

pCSA110 construct used in SAIL lines

Left border primers (LB1 is proximal, LB3 is distal and was used for sequencing):

LB1: 5'-GCCTTTTCAGAAATGGATAAATAGCCTTGCTTCC-3'

LB2: 5'-GCTTCCTATTATATCTTCCCAAATTACCAATACA-3'

LB3: 5'-TAGCATCTGAATTCATAACCAATCTCGATACAC-3'

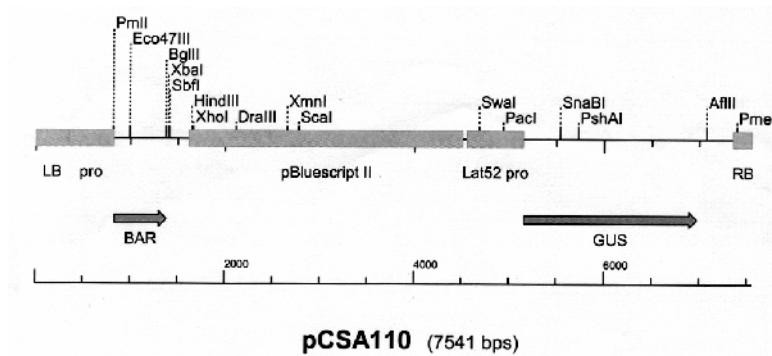
Right border primers:

QRB1: 5' - CAA ACT AGG ATA AAT TAT CGC GCG CGG TGT CA -3'

QRB2: 5' - GGT GTC ATC TAT GTT ACT AGA TCG GGA ATT GA - 3'

QRB3: 5' - CGC CAT GGC ATA TGC TAG CAT GCA TAA TTC - 3'

pCSA110 is described in a paper by John McElver et al in Genetics 159: 1751 (2001).



pCSA110 T-DNA

GGATCCTGATCACAGGCAGCAACGCTCTGTTCATCGTTACAATCAACATGCTACCCCTCCGC
GAGATCATCCGTGTTTCAAACCCGGCAGCTTAGTTGCCGTTCTTCCGAATAGCATCGGTA
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