

ANEXOS

1. Clon SM-1

T3_CACGCGCAGTACCTCACTAA

GGGACAAAAGCTGGGTACCGGGCCCCCCTCGAGGTCGACGGTATCGATAAGCTTG
ATATCGAATTCGATGACTCCCAGGGTCTGGCCGATGGACTTCAGGTATGTCCCAAGTC
ACTCCAGGCCACCCCCCTTCGAGTCAACGCCAGGTCAACGCGGAGTCAACGGTCA
GAACGGAATATTCCCAATATTCCGACTGTATCCCTGTTGACCGAGTCAAACGGCAT
ATACCGTTGGATTCCGCTCTTCCTTCCGATCGTGAATATGTGAGTTTCAACTCTGAACT
CCACACATGACTCGTGGAAAAGCGTGTATTCCGTTAAAACGGTGCCGCTCCCTCGAC
ATCGGCGACACACGCCGACAGCTCGCGGGATGGCTCTCCCCGACACAACCGCTGG
ACTCCGAGTATGTATGAGTGACCTCAGGCCCTCCACCACCACAGTCCACACACGGTC
AACGCGGGGTCAACGGTCAACGACGGAATATTCCGAGAATATTCCCTGCAAAGTGAA
CCACGTTTTTTCGTCGCCGGGTACGTACAAGTGATACAAGCATTGTGGACGTAATGCA
TGCATCATATCATGCCGCGTTCACAAGTTATCTCGCGTCAAAGGTACATCACCCGCCG
GACGAACGCTTACTGTCTTCGGCGATCCGCCGAAGTCATTCTCCGCGACTACCACTG
CTGTGGAGTAACCAACACCAGCTGTGCTACGTTTCGCACCCTTTCGAGCCAACGCTGC
CACGCCACTTCGCATGGCCGCCACACGGACTCGCGCCACGTGACAGAGCCAAAATA
TAAGTAGAGTCGGCACCGACACCGAGCTA**GGGTTAAGGTAGACTAGATGACTACA**
CCCGGCTGATGAGTCCCAAAAAGGACGAAACCGGTC **T**CGGGTTGTCTCTTGTCTCT
CCCTCGCTCACGAATGAATGAAAATTCATTCCCTCGCAAATTCATTCCCTCGCATCC
TCTACTTGCTCTCGATCACGCCGTTTCGACAACGGCACACGGAGAAATGTGGATTGC
ACAAAGTGGTGCCGCGATCTCGCTGGATTGGATCCACTAGTTCTAGATCGGGTTGTCT
CTTGTCTCTCCCTC

En negrita → Secuencia del promotor T3

En rojo → Secuencia HHR. El sitio de autocorte está marcado con una línea azul

En fondo amarillo → Mutación inactivadora de HHR (T a G)

Subrayado → Sitio XbaI

2. Clon SM-1A

CGCAATTAACCCTCACTAAAGGGGTAGAGTCGGCACCGAGAGTCGGTAGAGTCGGC
ACCGAGCTAGGGTTAAGGTAGACACTAGATGACTACACCCGGCTGATGAGTCCCAA
AAGGACGAAACCGGTCTCGGGTTGTCTCTTGTCTCTCCCTCGCTCACGAATGAATG
AAAATTCATTCCCTCGCATCCTCTACTTGCTCTCGATCACGCCGTTTCGACAACGGCA
CACGGAGAATGTGGATTGCACAAAGTGGTGCCGCGATCTCGCTGGATTGGATCCACT
AGTTCTAGA

3. Clon SM-1B

CGCAATTAACCCTCACTAAAGGGGTAGAGTCGGCACCCGAGCTAGGGTTAAGGTAGAC
ACTAGATGACTACACCCGGCTGATGAGTCCCAAAAAGGACGAAACCGGTCTCGGGTT
GTCTCTTGTCTCTCCCTCGCTCACGAATGAATGAAAATTCATTCCCTCGCATCCTCTA
CTTGCTCTCGATCACGCCGTTTCGACAACGGCACACGGAGAATGTGGATTGCACAAA
GTGGTGCCGCGATCTCGCTGGATTGGATCCACTAGTTCTAGA

4. Clon SM-1C

CGCAATTAACCCTCACTAAAGGGCCGAAGTCATTCTCCGCGACTACCACTGCTGTGG
AGTAACCAACACCAGCTGTGCTACGTTTCGCACCCTTGCGAGCCAACGCTGCCACGC
CACTTCGCATGGCCGCCACACGGACTCGCGCCACGTGACAGAGCCAAAATATAAGT
AGAGTCGGCACCCGAGCTAGGGTTAAGGTAGACACTAGATGACTACACCCGGCTGATG
AGTCCCAAAAAGGACGAAACCGGTCTCGGGTTGTCTCTTGTCTCTCCCTCGCTCAC
GAATGAATGAAAATTCATTCCCTCGCATCCTCTACTTGCTCTCGATCACGCCGTTTCGA
CAACGGCACACGGAGAATGTGGATTGCACAAAGTGGTGCCGCGATCTCGCTGGATTG
GATCCACTAGTTCTAGA