

1 **Supplementary Table 1.** Primers used in the site-directed mutageneses

Mutation	Primer	Sequence (5'→3')
G16E	Forward	cttagttacagtaaaatttagaggcacagctgatagaagcc
	Reverse	ggcttctatcagctgtccctctaatttactgtaactaag
G17E	Forward	cttagttacagtaaaatttagggAACAGCTGATAGAAGCC
	Reverse	ggcttctatcagctgtccctctaatttactgtaactaag
I64M	Forward	gacaatatgatcagatacttatggaaatttgtggaaaaaggc
	Reverse	gccttttccacaaattccataagtatctgatcatattgtc
K70R	Forward	cttata <u>gaaattt</u> tgggaaaaggctataggtagtacgtttaggg
	Reverse	cctactaac <u>actgt</u> acctatagcc <u>tttccacaaatt</u> tataag
I72V	Forward	gaaattt <u>tg</u> ggaaaaagg <u>ctgt</u> aggtagtacgtttaggg
	Reverse	cctactaac <u>actgt</u> acctac <u>aggc</u> tttccacaaatttc

2 * Underlined bases correspond to the changes made to the original BD6-15 clone.

Supplementary Table 2. Primers used in the oligonucleotide ligation assays

Protease codon	Upstream primer	Target	Downstream primer
17	5'-GTTACAGTAAAATTAGGG <u>G</u> A-3'	17E	5'-CAGCTGATAGAAGCCTTAT-3'
	5'-GTTACAGTAAAATTAGGG <u>G</u> -3'	17G	
64	5'-AATATGATCAGATA <u>C</u> TATG-3'	64M	5'-GAAATTGTGGAAAAAAAGGC-3'
	5'-AATATGATCAGATA <u>T</u> TA-3'	64I	
72	5'-AATTGTGGAAAAAAAGG <u>C</u> TG-3'	72V	5'-TAGGTACAGTGTAGTAGGA-3'
	5'-AATTGTGGAAAAAAAGG <u>C</u> A-3'	72I	

* Underlined bases correspond to the discriminating nucleotide position between the two variants present in each head-to-head competition.