

Supplementary Table 2. Primer pairs used in Sanger sequencing

Gene	Location	Position	AA change	Forward primer (5'-3')	Reverse primer (5'-3')
<i>CDH1</i>	chr16	68863688	p.N809fs	CGTTGCACCAACCCTCAT	GCCATGCTTTGGCTTCC
<i>ERBB2</i>	chr17	37872159	p.L494I	GGCCCTCATCCACCATAAC	ACTGGGCTCCCTGTCTTA
<i>ESR1</i>	chr6	152382171	p.M427I	AAATGTGTAGAGGGCATGGT	TTGAGGCACACAAACTCCT

AA change, amino acid change.