

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

Plant Molecular Biology

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**Online resource 7** Test amplification using primers MdProm-box1f and PycomC5r from several genotypes of *Pyrus communis* (Abbé Fétel, Kaiser, Max Red Bartlett), *P. pyrifolia* (Hayatama, Nijisseiki, Yakumo) and *P. bretschneideri* (Xue Hua Li) with apple 'Golden Delicious' used as control.

