

<b>Construct</b>	<b>Primer</b>	<b>Sequence</b>
<i>B/Man5_8</i>	sense <sup>a</sup>	CTAG <u>CTAGCGCGGACAATTCGCTCCACGTC</u>
	antisense <sup>a</sup>	<u>CGGGATCCTCAGCGAACCTTGCAGCTCGC</u>
<i>B/Man5_8-ΔCBM10</i>	sense <sup>b</sup>	CTCCAATTCGGTGGCT <u><i>A</i></u> GGGTAACACTGG
	antisense <sup>b</sup>	CCAGTGTTACCC <u><i>T</i></u> AGCCACCGGAATTGGAG
<sup>a</sup> The primers are from the 5' ends and the restriction sites are underlined. <sup>b</sup> The mutated basepair at position 1013 is underlined and highlighted in italics.		