

Supplementary Table S3. SSR marker and PCR multiplex conditions used. The set of six highly reproducible microsatellites suggested by the European group working within the grape GENRES projects is indicated in bold.

SSR marker	Forward primer	Reverse primer	Reference	Core repeat	Annealing T °C (max-min)	Dye	PCR multiplex
VVS2	CAG CCC GTA AAT GTA TCC ATC	AAATTCAAAATTCTAATTCAACTGG	Thomas and Scott (1993)	(GA) _n	57-50	YAKYE	3
VVMD5	CTAGAGCTACGCCAATCCAA	TATACCAAAAATCATATTCCTAAA	Bowers et al. (1996)	(CT) _n AT(CT) _n ATAG(AT) _n	62-55	AT565	2
VVMD6	ATCTCTAACCCCTAAAACCAT	CTGTGCTAAGACGAAGAAGA	Bowers et al. (1996)	(CT) _n C(CT) _n TTAG(CT) _n TAAT-(CT) _n C(CT) _n C(CT) _n	63-56	AT565	1
VVMD7	AGAGTTGCGGAGAACAGGAT	CGAACCTTCACACGCTTGAT	Bowers et al. (1996)	(CT) _n	62-55	AT550	2
VVMD17	TGACTCGCCAAAATCTGACG	CACACATATCATCACCACACGG	Bowers et al. (1999)	(CT) _n	57-50	AT565	3
VVMD21	GGTTGTCTATGGAGTTGATGTTGC	GCTTCAGTAAAAAGGGATTGCG	Bowers et al. (1999)	(CT) _n GAGAAGG(A) _n	62-55	YAKYE	4
VVMD24	GTGGATGATGGAGTAGTCACGC	GATTTAGGTTTCATGTTGGTGAAGG	Bowers et al. (1999)	(CT) _n	61-54	YAKYE	6
VVMD25	TTCCGTTAAAGCAAAAGAAAAAGG	TTGGATTGAAATTTATTGAGGGG	Bowers et al. (1999)	(CT) _n	57-50	AT550	3
VVMD27	GTACCAGATCTGAATACATCCGTAAGT	ACGGGTATAGAGCAAAACGGTGT	Bowers et al. (1999)	(CT) _n	62-55	FAM	2
VVMD28	AACAATTCAATGAAAAGAGAGAGAGAGA	TCATCAATTCGTATCTCTATTTGCTG	Bowers et al. (1999)	(CT) _n	62-55	YAKYE	2
VVMD32	TATGATTTTTAGGGGGGTGAGG	GGAAGATGGGATGACTCGC	Bowers et al. (1999)	(CT) _n	63-56	AT550	1
VrZAG62	GGTGAAATGGGCACCGAACACACGC	CCATGTCTCTCCTCAGTTCTCAG	Sefc et al. 1999	(GA) _n	63-56	YAKYE	1
VrZAG79	AGATTGTGGAGGAGGGAACAAACCG	TGCCCCCATTTTCAAACCTCCCTTC	Sefc et al. 1999	(GA) _n	63-56	FAM	1
VMC1b11	CTTTGAAAATTCCTTCGGGTT	TATTCAAAGCCACCCGTTCTCT	Zyprian and Töpfer (2005)	(GA) _n	62-55	FAM	4
VMC4f3.1	AAAGCACTATGGTGGGTGTAAA	TAACCAATACATGCATCAAGGA	Di Gaspero et al. (2000)	(CT) _n TT (CT) _n	60-53	FAM	5
VV1b01	TGACCCTCGACCTTAAATCTT	TGGTGAGTGCAATGATAGTAGA	Merdinoglu et al. (2005)	(CT) _n	60-53	AT565	5
VV1h54	CCGCACTTGTGTTGAATTTTACAG	CAAAACCGTTTTACACCCAGCAG	Merdinoglu et al. (2005)	(GA) _n	61-54	FAM	6
VV1n16	ACCTCTATAAGATCCTAACCTG	AAGGGAGTGTGACTGATATTTT	Merdinoglu et al. (2005)	(CA) _n CG (CA) _n	62-55	AT565	4
VV1n31	TACTTCACCTAACAATACAGCT	AATACATAAGGTGAAGATGCCT	Merdinoglu et al. (2005)	(CA) _n	60-53	AT550	5
VV1p31	TATCCAAGAGACAAATTTCCAC	TTCTCTGTTTCTGCAATATGG	Merdinoglu et al. (2005)	(GA) _n	61-54	AT550	6
VV1p60	GGGGAATAACTAAATTGAGGAT	GTATGAATGCGGATAGTTTGTG	Merdinoglu et al. (2005)	(TG) _n AT(GT) _n (GA) _n	57-50	FAM	3
VV1q52	TAAAAGGATGGTAGATGACAGA	ACAGGAAAGTGTCAATGGTTA	Merdinoglu et al. (2005)	(CT) _n	62-55	AT550	4
VV1v67	TATAACTTCTCATAGGGTTTCC	TTGGAGTCCATCAAATTCATCT	Merdinoglu et al. (2005)	(CA) _n AT(CA) _n (GA) _n TT(GA) _n (AG) _n	61-54	AT565	6

SSR markers with the same PCR multiplex number were amplified in a single PCR mix, all primers being pooled in the PCR mix and analyzed in the same sequencer run