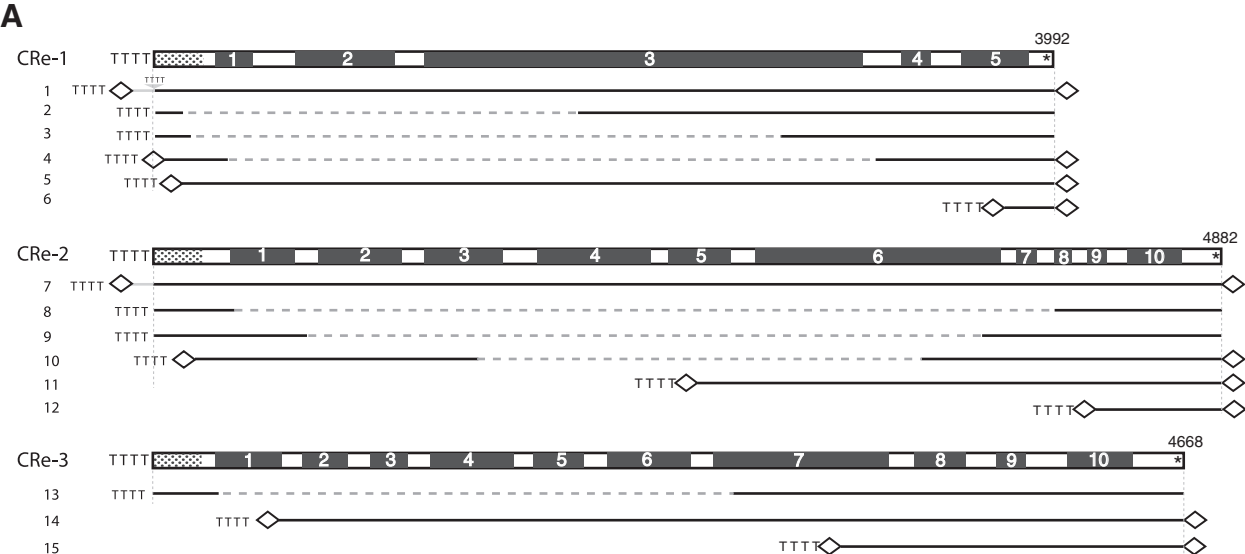




B

> 1 GCGTTTTTCTGGTCTGA<->TCTGGTCTGA

> 4 GTGTTTTTGGTGAAGC<->TGGTGAAGC

> 5 CGTTTTTTAGTGCATC<->TAGTGCATC

> 6 TGATTTTTGCCCGTATGTA<->TGCCCGTATGTA

> 7 GCCTTTTCCCTTGATGGTG<->CCCTTGATGGTG

> 10 TCCTTTTCTCCAGCTGGTG<->CTCCAGCTGGTG

> 12 GCATTTGCACGATCGATGGCAGTACG<->CACGATCGATGGCAGTACG

> 14 GCGTTTTTTTACCCCCCGC<->TTACCCCCCGC

> 15 ACATTTTCTAAACCAGCCCATAGCCG<->CTAATCCAGCCCATAGCCG

D

CRE-1 GGGAGGGGAGGGGCTCAGCCC--TTTTCCTGCCTCCCTAAGGC--AGCCACCT--CCTTGT

CRE-2 GTGGCGTGGCGGCWCAGCCCCACTTTTCC--CTTG--CAAGGGAGAGCCACCTTTCTTGT

CRE-3 ACTGTGGAGGGGCTCAGCCCC--TTTTC--CGTG--ACCGGG--GAA--CA-----CTTGT

Fanzor-1N1_CRe AGCGTACGGGGGCTCAGCCCA--TTTTC--AGCTGTACAAAGC--TGACACC--CCTTGT

Fanzor-1N2_CRe CGCTTCATGGGGCTCAGCCCA--TTTTC--AGCTGTACAAAGC--TGACACC--CCTTGT

Helitron-1_CRe ATCCCGCGGGGCTCAGCCCT--TTTTC--AGCTGTACAAAGC--TGACACC--CCTTGT

-45 -35



AACTGTTTTGCTTTT Helitron-1N1_CRe TGAATAATCCTTTTG

AACTGTTTTGCTTTT ----- TGAATAATCCTTTTG

ABCN01001933.1(17844-17160)

TACTGTGCTTTTTT Helitron-1N1_CRe TGTTTTTCGATTGCT

TACTGTGCTTTTTT ----- TGTTTTTCGATTGCT

ABCN01002965.1(18430-17746)

AGCACAAAGTATTTT Helitron-1N2_CRe CCAACGTACTAGCCG

AGCACAAAGTATTTT ----- CCAACGTACTAGCCG

ABCN01001984.1(12737-13439)

