

Characterization of 25 full-length *S-RNase* alleles, including flanking regions, from a pool of resequenced apple cultivars

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**Online Resource 1** List of primers used in this study.

Primer	5'-3' sequence
PycomC1f <sup>a</sup>	atTTTcaatttacgcagcaatatcagc
PycomC5r <sup>a</sup>	ctgcaaagwshgacctcaaccaattc
MdS8_5UTR	ttgcgcgtagagattccgat
MdS8_Ex1rev	actctgggtcaggtgcattc
MdS8_Ex1for	ggccttcgaacaggaatgc
MdS8_Ex2for	cgaacgggacaaacaggc
MdS8_3UTR	ggctggagtatggggtgtag
MdS23_5UTR	tgagactagcgtgttcccag
MdS23_3UTR	aggccaacattttacacgagt
MdS58_5UTR	ccacaaggctcatggtgaat
MdS58_3UTR	taactgcatttctctcgccc
MdS50_3UTR	gcgccgataagtttagggt
MdS6a_Ex1for	tgcaagactacagccctga
MdS6a_UTR1	ttatgtatactagccagccg
MdS6a_UTR2	cataactgtattcatctcttctat
MdS6a_UTR3	agtcttccaatttatgcattagaa
MdProm-box1f	arggcabtgacatga

<sup>a</sup> from Sanzol, 2009a.