Sequence: Sox17CR2em3 50 bp deletion.dna (Linear / 1031 bp) Features: 3 total

250 500 750 1000 -50 bp deletion predicted in founder F0-01 $\,$ crRNA rev for m5

Sox17CR2em3 50 bp deletion 1031 bp

TTTTCCTCCGAGGAACAGCACAGTGCTGCTGTCTTTGCACTTTATTTTACATTTTAATCTCGTGGTGTTTGCGATGGAATCATG	+ 85
CCTTGCAAGTCTTTAGAAATAAAACCAGATCTTGCCTCTTTTGCCGAACACACAAAAGGAGCTTTTGTTTTATATTTTTATGTT	- 170
CACACTGAAATTCCTCAGGGTTGACTTGAGCTCTTATTGAGCAGTGATATATTAGTGATATAGCATTTTATGTTAAACTTTGTT 	+ 255
TATTACAACTTGCTATTTACCTTGGATATTTTTATTATAATATAAGCATGATGGATG	+ 340
TTCTCTGAAAAATCATGCTAAGTTAATTACCTTCGGTAATATTTTTTTAAAAGAATGGAATAAATA	+ 425
TGTGAACAACCTTGGACTTGTTCTTCAATCTTCCGTTACAGTGAGCAGCACCTCCAGACATCTGAATTTCAGCCTTCCTATTTC	510
CCAAGAGGTCTTGGCGCCAGCGCCCGGCTCCAGCCAGTTTTCCCCAAGGCAGGTGTAGCCAGAACACTGCCGGGggAGGGGAAA	⊢ 595
GGGCGGGTGTGGGACTTGGGACTCGGACCACTGGCTGGGCggGGGCCTAAGGACTAGGGTCCGGTCC	680
-50 bp deletion predicted in founder F0-01 crRNA rev	for m5
GCCGCGTCCATAAAGAAAAggGCCCCGCGGCCCGGGGGGACACCCGCAGTGTCACTGGGTAGATCTCCAGCGCGGATTAGGCGA 	+ 765
-50 bp deletion predicted in founder F0-01 crRNA rev for m5	
GGGTTAGGTCTACCAGGAGAGCAGCGGGAGGGTAGCACCCGCTCAGCTTTACGAGTTCCTCTGGGCATCTCCAGGCTAGCTTCC	850
ATCCCTGCCTCAGGGTCGGGGGAAGCGGCGTGTCCCGTGGCCATAGCAGAGCTCGGGGTCGGTC	+ 935
GATGCGGGATACGCCAGTGACGACCAGAGCCAGCCCCGGAGCGCGCGC	1020
CTACGCCCTATGCGGTCACTGCTGGTCTCGGTCGGGGCCTCGCGCGTCGGGCGCCACTACCGTCCCAACCCGGGGACAGGGACC	

5 ′ 3 ′