGGAGGCGCGCCGTCGACGGATCCGGGCCAGGGGACGGT
 GGCTTCTACGTGCTTGGGACGTTCCCAGCCACCGTCCCATGTTCCCGGCGGGGGGCCAGCTGTCCCCACCGCC
 AGCCCAACTCAGCACTTGGTCAGGGTATCAGCTTGGTGGGGGGGCGTGAGCCAGCCCTTGGGGCGGCTCAGC
 CCATACAAGGCCATGGGGCTGGGCGCAAAGCATGCCGTGGGTTTCAGGGTGGGTATGGTGCAGGAGCAGGGAGGT
 GAGAGGCTCAGCTGCCCTCCAGAACTCCTCCCTGGGGACAACCCCTCCCAGCCAATAGCACAGCCTAGGTCCC
 CCTATATAAGGCCACGGCTGCTGGCCCTTCCCTTTGGGTGAGTGTACCTCCAGGATACAGACAGCCCCCCTTC
 AGCCAGCCAGCCAGGTACGGATCCGTCGACCGTACGGGCGCGCCACGCGTCAGTTTACTCACCAGGGATTCT
 AGAGGCAGCACTGCTGAACCCCTGAGCCCTTGGCACATCAGGTTGGCTGTCAGAAGTCGGCCCTTTGTACATACA
 CAGTTCCCTTGTGAGGCCCAGCTGCGTGTCTTAGGAGCGGGGCTCTCTCCACAGCAGAGCTCAGCCTCTCAA
 GTGTATGGACAGCACGGGTGCTGATGGGTGGATTTAGCCATGAGTTGAAGGTGGCTTGGGGAGAATGAGAGT
 TCTAGAGATAGGGAGAAGGGGTTGCCAATAGGAGAGTGAATTCCTGAGCACCTCGTCACAGGCAGCCGACAG
 AACATGAGCCGCAGGGGCCAGGCTATTTATACCTCGCCTGTCTACTATCAGGGTCCCCACAGCTCCCCCACCCT
 CCAGCCACACACAGCAGGTCCCTTTTGTCTTTCTGGTCCCTTCTCTACTCCTCCCCCTCCCTACCTAA