Fig. 1

(A) CSk-SH5 (SEQ ID NO:1)

CAGTTTACTCACCAGGGATTCAGAGGCAGCACTGCTGAACCCTGAGCCCTTGGCACATCAGGTTGGCTGTCAG
AAGTCGGCCTTTGTACATACACAGTTCCCTTGTGAGGCCCAGCTGCGTGTCCTAGGAGCGGGGCCTCTCTCCA
CAGCAGAGCTCAGCCTCTCAAGTGTATGGACAGCACGGGTGCCTGATGGGTGGATTTAGCCATGAGTTGAAGG
TGGCTTGGGGAGAATGAGAGTTCTAGAGATAGGGAGAAGGGGTTGCCAATAGGAGAGTGGAATTCCTGAGCAC
CTCGTCACAGGCAGCCGACAGAACATGAGCCGCAGGGCCCAGGCTATTTATACCTCGCCTGTCACTATCAGGG
TCCCCACAGCTCCCCCCACCTCCAGCCACACACAGCAGGTCCTTTTGCTCTTTCTGGTCCCTTCTCTACTCCT
CCCCCTCCCTACCTAA

(B) Dph-CRE02 (SEQ ID NO:2)

GACAGGTGCGGTTCCCGGAGCGCAGGCGCACACATGCACCCACCGGCGAACGCGGTGACCCTCGCCCCACCCC
ATCCCCTCCGGCGGGCAACTGGGTCGGGTCAGGAGGGGCAAACCCGCTAGGGAGACACTCCATATACGGCCCG
GCCCGCGTTACCTGGGACCGGGCCAACCCGCTCCTTCTTTGGTCAACGCAGGGGACCCGGGCGGGGGCCCAGG
CCGCGAACCGGCCGAGGGAGGGGGCTCTAGTGCCCAACACCCAAATATGGCTCGAGAAGGGCAGCGACATTCC
TGCGGGGTGGCGCGGAGGGAATGCCCGCGGGCTATATAAAACCTGAGCAGAGGGACAAGCGGCCACCGCAGCG
GACAGCGCCAAGTGAAGCCTCGCTTCCCCTCCGCGGCGACCAGGGCCCGAGCCGAGAGTAGCAGTTGTAGCTA
CCCGCCCAGGTAGG

(C) Dph-CRE04 (SEQ ID NO:3)

CCTTTTAGAGAATCCACACCTGTCCCAGTTGCTGGGTTCCACTACCAAAAGTGAATTGCAACTATTTTAGGAG
CACTTAAGCACATCCGAAAAATGAGTGATTCTGTTCTGGCCCACACCACATCACTGATGTACCCCCTTAAAGC
ATGTCCCTGAGTTCATCACAGAAGACTGCTCCTCCTGTGCCCTCCACAAGGTTAGAACTGTCCTTGTCTTAGG
GAAAAAGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGGGACAGGCACCAACT
GGGTAACCTCTGCTGACCCCCACTCTACTTTACCATAAGTAGCTCCAAATCCTTCTAGAAAATCTGAAAGGCA
TAGCCCCATATATCAGTGATATAAATAGAACCTGCAGCAGGCTCTGGTAAATGATGACTACAAGGTGGACTGG
GAGGCAGCCCGGCCTTGGCAGGCATCATCCTCTAAATATAAAGATGAGTTTGTTCAGCCTTTGCAGAAGGA

(D) Dph-CRE06 (SEQ ID NO:4)

GGGCCAGGGGACGGTGGCTTCTACGTGCTTGGGACGTTCCCAGCCACCGTCCCATGTTCCCGGCGGGGGGCCA
GCTGTCCCCACCGCCAGCCCAACTCAGCACTTGGTCAGGGTATCAGCTTGGTGGGGGGGCGTGAGCCCAGCCC
CTGGGGCGGCTCAGCCCATACAAGGCCATGGGGCTGGGCGCAAAGCATGCCTGGGTTCAGGGTGGGTATGGTG
CGGGAGCAGGGAGGTGAGAGGCTCAGCTGCCCTCCAGAACTCCTCCCTGGGGACAACCCCTCCCAGCCAATAG
CACAGCCTAGGTCCCCCTATATAAGGCCACGGCTGCTGGCCCTTCCTTTGGGTCAGTGTCACCTCCAGGATAC
AGACAGCCCCCCTTCAGCCCAGCCCAGCCAGGTAC

(E) Dph-CRE02-CRE04 (SEQ ID NO:5)

GACAGGTGCGGTTCCCGGAGCGCAGGCGCACACATGCACCCACCGGCGAACGCGGTGACCCTCGCCCCACCCC
ATCCCCTCCGGCGGGCAACTGGGTCGGGTCAGGAGGGGCAAACCCGCTAGGGAGACACTCCATATACGGCCCG
GCCCGCGTTACCTGGGACCGGGCCAACCCGCTCCTTCTTTGGTCAACGCAGGGGACCCGGGCGGGGGCCCAGG
CCGCGAACCGGCCGAGGGAGGGGGCTCTAGTGCCCAACACCCAAATATGGCTCGAGAAGGGCAGCGACATTCC
TGCGGGGTGGCGCGGAGGGAATGCCCGCGGGCTATATAAAACCTGAGCAGAGGGACAAGCGGCCACCGCAGCG
GACAGCGCCAAGTGAAGCCTCGCTTCCCCTCCGCGGCGACCAGGGCCCGAGCCGAGAGTAGCAGTTGTAGCTA
CCCGCCCAGGTAGGGGCGCGCCGTCGACGGATCCCCTTTTAGAGAATCCACACCTGTCCCAGTTGCTGGGTTC
CACTACCAAAAAGTGAATTGCAACTATTTTAGGAGCACTTAAGCACATCCGAAAAATGAGTGATTCTGTTCTGG
CCCACACCACATCACTGATGTACCCCCTTAAAGCATGTCCCTGAGTTCATCACAGAAGACTGCTCCTCCTGTG
CCCTCCACAAGGTTAGAACTGTCCTTGTCTTAGGGAAAAAGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGA
GAGAGAGAGAGAGAGAGAGGGACAGGCACCAACTGGGTAACCTCTGCTGACCCCCACTCTACTTTACCATAAG
TAGCTCCAAATCCTTCTAGAAAATCTGAAAGGCATAGCCCCATATATCAGTGATATAAATAGAACCTGCAGCA
GGCTCTGGTAAATGATGACTACAAGGTGGACTGGGAGGCAGCCCGGCCTTGGCAGGCATCATCCTCTAAATAT
AAAGATGAGTTTGTTCAGCCTTTGCAGAAGGA