

Supplementary Material

Table 1. Mutagenic oligomer pairs used to generate truncation and point mutant Tm-

Cyto constructs.

Name	Sequence
Δ 824–839	Top – CGCCA <u>ACT</u> GAAAGTCAAGTGA <u>AC</u> GGTCCCCAATGAAGCC Bottom – GGCTTCATTGGGGACCGT <u>TCA</u> CTTGACTTCAGTTGGCG
Δ 801–839	Top – CCACTGACAGAGCCCGAGTGA <u>A</u> AGGGTCCTGTAGAAAC Bottom – GTTCTACAGGACCCTT <u>TCA</u> CTCGGGCTCTGTCAAGTGG
Δ 791–839	Top – GGAAGCACACAGAGCCCTGA <u>A</u> ACGAGACCACACCAC Bottom – GTGGTGTGGTCTCGT <u>TCA</u> GGGCTCTGTGTGCTTCC
Δ 780–839	Top – GGAGGAGGAACGGACTTG <u>A</u> CCAAACCATGACGGAGG Bottom – CCTCCGTCATGGTTTGGT <u>CA</u> AGTCCGTTCCCTCCTCC
Δ 745–839	Top – CTGTGCGGCAAAGCGTGA <u>G</u> GGGCCCGGAGCC Bottom – GGCTCCGGGCCCT <u>CAC</u> GCTTTGCCGCACAG
T788A	Top – GAGGGAAGCAC <u>G</u> CAGAGCCCAACGAG Bottom – CTCGTTGGGCTCTG <u>C</u> GTGCTTCCCTC
T793A	Top – GAGCCCAACGAGG <u>C</u> CACACCACTGAC Bottom – GTCAGTGGTGTGG <u>C</u> CTCGTTGGGCTC
T794A	Top – CCCAACGAGACCG <u>C</u> ACCACTGACAGAG Bottom – CTCTGTCAGTGGT <u>G</u> CGGTCTCGTTGGG
T797A	Top – CCACACCACTGGCAGTGCCCGAGAAG Bottom – CTTCTCGGGCTCTG <u>C</u> CAGTGGTGTGG
T788D	Top – GACGGAGGGAAGCAC <u>GAC</u> GAGCCCAACGAGAC Bottom – GTCTCGTTGGGCTC <u>GTC</u> GTGCTTCCCTCCGTC
T794D	Top – GAGCCCAACGAGACCG <u>A</u> CCCACTGACAGAGCCC Bottom – GGGCTCTGTCAGTGGG <u>T</u> CGGTCTCGTTGGGCTC
T797D	Top – GAGACCACCACTGGAC <u>G</u> AGCCCGAGAAGGG Bottom – CCCTTCTCGGGCTC <u>GTC</u> CAGTGGTGTGGTCTC

Underlined nucleotides indicate added or changed bases.