

1 **Table S2.** Primers (F is forward, and R is reverse).

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Primer	Sequence (5'–3') ^a	5'- Restriction site
rFnBPA(194–511) F	CGC <u>GGATCC</u> GGCACAGATGTAACAAGTAAAG	BamHI
rFnBPA(194–511) R	GACGTCGACTTAATTTTTCTCATTTCGGTTCTC	SaI
rFnBPA(194–336) F	CGC <u>GGATCC</u> GGCACAGATGTAACAAGTAAAG	BamHI
rFnBPA(194–336) R	GACG <u>GTCGACT</u> TATTTAACATCTAATTCCTTTG	SaI
rFnBPA(337–511) F	CGC <u>GGATCC</u> TATAAAGATGGTATTGGG	BamHI
rFnBPA(337–511) R	GACGTCGACTTAATTTTTCTCATTTCGGTTCTC	SaI

3 ^aRestriction sites used for cloning are underlined.

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