

Supplemental Data for:

Migliaro D, De Lorenzis G, Di Lorenzo GS, De Nardi B, Gardiman M, Failla O, Brancadoro L and Crespan M. 2019.

Grapevine non-*vinifera* genetic diversity assessed by simple sequence repeat markers as a starting point for new rootstock breeding programs. *Am J Enol Vitic* 70:390-397. doi: 10.5344/ajev.2019.18054.

Supplemental Table 2 Three multiplex panels for PCR.

SSR^a marker	Multiplex panel	Dye	Labeled primer concn (μM)	Size range in basepairs
VVS2	1	6-Fam	0.20	123-171
VVMD7		6-Fam	0.20	227-265
VVMD27		6-Fam	0.20	175-223
VVMD28		Pet	0.24	219-285
VrZAG62		Vic	0.11	172-219
VrZAG79		Ned	0.11	236-280
VMC6E1		Vic	0.14	117-171
VVMD25	2	6-Fam	0.16	238-277
VrZAG64		6-Fam	0.20	127-193
VrZAG83		Vic	0.25	158-218
VChr4a		Pet	0.30	176-210
VMCNG4b9		Ned	0.14	138-180
VVMD5	3	Vic	0.48	218-276
VChr3a		6-Fam	0.20	131-302
VChr7b		Ned	0.15	167-211
VChr10b		Pet	0.20	114-150
VChr19a		Vic	0.20	120-165

^aSSR, simple sequence repeat.