

Table 1. Oligonucleotides used in this study.

Oligonucleotide	Sequence (5'-3') (length [nt]) ^a
1	ggcgat ggatcc gcgGTAA <u>ACTCAAAA</u> ACTCACAGGAGTAATTGAGTAGACTGTAACAGCTATATTTAAGTATGAAAGTGA TCccac gaattc gccc (98)
2	ggcg gaattc gtggGATC <u>ACTTT</u> CATACTTAAATATAGCTGTTACAGTCT <u>ACTCA</u> ATTACTCCTGTGAGTTTTTGAGTTTAC C gcatcc atcgcc (98)
3	ggcgat ggatcc gcgGTAATGTAACAGCTATACAGGAGT a ATTGAGTAGACACACAATTGAGTATTTAAGTATGAAAGTGA Cccac gaattc gccc (98)
4	ggcg gaattc gtggGATC <u>ACTTT</u> CATACTTAAATACTCAATTGTGTGTCTACTCAAT t ACTCCTGTATAGCTGTTACATTACC g catcc atcgcc (98)
5	ggcgat ggatcc gcgGTAA <u>ACTCAAAA</u> ACTCACAGGAGTAATTGAGTAGACACACAATTGAGTATTTTGTAACAGCTATGA TCccac gaattc gccc (98)
6	ggcg gaattc gtggGATCATAGCTGTTACAAAATACTCAATTGTGTGTCTACTCAATTACTCCTGTGAGTTTTTGAGTTTAC C gcatcc atcgcc (98)
7	CATACAAAGAGTATGGTAA <u>ACTCAAAA</u> ACTCACAACTAATCTAGCGTAGACACACAATTGAGTATTTAAGTATGAA <u>AGT</u> GATCTCATTCTGAATCCG (98)
8	CGGATTCAGAAATGAGATC <u>ACTTT</u> CATACTTAAATACTCAATTGTGTGTCTACGCTAGATTAGTTGTGAGTTTTTGAG <u>TTT</u> ACCATACTCTTTGTATG (98)
9	GCTAAGCTTGCATGCCTG (18)
10	ACGAATTCGTGGCATCACCG (20)
11	CCAGTGCCAAGCTTGCATG (19)
12	AACAGCTATGACCATGATTA (20)
13	CATACAAAGAGTATGGTAA (19)
14	CGGATTCAGAAATGAGATC (19)
15	GCACCCAGGCTTTACACTTTATGCTTC (28)
16	CGCCAGGTTTTCCAGTCACGAC (24)
17	CTGTGGCGCCGGTGATGC (18)
18	GGCCAGACAGGCGGTAAGGGC (21)
19	ggcgat ggatcc gcgGTAA <u>ACTCAAAA</u> ACTCACAGccac gaattc gccc (50)
20	ggcg gaattc gtggCTGTGAGTTTTTGAGTTTACCg catcc atcgcc (50)
21	ggcgat ggatcc gcgACAGGAGTAATTGAGTAGACccac gaattc gccc (50)

22	gggcggaattcgtggGTCTACTCAATTACTCCTGTcgcggatccatcgcc (50)
23	ggcgatggatccgcgAGACACACAATTGAGTATTTccacgaattccgccc (50)
24	gggcggaattcgtggAAATACTCAATTGTGTGTCTcgcggatccatcgcc (50)
25	ggcgatggatccgcgATTTAAGTATGAAAGTGATCccacgaattccgccc (50)
26	gggcggaattcgtggGATCACTTTCATACTTAAATcgcggatccatcgcc (50)
27	ggcgatggatccgcgAGACACACAATTGAGTATTTAAGTATGAAAGTGATCcggtgatccacgaattccgccc (74)
28	gggcggaattcgtggcatcaccgGATCACTTTCATACTTAAATACTCAATTGTGTgtctcgcggatccatcgcc (74)
29	ggcgatggatccgcgAGACACACAATTACTCATTccacgaattccgccc (50)
30	gggcggaattcgtggAAATGAGTAATTGTGTGTCTcgcggatccatcgcc (50)
31	AGTATGGTAAACTCAAAACTCACAAactaatcTaGctactataACgA (48)
32	TcGTtataggtagCtAgattagtTGTGAGTTTTTTGAGTTTACCATACT (48)
33	AGTATGGTAAACTCAAAACTCACAGGAGTAATTGAGTAGACACACAA (48)
34	TTGTGTGTCTACTCAATTACTCCTGTGAGTTTTTTGAGTTTACCATACT (48)
35	Cgccatggatcccccagtcacgacgttgtaaacgccacgaattccgccc (50)
36	Gggcgggaattcgtggcggtttacaacgtcgtgactgggggatccatggcg (50)
37	TCGTTATAGGTAGCTAGATTAGTTGTGAGTTTTTTGAGTTgACCATACT (48)
38	AGTATGGTcAACTCAAAACTCACAACTAATCTAGCTACCTATAACGA (48)
39	TCGTTATAGGTAGCTAGATTAGTTGTGAGTTTTTTGAGTgTACCATACT (48)
40	AGTATGGTAcACTCAAAACTCACAACTAATCTAGCTACCTATAACGA (48)
41	TCGTTATAGGTAGCTAGATTAGTTGTGAGTTTTTTGAGgTTACCATACT (48)
42	AGTATGGTAAcCTCAAAACTCACAACTAATCTAGCTACCTATAACGA (48)
43	TCGTTATAGGTAGCTAGATTAGTTGTGAGTTTTTTGAiTTTACCATACT (48)
44	AGTATGGTAAAaTCAAAACTCACAACTAATCTAGCTACCTATAACGA (48)
45	TCGTTATAGGTAGCTAGATTAGTTGTGAGTTTTTTGcGTTTACCATACT (48)
46	AGTATGGTAAAcgCAAAACTCACAACTAATCTAGCTACCTATAACGA (48)
47	TCGTTATAGGTAGCTAGATTAGTTGTGAGTTTTTTtAGTTTACCATACT (48)

48	AGTATGGTAA <u>ACT</u> aAAAACTCACA <u>ACT</u> AATCTAGCTACCTATAACGA (48)
49	TCGTTATAGGTAGCTAGATTAGTTGT <u>GAG</u> TTTTgGAGTTTACCATACT (48)
50	AGTATGGTAA <u>ACT</u> CcAAAACTCACA <u>ACT</u> AATCTAGCTACCTATAACGA (48)
51	TCGTTATAGGTAGCTAGATTAGTTGT <u>GAG</u> TTTgTGAGTTTACCATACT (48)
52	AGTATGGTAA <u>ACT</u> CAcAAAACTCACA <u>ACT</u> AATCTAGCTACCTATAACGA (48)
53	TCGTTATAGGTAGCTAGATTAGTTGT <u>GAG</u> TTgTTGAGTTTACCATACT (48)
54	AGTATGGTAA <u>ACT</u> CAAcAACTCACA <u>ACT</u> AATCTAGCTACCTATAACGA (48)
55	TCGTTATAGGTAGCTAGATTAGTTGT <u>GAG</u> TgTTTGAGTTTACCATACT (48)
56	AGTATGGTAA <u>ACT</u> CAAAcACTCACA <u>ACT</u> AATCTAGCTACCTATAACGA (48)
57	TCGTTATAGGTAGCTAGATTAGTTGT <u>GAG</u> gTTTTGAGTTTACCATACT (48)
58	AGTATGGTAA <u>ACT</u> CAAAAcCTCACA <u>ACT</u> AATCTAGCTACCTATAACGA (48)
59	TCGTTATAGGTAGCTAGATTAGTTGT <u>G</u> AtTTTTTGAGTTTACCATACT (48)
60	AGTATGGTAA <u>ACT</u> CAAAAAaTCACA <u>ACT</u> AATCTAGCTACCTATAACGA (48)
61	TCGTTATAGGTAGCTAGATTAGTTGT <u>G</u> cTTTTTGAGTTTACCATACT (48)
62	AGTATGGTAA <u>ACT</u> CAAAAACgCACA <u>ACT</u> AATCTAGCTACCTATAACGA (48)
63	TCGTTATAGGTAGCTAGATTAGTTGT <u>A</u> GTTTTTGAGTTTACCATACT (48)
64	AGTATGGTAA <u>ACT</u> CAAAA <u>ACT</u> aACA <u>ACT</u> AATCTAGCTACCTATAACGA (48)
65	TTGTGTGTCT <u>ACT</u> CAATTACTCCTGT <u>GAG</u> TTTTTGAGTTgACCATACT (48)
66	AGTATGGTcAACTCAAAA <u>ACT</u> CACAGGAGTAATTGAGTAGACACACAA (48)
67	TTGTGTGTCT <u>ACT</u> CAATTACTCCTGT <u>GAG</u> TTTTTGAGTgTACCATACT (48)
68	AGTATGGTAcACTCAAAA <u>ACT</u> CACAGGAGTAATTGAGTAGACACACAA (48)
69	TTGTGTGTCT <u>ACT</u> CAATTACTCCTGT <u>GAG</u> TTTTTGAGgTTACCATACT (48)
70	AGTATGGTAAcCTCAAAA <u>ACT</u> CACAGGAGTAATTGAGTAGACACACAA (48)
71	TTGTGTGTCT <u>ACT</u> CAATTACTCCTGT <u>GAG</u> TTTTGAtTTTACCATACT (48)
72	AGTATGGTAAAaTCAAAA <u>ACT</u> CACAGGAGTAATTGAGTAGACACACAA (48)

73	TTGTGTGTCT <u>ACTCAATTACTCCTGTGAGTTTTT</u> GcGTTTACCATACT (48)
74	AGTATGGTAAACgCAAAA <u>ACTCACAGGAGTAATTGAGTAGACACACAA</u> (48)
75	TTGTGTGTCT <u>ACTCAATTACTCCTGTGAGTTTTt</u> AGTTTACCATACT (48)
76	AGTATGGTAA <u>ACTa</u> AAAA <u>ACTCACAGGAGTAATTGAGTAGACACACAA</u> (48)
77	TTGTGTGTCT <u>ACTCAATTACTCCTGTGAGTTTTg</u> GAGTTTACCATACT (48)
78	AGTATGGTAA <u>ACTCc</u> AAAA <u>ACTCACAGGAGTAATTGAGTAGACACACAA</u> (48)
79	TTGTGTGTCT <u>ACTCAATTACTCCTGTGAGTTTg</u> TGAGTTTACCATACT (48)
80	AGTATGGTAA <u>ACTCAc</u> AAAA <u>ACTCACAGGAGTAATTGAGTAGACACACAA</u> (48)
81	TTGTGTGTCT <u>ACTCAATTACTCCTGTGAGTTg</u> TTGAGTTTACCATACT (48)
82	AGTATGGTAA <u>ACTCAAc</u> AA <u>ACTCACAGGAGTAATTGAGTAGACACACAA</u> (48)
83	TTGTGTGTCT <u>ACTCAATTACTCCTGTGAGTg</u> TTTGAGTTTACCATACT (48)
84	AGTATGGTAA <u>ACTCAAAc</u> ACTCACAGGAGTAATTGAGTAGACACACAA (48)
85	TTGTGTGTCT <u>ACTCAATTACTCCTGTGAGg</u> TTTTGAGTTTACCATACT (48)
86	AGTATGGTAA <u>ACTCAAAAc</u> CTCACAGGAGTAATTGAGTAGACACACAA (48)
87	TTGTGTGTCT <u>ACTCAATTACTCCTGTGAi</u> TTTTGAGTTTACCATACT (48)
88	AGTATGGTAA <u>ACTCAAAAAa</u> TCACAGGAGTAATTGAGTAGACACACAA (48)
89	TTGTGTGTCT <u>ACTCAATTACTCCTGTGc</u> GTTTTT <u>GAGTTTACCATACT</u> (48)
90	AGTATGGTAA <u>ACTCAAAAAc</u> gCACAGGAGTAATTGAGTAGACACACAA (48)
91	TTGTGTGTCT <u>ACTCAATTACTCCTGTt</u> AGTTTTT <u>GAGTTTACCATACT</u> (48)
92	AGTATGGTAA <u>ACTCAAAAAc</u> TaACAGGAGTAATTGAGTAGACACACAA (48)
93	TTGTGTGTaT <u>ACTCAATTACTCCTGTGAGTTTTT</u> GAGTgACCATACT (48)
94	AGTATGGTcAA <u>ACTCAAAAAc</u> ACTCACAGGAGTAATTGAGTAtACACACAA (48)
95	TTGTGTGTCg <u>ACTCAATTACTCCTGTGAGTTTTT</u> GAGTgTACCATACT (48)
96	AGTATGGTAc <u>ACTCAAAAAc</u> ACTCACAGGAGTAATTGAGTcGACACACAA (48)

97	TTGTGTGTCT <u>c</u> CTCAATTACTCCTGTGAGTTTTTGAGgTTACCATACT (48)
98	AGTATGGTAA <u>c</u> CTCAAAA <u>A</u> CTCACAGGAGTAATTGAGgAGACACACAA (48)
99	TTGTGTGTCT <u>Aa</u> TCAATTACTCCTGTGAGTTTTTGAtTTTACCATACT (48)
100	AGTATGGTAA <u>Aa</u> TCAAAA <u>A</u> CTCACAGGAGTAATTGAtTAGACACACAA (48)
101	TTGTGTGTCT <u>ACg</u> CAATTACTCCTGTGAGTTTTTGcGTTTACCATACT (48)
102	AGTATGGTAA <u>Ac</u> GCAAAA <u>A</u> CTCACAGGAGTAATTGcGTAGACACACAA (48)
103	TTGTGTGTCT <u>ACTa</u> AATTACTCCTGTGAGTTTTTtAGTTTACCATACT (48)
104	AGTATGGTAA <u>ACTa</u> AAAA <u>A</u> CTCACAGGAGTAATTtAGTAGACACACAA (48)
105	TTGTGTGTCT <u>ACTCc</u> ATTACTCCTGTGAGTTTTgGAGTTTACCATACT (48)
106	AGTATGGTAA <u>ACTCc</u> AAAA <u>A</u> CTCACAGGAGTAATgGAGTAGACACACAA (48)
107	TTGTGTGTCT <u>ACTCAc</u> TTACTCCTGTGAGTTTgTGAGTTTACCATACT (48)
108	AGTATGGTAA <u>ACTCAc</u> AA <u>A</u> CTCACAGGAGTAAgTGAGTAGACACACAA (48)
109	TTGTGTGTCT <u>ACTCAAg</u> TACTCCTGTGAGTTgTTGAGTTTACCATACT (48)
110	AGTATGGTAA <u>ACTCAAc</u> AA <u>A</u> CTCACAGGAGTAcTTGAGTAGACACACAA (48)
111	TTGTGTGTCT <u>ACTCAATg</u> ACTCCTGTGAGTgTTTGAGTTTACCATACT (48)
112	AGTATGGTAA <u>ACTCAAAc</u> ACTCACAGGAGTcATTGAGTAGACACACAA (48)
113	TTGTGTGTCT <u>ACTCAATTc</u> CTCCTGTGAGgTTTTGAGTTTACCATACT (48)
114	AGTATGGTAA <u>ACTCAAAAc</u> CTCACAGGAGgAATTGAGTAGACACACAA (48)
115	TTGTGTGTCT <u>ACTCAATTAa</u> TCCTGTGAtTTTTTGAGTTTACCATACT (48)
116	AGTATGGTAA <u>ACTCAAAAAa</u> TCACAGGAtTAATTGAGTAGACACACAA (48)
117	TTGTGTGTCT <u>ACTCAATTACg</u> CCTGTGcGTTTTTGAGTTTACCATACT (48)
118	AGTATGGTAA <u>ACTCAAAAACg</u> CACAGGcGTAATTGAGTAGACACACAA (48)
119	TTGTGTGTCT <u>ACTCAATTACTa</u> CTGTtAGTTTTTGAGTTTACCATACT (48)
120	AGTATGGTAA <u>ACTCAAAAACTa</u> ACAGtAGTAATTGAGTAGACACACAA (48)

121	gatcc atattaacctttactatacaaaagagtatGGTAA <i>tgT</i> <u><i>aAc</i></u> <i>Agctat</i> ACAGGAGTAATTGAGTAGAC (72)
122	<i>atagcTgTtAca</i> TTACCatactctttgatgagtaaaggttaatatgg (48)
123	<u>ACACAATTGAGTATTTAAGTATGAAAGT</u> GATCtcatttctgaatccgaaagg (52)
124	gatcc ctttcggattcagaaatgaGATCACTTTCATACTTAAATACTCAATTGIGTGTCTACTCAATTACTCCTGT (76)
125	gatcc atattaacctttactatacaaaagagtatGGTAA <u>ACTCAAAA</u> ACTCACAGGAGTAATTGAGTAGAC (72)
126	<u>GAGTTTTT</u> GAGTTTACCatactctttgatgagtaaaggttaatatgg (48)
127	<i>tgtaA</i> <u><i>cagctaT</i></u> ATTTAAGTATGAAAGT <u>GATC</u> tcatttctgaatccgaaagg (52)
128	gatcc ctttcggattcagaaatgaGATCACTTTCATACTTAAAT <u><i>AtagctgT</i></u> <u><i>taca</i></u> GTCTACTCAATTACTCCTGT (76)
129	<u>ACACAATTGAGTATTT</u> <i>tgtaATGgcta</i> TGATCtcatttctgaatccgaaagg (52)
130	gatcc ctttcggattcagaaatgaGATCA <u><i>atagcCAT</i></u> <u><i>taca</i></u> AAATACTCAATTGIGTGTCTACTCAATTACTCCTGT (76)

^aNucleotides derived from O_F are in upper case, and other nucleotides are in lower case. Restriction enzyme recognition sites or sticky ends are in bold, and subsites A-D are underlined. Nucleotides in bold and underlined in oligonucleotides 29/30 highlight the inverted half-site in subsite C. Mutations in oligonucleotides 121-130 are italicized.