Erratum to: Morphological Variability in Leaves and Molecular Characterization of Novel Table Grape Candidate Cultivars (*Vitis vinifera* L.)

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Table 7 of the original version of the article unfortunately contains an error. The presentation of values in Table 7 was incorrect and the corrected table is given below.

The online version of the original article can be found under doi:[10.1007/s12033-013-9729-6](https://doi.org/10.1007/s12033-013-9729-6).

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Table 7 Allele size in relative base pair distance to allele size *n* as coded by OIV descriptors list of twenty-six table grape genotypes at six microsatellite loci

	VVS2		VVMD5		VVMD7		VVMD27		VrZAG62		VrZAG79	
	OIV801		OIV802		OIV803		OIV804		OIV805		OIV806	
89IxU119	12	26	12	16	16	24	4	20	18	30	18	22
Alzey real	16	20	10	14	8	8	10	10	14	30	14	14
Bellini	28	32	6	16	8	16	4	10	14	18	2	24
Cardinal n.	12	12	4	14	18	18	4	10	12	12	14	18
Ceresa	10	26	6	18	18	18	10	20	12	20	10	14
Conegliano 218	10	28	14	16	16	18	4	20	12	18	18	20
Dalmaso VII-6	10	20	4	6	8	18	4	10	20	30	18	18
Dalmaso XIII-11	12	14	16	16	12	16	4	6	14	20	18	22
Dalmaso XXIII-12	12	14	4	16	16	18	4	6	18	30	18	22
Delizia di Vaprio	10	10	4	10	18	20	20	20	12	30	10	20
Gargiulo 102011	12	26	6	12	8	18	4	6	12	14	10	18
Gargiulo 88086	26	28	6	12	8	18	6	20	14	30	10	20
Gargiulo 88435	10	12	14	14	18	20	10	18	12	26	14	20
Gargiulo 90393	12	22	14	16	12	18	10	18	12	14	20	20
IxES 21	10	12	4	10	12	18	4	19	12	30	18	18
IxFT 87	10	26	10	12	8	12	4	10	14	30	14	18
ISV IxV93	10	28	6	16	8	16	4	10	18	20	2	18
Italia	10	26	10	16	12	16	4	19	18	30	18	20
Pirovano 100	10	14	6	16	18	18	6	19	28	30	8	20
Pirovano 244	10	12	10	14	16	18	4	19	18	30	20	22
Pirovano 432	10	26	10	16	18	18	4	4	12	30	18	18
Pirovano 727	12	26	6	10	8	16	6	19	14	18	20	20
Prosperi 190	12	26	10	16	8	16	10	19	14	18	6	20
Sugraone	12	12	4	14	8	18	4	6	12	14	18	22
Unterkofler 50	10	20	4	16	20	32	4	8	20	30	10	18
Valsesia 54	10	12	10	18	16	18	2	18	18	30	18	20
H_o^a	0.88		0.92		0.81		0.88		0.96		0.77	
H_E^b	0.81		0.84		0.76		0.82		0.81		0.79	
Na^c	9		7		7		8		7		9	
NDG ^d	13		16		11		14		13		15	

^a Observed heterozygosity^b Expected heterozygosity^c Observed number of alleles^d Number of different genotypes

Homozygous loci are reported in bold