

Vitis 46 (4), 192–194 (2007)

'Vitouska' is the progeny of 'Prosecco tondo' and 'Malvasia bianca lunga'

M. Crespan, G. Crespan, S. Giannetto, S. Meneghetti, and A. Costacurta

C.R.A. Centro di ricerca per la Viticoltura, Susegana (TV), Italy

Summary

'Vitouska' is a minor white wine grape variety, cultivated in the Kars region and recently recovered and revaluated by local viticulturists. Its obscure origins, traditionally linked to this land, are now supported by the results obtained from the present research, which identified the two parents, 'Malvasia bianca lunga' alias 'Malvasia del Chianti' and 'Prosecco tondo'. The latter cultivar has an ancient and well-documented presence in the same growing region as 'Vitouska'. Molecular analyses have been performed with 37 nuclear microsatellite loci.

K e y w o r d s : 'Vitovska', 'Garganija', parentage, SSR markers, minor cultivar.

Introduction

'Vitouska', known also as 'Vitovska', 'Vitovska-Garganija', 'Gargania', is a minor white wine grape cultivar. It is a vigorous and late-ripening variety, with compact cluster and big berries, which found an ideal environment in the red soils of the Kars region (North East Italy and West Slovenia). It proved to have an optimal equilibrium between vegetative and productive activity in this ferrous and poor fertile land, giving refined and sapid wines. Almost rustic, it can tolerate the cold winters with the strong Bora winds, as well as summer droughts; however, its medium-early shooting makes it susceptible to late frosts. 'Vitouska' sometimes shows the phenomenon of gigantism, as clusters can reach 1 kg in some cases; the mean weight is around 250-300 g. With the aim of recovering and promoting the most worthy local grapevines, 'Vitouska' was registered in the Italian Catalogue in 1990 (code no. 320) and it belongs to the few varieties selected for Kars D.O.C. (Controlled Denomination Origin). Although opinion is widespread that 'Vitouska' originated in the Kars area, there is little information to support the assertion that this cultivar is strictly localized in the region and there are no traces of its presence elsewhere (DEL ZAN et al. 2004). The comparison of 11 SSR data, obtained for 'Vitouska' molecular identification, with the Istituto Sperimentale per la Viticoltura database suggested that this variety could be an offspring of 'Malvasia bianca lunga' alias 'Malvasia del Chianti' and 'Prosecco tondo'. By extending molecular analysis to 37 nuclear SSR loci, we provided useful support to the knowledge on the origins of 'Vitouska', since the additional data confirmed the two putative parents indicated above.

Material and Methods

Plant material: 'Malvasia bianca lunga', 'Prosecco tondo' and 'Vitouska' samples came from the Istituto Sperimentale per la Viticoltura collections at Conegliano (Italy). Two additional 'Vitouska' samples were kindly provided by Augusto Fabbro (ERSA of Gorizia).

Microsatellite analysis: The thirty nuclear SSR used in Crespan et al. (2006) were analysed, plus five additional ones from Vitis Microsatellite Consortium, UCH11 (Lefort et al. 2002) and scu05 (Scott et al. 2000). Multiplex PCR of two or three SSR loci were suitably arranged based on expected allele lengths. Electrophoresis was performed on denaturating polyacrylamide gel and silver staining for signal revelation; alleles of reference varieties were used for unknown allele sizing and manual scoring, as reported in Crespan and Milani (2001).

Statistical analysis: Molecular data significance was estimated with the Identity 1.0 program, freely available on http://www.boku.ac.at/zag/forsch/MANUAL. rtf using the molecular database of our centre.

Results and Discussion

The molecular profiles obtained for the three analysed varieties are reported in Tab. 1, along with a further three reference cultivars for easier data comparison. 'Vitouska' shows one allele derived from each of the presumed parents at all 37 nuclear SSR loci, strongly supporting the hypothesis that 'Vitouska' is truly an offspring of 'Prosecco tondo' and 'Malyasia bianca lunga'.

Cumulative likelihood ratios of 'Vitouska' being the progeny of 'Prosecco tondo' (1) and 'Malvasia bianca lunga' alias 'Malvasia del Chianti' (2), versus alternative parents, including close relatives, combined over 36 nuclear SSR loci are reported in Tab. 2. VVMD8 locus was excluded from computation, due to the known presence of null alleles. This estimate shows that the indicated parents are more highly probable than other presumable combinations (1.04 x 10¹⁵ with 95 % upper confidence limit).

'Prosecco tondo' takes its name from the village of Prosecco, in Trieste province, where it is also known as 'Glera'. The 'Glera's' are an ensemble of different ancient varieties cultivated in this province. Recent research has shown that the name Glera refers mostly to 'Prosecco lungo' and less frequently to 'Prosecco tondo' or other minor varieties (Crespan *et al.* 2007).

'Malvasia bianca lunga' is usually grown in the Tre Venezie area, even if under a different name from the most common one: for example it is known as 'Prosecco nos-

 $$\rm T\,a\,b\,l\,e\,\,\,1\,$$ SSR data of 'Vitouska' and its proposed parents, plus three reference varieties. Allele lengths are in bp

Variety	Vitouska	Prosecco tondo	Malvasia bianca lunga	Cabernet Sauvignon	Moscato bianco	Sultanina	Variety	Vitouska	Prosecco tondo	Malvasia bianca lunga	Cabernet Sauvignon	Moscato bianco	Sultanina
VVS1	181 181	181 181	180 181	181 181	181 181	181 188	VrZAG79	242 248	248 258	242 250	246 246	250 254	246 258
VVS2	133 145	133 143	145 145	139 151	133 133	145 151	VrZAG83	194 194	188 194	190 194	200 200	188 188	188 194
VVS29	171 171	171 171	171 171	179 181	171 171	171 179	ISV2 (VMC6e1)	151 165	141 151	143 165	141 165	141 143	143 143
VVMD5	226 240	226 246	226 240	232 240	228 236	234 234	ISV3 (VMC6f1)	133 133	133 139	133 139	133 139	133 139	133 139
VVMD7	239 247	239 247	239 253	239 239	233 249	239 253	ISV4 (VMC6g1)	177 197	169 197	177 177	169 191	169 187	191 193
VVMD8	147 147	143 147	147 157	143 157	141 141	145 157	VMC1e12	250 260	246 250	250 260	240 250	260 260	244 260
VVMD17	222 222	221 222	221 222	221 222	220 222	222 222	VMCNG4b9	166 176	166 176	150 176	168 176	158 166	138 158
VVMD21	243 249	243 249 214	249 249	249 258	249 266	249 256	VMC4g6	125 133	125 143	133 143	127 133	127 143	125 129
VVMD24	210 214 243	218 243	210 210 243	210 219 243	214 219 245	210 219 243	VMC2h9	117 117 161	117 123 161	117 117 163	119 123 159	121 123 161	123 123 161
VVMD25	245 249	247 249	245 249	253 249	253 251	253 249	VMC3d7	171 119	163 119	171 119	161 121	163 123	163 125
VVMD26	251 179	251 179	251 179	251 175	251 179	251 181	VMC2g2	119 198	125 206	119 198	125 212	125 200	125 204
VVMD27	194 239	194 239	179 251	189 237	194 249	194 221	VMC2h4	214 95	214 93	200 95	220 111	214 111	214 121
VVMD28 VVMD31	257 214	247 212	257 212	239 206	271 212	247 212	VMC6e10 VMC4h6	119 158	119 158	97 158	121 152	113 152	127 162
VVMD31	216 253	216 263	214 253	210 241	216 265	212 251	VMC4c6	158 160	158 160	158 160	162 163	162 157	164 151
VVMD36	263 254	265 252	257 254	241 254	273 244	251 250	UCH11	163 242	163 242	163 242	175 244	157 244	157 242
VrZAG21	254 190 206	254 190 200	254 204 206	264 200 206	264 206 206	268 190 202	SCU05	246 165 169	246 169 169	262 163 165	262 165 168	248 160 165	244 165 171
VrZAG62	195 203	187 203	195 199	187 193	185 195	187 187	VMC5g6.1	138 139	138 142	139 165	142 155	139 151	171 139 139
VrZAG64	159 163	143 163	137 159	139 159	141 159	143 159		137	112	100	100	101	137

Table 2

Cumulative likelihood ratios of 'Vitouska' being the progeny of 'Malvasia bianca lunga' *alias* 'Malvasia del Chianti' (1) and 'Prosecco tondo' (2), versus alternative parents, including close relatives, combined over 36 nuclear SSR loci. VVMD8 locus was excluded for null allele presence

Parents combinations	(1) x (2)	(1) x X	rel (2) x (1)	(2) x X	rel (1) x (2)
with observed allele frequencies	1.72 x 10 ²⁴	8.76 x 10 ¹³	7.16 x 10 ⁴	2.91 x 10 ¹⁵	1.31 x 10 ⁵
with 95 % upper confidence limit	4.51×10^{17}	6.13×10^{10}	8.58×10^3	1.57×10^{12}	1.74×10^4

trano of Conegliano' (CALÒ et al. 2000). We have no direct evidence of 'Malvasia bianca lunga' cultivation in Kars, but it is highly probable. The well-known and documented presence of at least one putative parent in the same cultivation region as 'Vitouska' is an important additional element in support of the indication came out from molecular data. It is therefore very likely that 'Vitouska' originated in the Kars area. A concise ampelographic description of 'Vitouska' is reported in DEL ZAN et al. (2004), with pictures of shoot, mature leaf and cluster. 'Vitouska' is quite distinct from 'Garganega' of the Veneto region. It is also morphologically different from another 'Gargania', cited and briefly described among the grapes grown in Friuli region in the first half of the 1800s, both in the "Catalogo delle varietà delle viti del Regno veneto" (Grape cultivar catalogue of Veneto Kingdom) written by P. DA MANIAGO in 1823 and in that of "Mostra delle uve dell'Associazione agraria friulana" (Grape exhibition of Friuli Agronomy Association) of 1863 (CALÒ and COSTACURTA 1991). This latter is described as a mediocre cultivar, for table and wine use, with very long cluster and highly sparse berries, yellow-red or yellow-pink coloured and with a thick skin.

Interest in 'Vitouska' has increased in recent years due to the ability and determination of some local viticulturists, who believe in this variety and are firmly resolved to recover it in culture, despite the difficulty linked to the need for careful canopy management to reach full berry ripening. 'Vitouska' grape vinification produces fresh, dry or slightly sweetish wines, characterized by flavours such vegetable, hay, sage, with William pear and lemon aromas. The wine has a pale yellow colour, with a slightly sour and sapid taste and medium structure. Vinification in blends could prove an alternative and interesting use: tests are under way using local white varieties, such as 'Malvasia istriana'.

Acknowledgements

This research was supported by Trattato Internazionale FAO, Progetto Risorse Genetiche Vegetali, funded by the Italian Ministero delle Politiche Agricole, Alimentari e Forestali.

References

- Calò, A.; Costacurta, A.; 1991: Delle Viti in Friuli. Ed. Arti Grafiche Friulane (Udine).
- Calò, A.; Costacurta, A.; Cancellier, S.; Crespan, M.; Milani, N.; Carraro, R.; Giust, M.; Sartori, E.; Anaclerio, F.; Forti, R.; Ciprian, L.; Di Stefano, R.; Pigella, R.; Bottero, S.; Gentillini, N.; 2000: Delle viti Prosecche. Ovvero della Distinzione fra Prosecco tondo e Prosecco lungo. Libra Ed.
- CRESPAN, M.; MILANI, N.; 2001: The Muscats: A molecular analysis of synonyms, homonyms and genetic relationships within a large family of grapevine cultivars. Vitis 40, 23-30.
- Crespan, M.; Cancellier, S.; Chies, R.; Giannettio, S.; Meneghetti, S.; 2006: The parents of Raboso veronese were discovered: a new hypothesis on its origin. Riv. Vit. Enol. 1, 3-12.
- Crespan, M.; Cancellier, S.; Chies, R.; Giannetto, S.; Meneghetti, S.; Costacurta, A.; 2007: Molecular contribution to the knowledge of two ancient varietal populations: Rabosi and Glere. Acta Hortic. (accepted for publication).
- Del Zan, F.; Failla, O.; Scienza, A.; 2004: La vite e l'Uomo. Dal Rompicapo delle Origini al Salvataggio delle Reliquie. Ed. ERSA Agenzia regionale per lo sviluppo rurale, Gorizia.
- LEFORT, F.; KYVELOS, C. J.; ZERVOU, M.; EDWARDS, K. J.; ROUBELAKIS-AN-GELAKIS, K. A.; 2002: Characterization of new microsatellite loci from *Vitis vinifera* and their conservation in some *Vitis* species and hydrids. Mol. Ecol. Notes 2, 20-21.
- Scott, K. D.; Eggler, P.; Seaton, G.; Rossetto, M.; Ablett, E. M.; Lee, L. S.; Henry, R. J.; 2000: Analysis of SSRs derived from grape ESTs. Theor. Appl. Genet. 100, 723-726.

Received April 26, 2007