

Table S2. Primers used in this study

Name	Orientation	Sequence (5' to 3')	Description
NF1558	Forward	GCTTTTGTAACTATGGAAAAA TAACTAACTTAAAGG	T413A mutation in CwpV cleavage site by inverse PCR
NF1559	Forward	TGTTTTGTAACTATGGAAAAA TAACTAACTTAAAGG	T413V mutation in CwpV cleavage site by inverse PCR
NF1560	Forward	AGTTTTGTAACTATGGAAAAA TAACTAACTTAAAGG	T413S mutation in CwpV cleavage site by inverse PCR
NF1561	Forward	GTTTTTGTAACTATGGAAAAA TAACTAACTTAAAGG	T413C mutation in CwpV cleavage site by inverse PCR
NF1562	Reverse	TCCATCTTTTACATCATCTATAA CTACATC	Mutation in CwpV cleavage site by inverse PCR. Universal reverse primer
NF1599	Forward	CCATGGGACAAACTGTGGC	Amplification of CwpV fragment (aa 28-626) for pET28a cloning
NF1600	Reverse	CTCGAGTGAATCTGTAGCTCC	Amplification of CwpV fragment (aa 28-626) for pET28a cloning
NF1605	Forward	GGAGGAGGAACTTTTGTAACT ATGGAAAAATAACTAACTAAA GG	K411Gly; D412G mutation in CwpV cleavage site by inverse PCR
NF1606	Reverse	TACATCATCTATAACTACATCA TCCACTTTTGCATAGTTATTAC ATC	K411G; D412G mutation in CwpV cleavage site by inverse PCR
NF1622	Forward	CAATTTGAAAAATGAGATCCGG CTGCTAACAAAG	<i>Strep</i> -tagging a CwpV fragment (aa 28-611)
NF1623	Reverse	TGGATGACTCCACACACTTACA TTATTTAAATCATTTTGTATT	<i>Strep</i> -tagging a CwpV fragment (aa 28-611)