# Genetic characterization and relationships of traditional grape cultivars from Transcaucasia and Anatolia 

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#### Abstract

We present here the first large-scale genetic characterization of grape cultivars from Transcaucasia and Anatolia. These regions where wild grapes still grow in nature have been cultivating wine and table grapes for thousands of years and are considered the cradles of viticulture. Using 12 nuclear microsatellite markers, we genotyped 116 accessions of traditional grape cultivars from Armenia, Georgia and Turkey and we detected 17 identical genotypes and six homonymy cases, mainly within each national germplasm. Neighbour-joining analysis of genetic distance showed that each germplasm could have multiple origins and although they are now separated, they might have some common ancestors. In addition, four varieties from Western Europe included as outgroups turned out to be more related to Georgian cultivars than other germplasms, suggesting a possible ancient origin in Georgia. This work represents a first step towards germplasm management of this rich ampelographic heritage.


Keywords: Armenia; genetic relationship; Georgia; microsatellite; Turkey; Vitis vinifera

## Introduction

Transcaucasia and Anatolia have long been regarded as likely homelands of viticulture and the earliest 'wine culture' (Vavilov, 1926; Negrul, 1938; Levadoux, 1956; Olmo, 1995; Zohary and Hopf, 2000; McGovern, 2003). The wild Vitis vinifera L. subsp. silvestris continues to thrive in these regions, where today hundreds of cultivars (V. vinifera L. subsp. vinifera) are grown for wine and table grapes. Based on recent archaeological and chemical

[^0]evidence, a 'wine culture' had been established as early as 6000 BC in the upland region of the Taurus Mountains in Eastern Anatolia, the Caucasus Mountains (including Transcaucasia) and the northern Zagros Mountains of Iran (see McGovern, 2003 for discussion and references). Recent chemical analyses (P. McGovern, in preparation) of Neolithic pottery from Georgia (Shulaveris-Gora) and Eastern Anatolia (Cayönü), dating back to the early 6th millennium BC , corroborate that the same beverage was being produced over a broad area of the mountainous Near East.
The present work will focus on DNA typing of grape cultivars from these areas. In Georgia, more than 500 indigenous wine and table grape cultivars have been described
(Ketskhoveli et al., 1960), including centuries-old cultivars like 'Rkatsiteli', 'Mtsvane' or 'Khikhvi' (Chkhartishvili and Tsertsvadze, 2003). However, only half of these cultivars have been conserved in four national collections and today only a small number of local varieties are still cultivated (Chkhartishvili, 2003; Maghradze, 2003). The most important autochthonous cultivars for winemaking are 'Rkatsiteli' (white) and 'Saperavi' (red). In Armenia, the Merdzavan ampelographic collection used to contain more than 800 accessions of indigenous and introduced varieties (including various clones) until 1993, but it has been unfortunately destroyed after land privatization (Gasparyan and Melyan, 2003). Less than 100 accessions are now available from three national collections, of which very few are autochthonous cultivars. The most important indigenous wine varieties are the white 'Voskeat' and 'Mskhali' (used for brandy) and the red 'Areni Chernyi' and 'Kachet'. In Azerbaijan, more than 500 grape cultivars are kept in collection, half of them being considered local varieties (Musayev, 2003). The most important indigenous varieties are the white 'Bajac Shirei' and the red 'Shahani'. In Turkey, more than 1000 grape accessions exist in the National Germplasm Repository Vineyard at Tekirdağ Viticulture Research Institute in Thrace (Ağaoğlu and Celik, 1986; Ergül et al., 2002), most of them being considered indigenous to Anatolia. The most important indigenous varieties are the white 'Sultani Çekirdeksiz' ('Sultanina' or 'Thompson Seedless', especially for table grape production), 'Emir', 'Narince' and 'Misket' and the red 'Öküzgözü' and 'Boğazkere'. The genetic relationships among and between these gene pools of grape cultivars were investigated here by DNA profiling.

Since their first application to grapevine (Thomas and Scott, 1993), microsatellites have been widely used for cultivar identification (Grando and Frisinghelli, 1998; Sefc et al., 1998a; Meredith et al., 1999) and analysis of genetic relationships (Lefort and Roubelakis-Angelakis, 2001; Aradhya et al., 2003). However, only a fraction of the 8000-10,000 grape cultivars existing worldwide (Alleweldt, 1997) have been genotyped with microsatellites, most of them coming from occidental Europe. The greater part of the huge germplasm of grape cultivars from the Near and Middle East remains to be genotyped. To our knowledge, there are no microsatellite data available in the literature for cultivars from Armenia or Georgia, with the exception of the widespread 'Rkatsiteli' and 'Saperavi' in Lamboy and Alpha (1998). From Turkey, Aradhya et al. (2003) analysed five cultivars and Benjak et al. (2005) genotyped nine cultivars, of which three are analysed in the present paper ('Erik Kara'/'Kara Erik', 'Hatun Parmağ'' and 'Kabarcık') with five loci overlapping. For each country, we selected the most ancient and traditional grape cultivars in order to obtain a representative sampling of the whole germplasm. We used 12
nuclear microsatellite markers to characterize 116 accessions of traditional grape cultivars from Armenia (13), Georgia (41) and Turkey (62) and searched for synonyms, homonyms and genetic relationships. This work represents the first microsatellite characterization of germplasms from Transcaucasia and Anatolia, and it is a first step towards germplasm management of this rich ampelographic heritage.

## Materials and methods

## Plant material

Genomic DNA was extracted with Qiagen DNeasy Plant Mini Kit or according to Lodhi et al. (1994) from dried leaves of 116 accessions sampled in vineyards or ampelographic collections and putatively corresponding to 98 grape cultivars (Table 1) from Georgia, Armenia and Turkey (Azerbaijan could unfortunately not be included in the sampling).

## Microsatellite analysis

We analysed 12 nuclear microsatellite markers: VVMD5 and 7 (Bowers et al., 1996), VVMD24, 28, 31 and 32 (Bowers et al., 1999b), VrZAG62 and 79 (Sefc et al., 1999), VVS2 (Thomas and Scott, 1993), and VMC2C3, 2H4 and 5A1 (Vitis Microsatellite Consortium, www.agrogene.com). Five of these markers belong to the 'core set' chosen by the international grape community (This et al., 2004): this will allow comparison of our data with most other germplasms. PCR amplifications were performed in $10 \mu \mathrm{l}$ reaction mixtures with Qiagen HotStarTaq Master Mix Kit. Microsatellite markers were labelled with three possible fluorescent dyes (6-FAM, HEX and NED, Applied Biosystems). PCR conditions were: 15 min at $95^{\circ} \mathrm{C}$ (initial activation step for HotStarTaq DNA Polymerase), 35 cycles comprised of 60 s at $94^{\circ} \mathrm{C}$ (denaturation), 30 s at 52 or $56^{\circ} \mathrm{C}$ according to literature (annealing), 90 s at $72^{\circ} \mathrm{C}$ (extension), followed by 10 min at $72^{\circ} \mathrm{C}$ (final extension). Every accession was amplified at least twice or more if necessary in order to avoid typing errors. PCR products were electrophoresed in an ABI PRISM 3100 DNA Sequencer (Applied Biosystems). Allele sizes were assigned against Genescan ROX 400 internal size standard and individuals were genotyped using Genescan analysis software and Genotyper software version 3.7 (Applied Biosystems). In each run, we have included four well-known cultivars from Western Europe ('Chasselas' and 'Pinot Noir' from Agroscope RAC Changins, Centre viticole du Caudoz, Pully,

Table 1. Cultivars included in this study: the 116 cultivars are grouped by country of origin and numbered alphabetically

| Name ${ }^{\text {a }}$ |  | Sampling ${ }^{\text {b }}$ | Cultivation area ${ }^{\text {c }}$ |
| :---: | :---: | :---: | :---: |
| Armenia |  |  |  |
| 1 | Ak-kaltak (b,wt) | YR | Uzbekistan |
| 2 | Areni Chernyi ( $\mathrm{n}, \mathrm{w}$ ) | N | Vajots Dzor (Armenia) |
| 3 | Areni Chernyi ( $\mathrm{n}, \mathrm{w}$ ) | Areni ${ }^{+}$ | Vajots Dzor (Armenia) |
| 4 | Chilar (b,w) | YR | Armenia |
| 5 | Garandmak (b,w) | YR | Armenia, Crimea and Caucasus |
| 6 | Kachet ( $\mathrm{n}, \mathrm{w}$ ) | Areni ${ }^{+}$ | Armenia |
| 7 | Karmir Kakhani (n,t) | YR | Armenia |
| 8 | Khatun Khardjzhi (b,w) | Areni ${ }^{+}$ | Armenia |
| 9 | Mskhali (b,wt) | YR | Armenia |
| 10 | Mskhali (b,wt) | Areni ${ }^{+}$ | Armenia |
| 11 | Tozot ( $\mathrm{n}, \mathrm{w}$ ) | Areni ${ }^{+}$ | Armenia |
| 12 | Vardagujn Jerevani (r,t) | YR | Erevan (Armenia) |
| 13 | Voskeat (b,w) | YR | Echmiadzin, Erevan (Armenia) |
| Georgia |  |  |  |
| 14 | Alabeuri Chavi* ( $\mathrm{n}, \mathrm{w}$ ) | Racha-Lechkumi ${ }^{+}$ | Racha-Lechkumi and Crimea |
| 15 | Aladasturi ( $\mathrm{n}, \mathrm{w}$ ) | TVI | Imereti and Crimea |
| 16 | Aleksandrouli ( $\mathrm{n}, \mathrm{w}$ ) | TVI | Georgia and Crimea |
| 17 | Buera (b,wt) | TVI | Georgia and former USSR |
| 18 | Chavkapito ( n , w) | Kharti ${ }^{\text {+ }}$ | Ossetia |
| 19 | Chinuri (b,w) | TVI | Georgia |
| 20 | Chitistvala Bodburi (b,w) | Khakheti ${ }^{+}$ | Khakheti |
| 21 | Chkhaveri ( $\mathrm{n}, \mathrm{w}$ ) | TVI | Mingrelia |
| 22 | Chkovra* (?, w) | Imereti ${ }^{\dagger}$ | Imereti |
| 23 | Dondglabi (b,w) | TVI | Imereti |
| 24 | Dondglabi Tetri* (n,w) | Imereti ${ }^{\dagger}$ | Imereti |
| 25 | Dzvelshavi Sachkheris (n,w) | TVI | Imereti and former USSR |
| 26 | Gorula (or Ananovra) (b,wt) | TVI | Khartli |
| 27 | Grdzelmtevana (b,w) | Gurdjani ${ }^{+}$ | Khakheti, Imereti and former USSR |
| 28 | Kapistoni Imeretinskii (b,w) | Imereti ${ }^{\text {+ }}$ | Imereti |
| 29 | Kharistvala Tetri (b,wt) | TVI | Georgia |
| 30 | Khikhvi (b,w) | TVI | Khakheti, Moldavia, Crimea and Daghestan |
| 31 | Khounalige* (b,w) | TVI | Abkhazia |
| 32 | Khupishizh ( $\mathrm{n}, \mathrm{w}$ ) | TVI | Abkhazia |
| 33 | Kichouri* (b,wt) | TVI | Khartli |
| 34 | Kisi (b,wt) | TVI | Khakheti |
| 35 | Krakhuna (b,wt) | TVI | Imereti and Crimea |
| 36 | Kundza (b,w) | TVI | Imereti, Khakheti |
| 37 | Kvira (r,w) | TVI | Racha-Lechkumi, Imereti |
| 38 | Maglari Tvrina (r,w) | TVI | Georgia |
| 39 | Mamukas Sapere ( $\mathrm{n}, \mathrm{w}$ ) | TVI | Georgia |
| 40 | Meskhuri Chitiskvertskha* (n,w) | Kharti ${ }^{+}$ | Khartli |
| 41 | Meskhuri Shavi (r,wt) | TVI | Khartli |
| 42 | Mudzhuretuli ( $\mathrm{n}, \mathrm{w}$ ) | TVI | Georgia |
| 43 | Odzhaleshi ( $\mathrm{n}, \mathrm{wt}$ ) | TVI | Samegrelo, Mingrelia, Imereti, Khakheti |
| 44 | Otskhanuri Sapere ( $\mathrm{n}, \mathrm{w}$ ) | Imereti ${ }^{\text {+ }}$ | Imereti |
| 45 | Rkatsiteli (b,wt) | TVI | Khakheti and former USSR |
| 46 | Saperavi Mrgvalmarzvala* ( $\mathrm{n}, \mathrm{w}$ ) | Khakheti ${ }^{\dagger}$ | Khakheti and former USSR |
| 47 | Saperavi Pachkha ( n , w) | Khakheti ${ }^{+}$ | Khakheti and former USSR |
| 48 | Skhilatubani ( $\mathrm{n}, \mathrm{w}$ ) | TVI | Georgia |
| 49 | Tavkara (n,w) | Khakheti ${ }^{+}$ | Khakheti |
| 50 | Tavkveri (n,wt) | TVI | Khakheti and Azerbaijan |
| 51 | Tkupkvirta ( n , w) | Khakheti ${ }^{+}$ | Khakheti, Imereti |
| 52 | Tsitska (b,wt) | Imereti ${ }^{+}$ | Imereti |
| 53 | Tsolikouri (b,w) | Imereti ${ }^{+}$ | Imereti |
| 54 | Uriatubanskii ( $\mathrm{n}, \mathrm{w}$ ) | Vardisubani ${ }^{+}$ | Russia |
| Turkey |  |  |  |
| 55 | Abderi* (b,tr) | TNGRV | Çermik/Diyarbakır (SE Anatolia) |
| 56 | Asseri (b,t) | TNGRV | Nevşehir (Cappadocia) |
| 57 | Azezi (b,wtr) | TNGRV | Adıyaman (SE Anatolia) |

Table 1. Continued

| Name ${ }^{\text {a }}$ |  | Sampling ${ }^{\text {b }}$ | Cultivation area ${ }^{\text {c }}$ |
| :---: | :---: | :---: | :---: |
| 58 | Ballıboz (b,t) | Adıyaman ${ }^{+}$ | Adıyaman (SE Anatolia) |
| 59 | Belelük* ( $\mathrm{n}, \mathrm{w}$ ) | Çüngüş/Diyarbakır ${ }^{+}$ | Çüngüş/Diyarbakır (SE Anatolia) |
| 60 | Besni (b,tr) | TNGRV | SE and E Anatolia |
| 61 | Besni (b,tr) | AUKVRES | SE and E Anatolia |
| 62 | Boğazkere (n,wt) | TNGRV | Elazığ, Malatya, Gaziantep (SE Anatolia) |
| 63 | Boğazkere ( n , w) | AUKVRES | Elazığ, Malatya, Gaziantep (SE Anatolia) |
| 64 | Burdur Dimriti (n,wt) | TNGRV | Eğridir/Isparta (Mediterranean) |
| 65 | Burdur Dimriti (b,wr) | TNGRV | Eğridir/Isparta (Mediterranean) |
| 66 | Dımışkı (b,t) | TNGRV | SE Anatolia |
| 67 | Dökülgen (b,wt) | Kahramanmaraş ${ }^{+}$ | Gaziantep, Kahramanmaraş (SE Anatolia) |
| 68 | Dökülgen (b,w) | Konya ${ }^{+}$ | Hadım/Konya (C Anatolia) |
| 69 | Dökülgen (b,w) | AUKVRES | E Turkey, Kalecik |
| 70 | Ekşi Kara (n,wtr) | TNGRV | Narlıdere/Karaman (C Anatolia) |
| 71 | Ekşi Kara ( n , wtr) | TNGRV | Morcalı/Karaman (C Anatolia) |
| 72 | Emir (b,wt) | Nevşehir ${ }^{\dagger}$ | Nevşehir, Kayseri (C Anatolia) |
| 73 | Emir (b,wt) | AUKVRES | Cappadocia (C Anatolia) |
| 74 | Erik Kara (n,wt) | TNGRV | Sivas, Erzincan, Malatya (E Anatolia) |
| 75 | Gemre Siyah (n,t) | Isparta ${ }^{+}$ | C/Isparta (Mediterranean) |
| 76 | Gemre Siyah ( $\mathrm{n}, \mathrm{t}$ ) | Isparta ${ }^{+}$ | C/Isparta (Mediterranean) |
| 77 | Gemre Siyah ( $\mathrm{n}, \mathrm{t}$ ) | Isparta ${ }^{+}$ | C/Isparta (Mediterranean) |
| 78 | Gök Üzüm (b,wt) | Morcalı ${ }^{+}$ | Morcalı/Karaman (C Anatolia) |
| 79 | Hasandede Beyazı (b,wt) | AUKVRES | Ankara, Kırıkkale |
| 80 | Hasipali* (b,t) | Konya ${ }^{+}$ | Çatalhöyük/Konya (C Anatolia) |
| 81 | Hatun Parmagı** n , t$)$ | Çermik ${ }^{+}$ | Çermik/Diyarbakır (SE Anatolia) |
| 82 | Hönüsü (n,t) | AUKVRES | SE Anatolia |
| 83 | Hönüsü ( $\mathrm{n}, \mathrm{w}$ ) | TNGRV | SE Anatolia |
| 84 | İri Beyaz (b,t) | Çatalhöyük ${ }^{\dagger}$ | Çatalhöyük/ Konya (C Anatolia) |
| 85 | İri Daneli Ak Üzüm (b,t) | Bozkır ${ }^{+}$ | Bozkır/Konya (C Anatolia) |
| 86 | İri Kara (r,wtr) | Hadim ${ }^{+}$ | Hadım/Konya (C Anatolia) |
| 87 | İri Siyah* (n,t) | Çatalhöyük ${ }^{+}$ | Çatalhöyük/Konya (C Anatolia) |
| 88 | Kabarcık (b,wt) | TNGRV | E Turkey, Kalecik |
| 89 | Kabarcık (b,wt) | Kahramanmaraş ${ }^{+}$ | Kahramanmaraş, Gaziantep, Malatya |
| 90 | Kayseri Karası (n,w) | Kayseri ${ }^{+}$ | Kayseri (C Anatolia) |
| 91 | Kirkit (b,t) | Diyarbakır ${ }^{+}$ | Diyarbakır (SE Anatolia) |
| 92 | Kırmızı Dimrit (n,w) | Nevşehir ${ }^{+}$ | Nevşehir (Cappodochia) |
| 93 | Kızıl Üzüm (r,t) | TNGRV | Çatalhöyük/Konya (C Anatolia) |
| 94 | Künefi (n,t) | Gaziantep ${ }^{+}$ | Gaziantep (SE Anatolia) |
| 95 | Luvanek* (b,tr) | Çermik ${ }^{+}$ | Çermik/Diyarbakır (SE Anatolia) |
| 96 | Mor Üzüm* (b,t) | Uçhisar ${ }^{+}$ | Uçhisar/Nevşehir (C Anatolia) |
| 97 | Morek* (n,t) | Ergani ${ }^{\text {r }}$ | Ergani-Diyarbakır (SE Anatolia) |
| 98 | Muhammediye (b,wt) | TNGRV | Mardin (SE Anatolia) |
| 99 | Narince (b,w) | AUKVRES | C Turkey |
| 100 | Öküzgözü (n,w) | Elazığ ${ }^{+}$ | Elazığ, Malatya, Gaziantep (SE Anatolia) |
| 101 | Öküzgözü (n,w) | AUKVRES | E Turkey |
| 102 | Parmak (b,t) | Nevşehir ${ }^{+}$ | Nevşehir, Kayseri, Konya (C Anatolia) |
| 103 | Razakı (b,t) | Isparta ${ }^{+}$ | C/Isparta (Mediterranean) |
| 104 | Şaraplık Siyah (n,w) | Çermik ${ }^{+}$ | Çermik/Diyarbakır (SE Anatolia) |
| 105 | Şıralık (b,w) | Siverek ${ }^{+}$ | Siverek/Urfa (SE Anatolia) |
| 106 | Şıralık (b,w) | Siverek ${ }^{+}$ | Siverek/Urfa (SE Anatolia) |
| 107 | Şıralık (b,w) | Siverek ${ }^{+}$ | Siverek/Urfa (SE Anatolia) |
| 108 | Şıralık (b,w) | Siverek ${ }^{+}$ | Siverek/Urfa (SE Anatolia) |
| 109 | Şıralık (b,w) | Eğil ${ }^{+}$ | Eğil/ Diyarbakır (SE Anatolia) |
| 110 | Siyah* (n,w) | Yalvaç ${ }^{+}$ | Yalvaç/lsparta (Mediterranean) |
| 111 | Sungurlu (b,w) | AUKVRES | Kırıkkale, Çorum (C Anatolia) |
| 112 | Tahannebi (b,t) | AUKVRES | E Turkey |
| 113 | Tahannabi (b,t) | TNGRV + | SE Anatolia |
| 114 | Timbo* (n,wt) | Adiyaman ${ }^{+}$ | Adıyaman (SE Anatolia) |
| 115 | Vanki (b,wt) | Çermik ${ }^{+}$ | Çermik-Diyarbakır (SE Anatolia) |
| 116 | Vilki* (n,t) | Çermik ${ }^{+}$ | Çermik/Diyarbakır (SE Anatolia) |

Switzerland and 'Syrah' and 'Nebbiolo' from Istituto Agrario di San Michele all'Adige, Italy). They served as standards in order to have consistent allele sizes over all runs and they allowed allele size comparison with other germplasms.

## Identical genotypes and homonyms

The genotypes were compared to those of more than 1700 grape cultivars from all over the world put together and standardized from different databases (University of California, Davis; Grape Microsatellite Collection, IASMA, Italy, http://www.ismaa.it/areabioav/gmc.html; Greek Vitis Database, University of Crete, Heraklion, Greece, http://www.biology.uoc.gr/gvd/; Bulgarian Grape nSSR Database, http://bulgenom.abi.bg/ Grape\%20nSSR\%20Database.html) and from various references in the literature. We checked for the presence of identical genotypes within the accessions with the program 'DNA-Data' (B. H. Prins, unpublished). This program offers the option of a user-defined level of discrepancy, in order to ascertain possible identities despite the presence of a few allelic mismatches. This is particularly useful with mutations or null alleles.

## Genetic analysis

Standard genetic parameters were calculated using Microsat (Minch et al., 1995). Probability of identity was estimated with the program Identity (Wagner and Sefc, 1999). Populations (version 1.2.28) (Langella, 2002) was used to calculate Nei et al.'s (1983) $D_{A}$ pairwise genetic distance between individuals and construct the neigh-bour-joining tree of individuals presented in Fig. 1 and displayed with Treeview (Page, 1996). According to Takezaki and Nei (1996), the $D_{A}$ genetic distance is more efficient than Nei's (1972) standard genetic distance $\left(D_{s}\right)$, Nei's (1973) minimum genetic distance ( $D_{m}$ ) and Rogers' (1972) distance ( $D_{r}$ ) in obtaining the correct
topology, either under the infinite-allele model (IAM) or the stepwise mutation model (SMM) of microsatellites evolution. The program Populations accepts input file in Genepop (Raymond and Rousset, 1995) format. Hence we used the software Genetix (Belkhir et al., 19962002) to convert microsatellite allelic data computed in Microsoft Excel into Genepop format.

## Results and discussion

The analysis of 116 accessions from Armenia (13), Georgia (41) and Turkey (62) at 12 microsatellite markers (Table 2) generated 138 alleles. The number of alleles per locus ranged from six (VVMD24) to 16 (VVMD28 and 32) with a mean number of 11.9 (Table 3), which is much higher than values previously reported for the same loci (Lopes et al., 1999; Maletic et al., 1999; Lefort and Roubelakis-Angelakis, 2001; Costantini et al., 2005). Average observed heterozygosity was high at 0.796, slightly above Aradhya et al. (2003) who obtained 0.771 with 244 Vitis vinifera accessions analysed at eight microsatellite loci (of which five were analysed in the present work) and Sefc et al. (2000) who obtained 0.785 with 164 cultivars analysed at nine SSR markers (of which five were analysed in the present work). Such a high rate of heterozygosity is commonly observed among clonally propagated, outbreeding, perennial species (Aradhya et al., 2003). Total probability of identity (PI) was very low at $1.67 \mathrm{e}-12$. The most informative locus turned out to be VMC2H4 ( 12 alleles, PI $=0.054$ ) and the least informative was VVMD24 (seven alleles, PI $=0.231$ ). A few accessions exhibited three alleles at some loci (especially at VVS2 locus; see Table 2). Since one of the alleles usually had a weaker amplification signal, we suggest the presence of chimeric alleles at these loci. Chimerism has already been reported in grape cultivars (Franks et al., 2002; Riaz et al., 2002), especially for ancient cultivars (e.g. 'Pinot'). This suggests that cultivars showing chimeras in Table 2 ('Kachet' and 'Tozot' from Armenia; 'Dzvelshavi Sachkheris' from Georgia; 'Dımışkı',

[^1]
'Luvanek', 'Morek', 'Sungurlu' and 'Vilki' from Turkey) might be very old.

## Identical genotypes and homonyms

The Anatolian and Transcaucasian genotypes were checked against a database of more than 1700 cultivars (mostly from Western Europe) and wild grapes (from Western Europe and Near East) at 6-12 microsatellite markers (depending on the data available). We found 17 cases of identical genotypes (Table 4) and six cases of homonymy (Table 5). Referring to Table 1, several identical genotypes occurred between accessions of different colours and/or different uses (Table 4, pairs in italic). In such cases, it has not been possible to determine whether the cause of identical genotypes was synonymy, misnaming or mutations, because this would require detailed ampelographic analysis with independent accessions when possible. Three cases of identical genotypes were found outside the study area:

- 'İri Daneli Ak Üzüm' and 'Italia': 'Italia' is an artificial cross between 'Bicane' and 'Muscat de Hambourg' obtained in 1911 (Galet, 2000). 'İri Daneli Ak Üzüm’ literally means 'white grape with large berries'. It matched the genotype of 'Italia' from (i) IASMA collection at all loci; (ii) Crespan et al. (1999) at the six shared loci; (iii) Montpellier (Vassal, France) collection at the five shared loci; and (iv) Sanchez-Escribano et al. (1999) at the three shared loci. In addition, this genotype shares at least one allele at each available locus with 'Muscat de Hambourg' accessions from various sources (e.g. IASMA; University of California, Davis; Crespan, 2003). As a result, this genotype is most likely the true-to-type 'Italia'. Therefore, 'Italia' must have been introduced later to Turkey where it was given the name 'İri Daneli Ak Üzüm' after its berry colour and size.
- 'Parmak' and 'Jerusalem Bleu': 'Jerusalem Bleu' is a black-berried grape kept in a collection at Montpellier (Vassal, France) and supposedly introduced from Germany, but its origin is unknown (Galet, 2000). 'Parmak' is a white-berried grape cultivated in Central Anatolia. Careful ampelographic analysis would be
required to determine whether mislabelling or colour mutation could explain their having identical genotypes.
- 'Mor Üzüm' and 'Tsaousi': according to Galet (2000), the Greek 'Tsaousi' is the same as 'Chaouch', a white table grape very widespread in the Near East. Indeed, we observed that the genotype of 'Tsaousi' in Lefort and Roubelakis-Angelakis' (2001) is identical to the genotype of 'Chaouchi Politico' in Bowers et al. (1996). Thus, 'Mor Üzüm', 'Tsaousi' and 'Chaouch' are synonyms, which is consistent with the suggestion of Lefort and Roubelakis-Angelakis (2001) that 'Tsaousi' might be of Eastern origin.

The other identical genotypes occurred within the studied areas: the genotype of 'Saperavi Mrgvalmarzvala' and 'Saperavi Pachkha' matched the 'Saperavi' accessions in Lamboy and Alpha (1998) and in the collection of the University of California at Davis. As a result, and although Galet (2000) considers them as distinct cultivars, our data provided evidence that 'Saperavi Mrgvalmarzvala' and 'Saperavi Pachkha' are clones of the same cultivar.

For most homonymy cases in Table 5, it has been for now impossible to determine which cultivar is true-totype, because referring to ampelographic descriptions would be required, which is outside of the scope of this paper. The genotype of 'Areni Chernyi' (no. 2) sampled from a grape collection was different from the genotype of 'Areni Chernyi' (no. 3) that was sampled directly from the vineyards producing the nationally famous Areni wines; for that reason, we suggest that accession 3 is more likely the true-to-type 'Areni Chernyi', but analysing more independent samples and ampelographic descriptions would clear up this dilemma completely. Similarly, it has been impossible to determine the true-to-type accession of 'Burdur Dimriti' (Table 5). Since 'Dökülgen' 68 and 'Ekşi kara' 70 (Table 4) have the same genotype but different berry colours, 'Dökülgen' 68 is likely to be a misnaming and accessions 67 and 69 should be true-to-type 'Dökülgen'. However, because of identical genotypes with different names, we could not determine the true-to-type 'Ekşi kara' (see Genetic relationships). This would require careful study of ampelographic descriptions, etymology and local literature, which is out of the scope of the present

Fig. 1. Neighbour-joining tree of 89 distinct grape cultivars (identical genotypes were merged) from Anatolia and Transcaucasia and four Western European standards ('Chasselas', 'Nebbiolo', 'Pinot Noir' and 'Syrah') constructed from 12 microsatellite markers with Nei's $D_{A}$ genetic distance. Every cultivar is shown with its country of origin (A, Armenia; G, Georgia; T, Turkey) and its accession number in Table 1. Five distinct clusters were isolated, and the main cluster (no. 2) was subdivided (see text for details). Germplasms from each country are well separated and might have multiple origins, although all three germplasms are likely to have common ancestors. The Western European standards turned out to be closer to Georgian cultivars than Turkish or Armenian, suggesting they could have some Georgian ancestors.

| Cultivar | VVMD5 | VVMD7 | VVMD24 | VVMD28 | VVMD31 | VVMD32 | VVS2 | VrZAG62 | VrZAG79 | VMC2C3 | VMC2H4 | VMC5A1 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Ak-kaltak (1) | 240-234 | 253-243 | 223-214 | 249-235 | 212-196 | 257-251 | 157-151 | 189-189 | 257-247 | 170-162 | 218-208 | 171-161 |
| Areni Chernyi (2) | 236-236 | 245-235 | 219-210 | 261-237 | 222-210 | 253-249 | 143-135 | 195-189 | 249-237 | 170-165 | 212-212 | 169-165 |
| Areni Chernyi (3) | 240-236 | 249-249 | 218-210 | 247-247 | 216-210 | 273-273 | 133-133 | 201-199 | 247-243 | 179-170 | 222-218 | 167-165 |
| Chilar (4) | 236-236 | 253-249 | 218-214 | 279-279 | 212-196 | 273-267 | 135-133 | 205-197 | 251-241 | 179-170 | 222-210 | 167-165 |
| Garandmak (5) | 236-234 | 249-247 | 212-210 | 281-279 | 212-212 | 267-265 | 143-125 | 195-191 | 251-239 | 170-165 | 210-206 | 167-161 |
| Kachet (6) | 240-238 | 253-239 | 219-210 | 239-239 | 216-212 | 273-251 | 157-143-135 | 203-201 | 249-237 | 195-165 | 212-210 | 171-161 |
| Karmir Kakhani (7) | 240-240 | 253-243 | 210-210 | 247-241 | 216-214 | 273-251 | 151-135 | 197-189 | 257-247 | 170-165 | 220-218 | 171-171 |
| Khatun <br> Khardjzhi (8) | 246-240 | 249-239 | 218-218 | 247-247 | 216-210 | 273-273 | 133-125 | 197-189 | 247-239 | 179-170 | 218-208 | 171-165 |
| Mskhali (9)/ Mskhali (10) | 238-238 | 249-247 | 214-210 | 251-239 | 210-210 | 273-257 | 151-135 | 201-189 | 247-247 | 165-165 | 208-204 | 171-171 |
| Tozot (11) | 234-234 | 255-239 | 218-218 | 273-243 | 210-202 | 273-263 | 157-145-125 | 205-189 | 247-239 | 169-165 | 208-206 | 171-169 |
| Vardagujn Jerevani (12) | 234-228 | 247-239 | 219-208 | 279-219 | 224-212 | 273-251 | 151-143 | 205-189 | 259-251 | 170-162 | 208-206 | 171-161 |
| Voskeat (13) | 236-236 | 249-247 | 218-210 | 261-261 | 212-212 | 275-267 | 145-143 | 195-191 | 249-239 | 179-165 | 210-204 | 171-167 |
| Alabeuri Chavi (14) | 228-226 | 247-245 | 218-210 | 261-251 | 212-210 | 263-251 | 153-143 | 205-203 | 251-251 | 165-165 | 218-206 | 171-161 |
| Aladasturi (15) | 232-232 | 249-233 | 210-210 | 237-237 | 214-214 | 257-247 | 145-133 | 205-201 | 251-247 | 165-165 | 212-208 | 169-169 |
| Aleksandrouli (16) | 240-226 | 263-239 | 210-210 | 239-231 | 216-206 | 259-241 | 147-139 | 205-195 | 259-247 | 198-179 | 218-216 | 169-169 |
| Buera (17) | 238-236 | 259-247 | 210-210 | 247-247 | 212-196 | 251-247 | 145-137 | 197-191 | 251-251 | 170-165 | 218-210 | 169-165 |
| Chavkapito (18) | 240-228 | 247-239 | 210-208 | 261-237 | 210-196 | 273-251 | 145-135 | 201-191 | 251-251 | 195-165 | 218-212 | 171-161 |
| Chitistvala Bodburi (20) | 240-236 | 249-249 | 219-210 | 261-247 | 214-212 | 263-253 | 145-143 | 205-201 | 251-251 | 170-170 | 218-210 | 171-171 |
| Chkhaveri (21) | 240-232 | 263-247 | 210-210 | 237-229 | 216-214 | 263-251 | 133-133 | 205-203 | 253-239 | 200-165 | 208-206 | 169-165 |
| Chkovra (22)/ <br> Krakhuna (35)/ <br> Dondglabi <br> Tetri (24) | 234-232 | 253-239 | 214-210 | 239-239 | 216-212 | 253-251 | 145-133 | 197-195 | 253-239 | 195-165 | 212-208 | 169-169 |
| Dondglabi (23)/ <br> Kapistoni Im. <br> (28)/Gorula <br> (26)/Tavkara (49) | 236-234 | 253-247 | 210-210 | 247-237 | 216-196 | 273-251 | 143-141 | 201-191 | 251-237 | 165-165 | 210-210 | 175-169 |
| Dzvelshavi Sachkheris (25) | 240-226 | 251-247-239 | 214-210 | 239-239 | 214-212 | 273-251 | 145-143 | 201-195 | 261-255 | 192-165 | 208-204 | 171-171 |
| Grdzelmtevana (27) | 234-228 | 259-249 | 219-210 | 243-239 | 212-212 | 251-245 | 145-135 | 203-197 | 259-255 | 195-170 | 218-206 | 161-161 |
| Kharistvala Tetri (29) | 240-238 | 253-249 | 210-210 | 277-261 | 216-210 | 273-253 | 141-137 | 201-201 | 251-245 | 165-165 | 216-210 | 171-165 |
| Khikhvi (30) | 240-236 | 259-253 | 218-210 | 237-237 | 212-210 | 263-261 | 141-135 | 201-197 | 249-237 | 165-165 | 218-210 | 169-161 |
| Khounalige (31)/ Chinuri (19) | 240-238 | 253-249 | 210-210 | 261-239 | 216-210 | 273-263 | 141-137 | 201-201 | 251-249 | 170-156 | 224-210 | 165-161 |
| Khupishizh (32) | 238-232 | 251-249 | 210-210 | 237-237 | 212-210 | 263-253 | 155-151 | 201-195 | 255-237 | 170-170 | 222-200 | 167-161 |
| Kichouri (33) | 236-234 | 247-239 | 219-210 | 261-239 | 212-196 | 273-271 | 145-141 | 201-191 | 251-237 | 170-170 | 210-204 | 175-161 |
| Kisi (34) | 240-234 | 253-247 | 219-210 | 239-239 | 216-210 | 263-261 | 143-141 | 201-191 | 251-237 | 156-156 | 224-208 | 169-161 |
| Kundza (36) | 232-226 | 253-253 | 214-210 | 261-239 | 216-212 | 273-263 | 155-143 | 201-195 | 251-239 | 195-170 | 218-212 | 171-169 |
| Kvira (37) | 240-240 | 253-247 | 210-210 | 239-237 | 216-212 | 263-251 | 143-141 | 205-201 | 243-235 | 195-165 | 224-210 | 161-161 |

Table 2. Continued

| Cultivar | VVMD5 | VVMD7 | VVMD24 | VVMD28 | VVMD31 | VVMD32 | VVS2 | VrZAG62 | VrZAG79 | VMC2C3 | $\mathrm{VMC2H} 4$ | VMC5A1 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Maglari Tvrina (38) | 232-228 | 249-239 | 214-210 | 261-239 | 216-212 | 273-241 | 143-137 | 197-195 | 251-251 | 170-165 | 224-210 | 169-161 |
| Mamukas Sapere (39) | 240-232 | 253-233 | 210-210 | 239-237 | 216-214 | 263-255 | 151-141 | 205-201 | 259-239 | 179-165 | 208-204 | 165-165 |
| Meskhuri Chitiskvertskha (40) | 246-234 | 253-249 | 210-208 | 261-237 | 212-210 | 259-251 | 141-133 | 201-201 | 251-237 | 170-156 | 224-202 | 171-169 |
| Meskhuri Shavi (41) | 240-240 | 253-247 | 218-210 | 251-239 | 216-216 | 263-251 | 145-141 | 205-201 | 249-243 | 165-165 | 224-210 | 169-161 |
| Mudzhuretuli (42) | 228-226 | 245-239 | 218-210 | 261-251 | 212-210 | 263-251 | 155-143 | 205-205 | 251-245 | 165-165 | 218-206 | 171-161 |
| Odzhaleshi (43) | 240-234 | 247-239 | 214-210 | 251-243 | 216-212 | 273-263 | 145-143 | 205-189 | 257-251 | 165-165 | 210-210 | 171-165 |
| Otskhanuri Sapere (44) | 232-228 | 263-247 | 214-210 | 261-239 | 216-212 | 273-241 | 139-133 | 205-195 | 247-245 | 170-165 | 224-210 | 169-161 |
| Rkatsiteli (45) | 240-234 | 253-247 | 210-210 | 247-239 | 210-210 | 273-263 | 141-133 | 201-191 | 259-249 | 170-165 | 210-210 | 165-161 |
| Saperavi Pachkha <br> (47)/Saperavi <br> Mrgvalmarzvala (46) | 240-224 | 239-239 | 219-214 | 247-237 | 212-212 | 251-245 | 145-133 | 201-189 | 259-243 | 165-165 | 224-208 | 171-167 |
| Skhilatubani (48) | 232-232 | 249-241 | 210-210 | 237-237 | 216-212 | 263-255 | 151-151 | 201-195 | 251-239 | 198-165 | 218-208 | 169-169 |
| Tavkveri (50) | 238-228 | 243-239 | 218-216 | 239-221 | 216-216 | 273-241 | 151-137 | 195-189 | 245-239 | 198-170 | 204-204 | 167-157 |
| Tkupkvirta (51) | 240-238 | 253-247 | 210-210 | 239-239 | 216-216 | 259-251 | 143-141 | 205-201 | 257-237 | 165-156 | 212-212 | 171-165 |
| Tsitska (52) | 234-226 | 253-239 | 210-210 | 261-239 | 216-212 | 273-263 | 145-143 | 197-195 | 251-251 | 179-165 | 212-212 | 171-169 |
| Tsolikouri (53) | 232-226 | 253-247 | 214-210 | 239-237 | 216-216 | 263-249 | 145-143 | 209-195 | 239-239 | 195-165 | 212-208 | 171-169 |
| Uriatubanskii (54) | 240-236 | 247-247 | 210-210 | 239-237 | 216-216 | 273-263 | 145-137 | 205-191 | 237-237 | 195-165 | 210-204 | 175-171 |
| Aşeri (56)/ Hasandede Beyazı (79) | 240-236 | 253-249 | 210-210 | 261-239 | 212-212 | 257-257 | 149-143 | 189-189 | 247-247 | 170-165 | 224-218 | 171-165 |
| Azezi (57) | 234-226 | 249-249 | 214-210 | 239-239 | 214-210 | 273-253 | 135-133 | 205-193 | 257-251 | 170-170 | 210-202 | 173-171 |
| Ballıboz (58) | 246-234 | 253-251 | 219-214 | 261-261 | 212-196 | 273-263 | 143-139 | 205-205 | 247-243 | 165-165 | 218-210 | 171-171 |
| Belelük (59) | 232-232 | 249-249 | 214-210 | 247-237 | 212-204 | 259-241 | 155-141 | 195-195 | 249-249 | 170-165 | 218-204 | 171-165 |
| Besni (60)/ <br> Besni (61) | 238-232 | 247-247 | 219-210 | 237-237 | 216-210 | 273-263 | 151-141 | 205-193 | 257-257 | 179-165 | 210-204 | 171-157 |
| Boğazkere (62 and 63)/ Şaraplık Siyah (104) | 238-232 | 255-249 | 210-210 | 247-237 | 216-212 | 273-259 | 155-151 | 205-195 | 249-247 | 165-165 | 210-204 | 171-167 |
| Burdur Dimriti (64) | 236-236 | 249-247 | 210-210 | 247-237 | 216-210 | 257-257 | 143-137 | 205-201 | 255-247 | 195-170 | 222-218 | 171-169 |
| Burdur Dimriti (65) | 234-234 | 255-239 | 219-210 | 247-237 | 212-212 | 251-251 | 151-143 | 201-189 | 259-247 | 165-162 | 218-208 | 171-171 |
| Dımışkı (66) | 240-236-226 | 249-239 | 214-212 | 261-247 | 214-214 | 257-253 | 149-133 | 205-189 | 257-257 | 179-170 | 218-202 | 171-171 |
| Dökülgen (67 and 69) | 246-226 | 247-247 | 214-212 | 261-247 | 216-210 | 273-253 | 143-135 | 205-193 | 251-247 | 165-165 | 220-204 | 171-169 |
| Ekşi Kara (70)/ <br> Dökülgen (68) | 236-228 | 249-239 | 210-208 | 261-251 | 212-212 | 273-253 | 139-137 | 189-189 | 251-247 | 170-165 | 222-202 | 165-161 |
| Ekşi Kara (71) | 240-236 | 249-243 | 219-210 | 261-237 | 212-210 | 273-273 | 143-137 | 201-189 | 251-247 | 170-165 | 222-202 | 169-165 |
| Emir (72)/Emir (73) | 246-236 | 249-247 | 210-208 | 251-243 | 214-196 | 263-251 | 133-133 | 205-197 | 249-243 | 170-165 | 218-208 | 171-167 |
| Gemre Siyah (75)/ Gemre Siyah (76) | 240-236 | 249-247 | 212-210 | 247-247 | 216-196 | 253-253 | 143-137 | 205-189 | 255-255 | 195-170 | 220-218 | 171-165 |
| Gemre Siyah (77) | 236-236 | 253-249 | 219-210 | 247-237 | 210-210 | 251-249 | 133-131 | 201-189 | 259-251 | 170-165 | 220-204 | 171-165 |
| Gök Üzüm (78) | 238-234 | 251-247 | 219-210 | 247-247 | 224-196 | 263-257 | 143-143 | 205-205 | 247-245 | 170-170 | 218-202 | 173-171 |

Table 2. Continued

| Cultivar | VVMD5 | VVMD7 | VVMD24 | VVMD28 | VVMD31 | VVMD32 | VVS2 | VrZAG62 | VrZAG79 | VMC2C3 | VMC2H4 | VMC5A1 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Hasipali (80) | 240-226 | 253-239 | 218-210 | 261-251 | 212-212 | 273-259 | 145-143 | 201-189 | 259-243 | 165-165 | 218-204 | 171-171 |
| Hatun Parmağı (81) | 246-236 | 247-239 | 218-210 | 237-237 | 210-196 | 273-257 | 145-141 | 205-189 | 247-247 | 179-170 | 222-206 | 171-169 |
| Hönüsü (82)/ Hönüsü (83) | 232-228 | 253-239 | 219-210 | 247-247 | 210-204 | 275-251 | 149-143 | 189-189 | 259-257 | 195-179 | 208-202 | 165-161 |
| İri Daneli Ak <br> Üzüm (85) | 238-232 | 247-243 | 214-210 | 247-237 | 214-212 | 273-253 | 149-133 | 205-193 | 257-255 | m.d. | m.d. | m.d. |
| İri Kara (86) | 240-238 | 263-243 | 219-210 | 261-237 | 212-210 | 273-241 | 143-137 | 205-201 | 247-243 | 198-165 | 218-202 | 169-169 |
| İri Siyah (87) | 236-236 | 249-247 | 210-210 | 261-237 | 216-212 | 275-275 | 143-143 | 205-189 | 251-247 | 198-170 | 218-202 | 171-165 |
| Kayseri Karası <br> (90)/İri Beyaz (84) | 240-236 | 247-239 | 218-208 | 251-247 | 214-212 | 273-259 | 145-137 | 205-189 | 259-251 | 165-165 | 218-204 | 171-171 |
| Kirkit (91) | 236-226 | 247-239 | 219-210 | 261-261 | 210-210 | 257-253 | 143-143 | 193-189 | 251-247 | 170-165 | 212-202 | 175-171 |
| Kırmızı Dimrit (92) | 240-236 | 253-249 | 218-210 | 261-261 | 212-210 | 273-259 | 145-137 | 201-201 | 259-243 | 198-165 | 224-204 | 171-171 |
| Kızıl Üzüm (93) | 236-236 | 249-243 | 210-210 | 261-261 | 216-210 | 257-257 | 143-133 | 205-201 | 251-251 | 170-165 | 218-202 | 167-165 |
| Künefi (94) | 236-234 | 247-243 | 214-210 | 247-237 | 216-212 | 263-241 | 135-133 | 193-189 | 249-249 | 170-165 | 216-204 | 171-171 |
| Luvanek (95) | 238-236 | 249-247 | 212-210 | 261-261 | 210-202 | 273-263 | 155-141-133 | 195-191 | 257-249 | 192-165 | 210-210 | 171-165 |
| Mor Üzüm (96) | 238-228 | 249-247 | 219-208 | 261-261 | 224-210 | 273-253 | 151-135 | 205-189 | 249-247 | 179-179 | 216-206 | 171-161 |
| Morek (97) | 238-232 | 255-247 | 210-210 | 247-247 | 216-212 | 259-259 | 155-151-135 | 205-195 | 249-247 | 165-165 | 210-204 | 171-171 |
| Muhammediye (98)/Kabarcık (88 and 89) | 234-234 | 253-247 | 219-210 | 261-261 | 212-210 | 263-251 | 143-133 | 205-193 | 247-247 | 165-165 | 218-206 | 173-171 |
| Narince (99) | 240-236 | 249-243 | 210-210 | 261-237 | 212-212 | 273-253 | 145-143 | 201-189 | 251-251 | 165-165 | 224-224 | 171-165 |
| Öküzgözü (100)/ Öküzgözü (101)/ Erik Kara (74) | 234-234 | 249-247 | 219-219 | 247-237 | 216-216 | 257-241 | 151-143 | 205-205 | 257-251 | 192-179 | 210-204 | 171-169 |
| Parmak (102) | 236-236 | 249-247 | 210-210 | 247-237 | 212-210 | 275-257 | 143-137 | 201-189 | 251-247 | 198-198 | 224-218 | 173-171 |
| Razakı (103) | 236-236 | 249-247 | 210-210 | 247-247 | 212-196 | 275-253 | 137-137 | 189-189 | 255-251 | 195-195 | 224-218 | 171-171 |
| Şıralık (105) | 240-236 | 249-243 | 210-210 | 247-237 | 210-196 | 273-263 | 143-133 | 191-189 | 249-247 | 170-165 | 210-210 | 171-169 |
| $\begin{aligned} & \text { Şıralık (107, } \\ & 108 \text { and 109) } \end{aligned}$ | 236-236 | 249-247 | 212-210 | 261-261 | 210-202 | 273-263 | 133-133 | 205-201 | 249-247 | 192-165 | 210-208 | 167-165 |
| Siyah (110) | 236-226 | 249-239 | 214-210 | 257-247 | 210-204 | 273-257 | 145-145 | 201-187 | 251-243 | 165-165 | 218-218 | 171-171 |
| Sungurlu (111) | 240-236-232 | 253-249-239 | 210-210 | 261-239 | 212-204 | 273-257 | 149-143-135 | 189-189 | 247-243 | 165-165 | 224-218 | 165-161 |
| Tahannebi (112)/ Tahannebi (113) | 228-226 | 247-239 | 218-214 | 247-239 | 210-210 | 275-273 | 143-135 | 193-189 | 257-257 | 165-165 | 202-202 | 171-165 |
| Timbo (114) | 232-232 | 255-247 | 214-210 | 261-247 | 216-196 | 273-257 | 151-143 | 205-203 | 247-239 | 170-165 | 218-204 | 171-171 |
| Vanki (115) | 240-236 | 249-249 | 210-210 | 261-247 | 210-196 | 273-251 | 143-133 | 191-189 | 251-247 | 192-165 | 210-210 | 171-169 |
| Vilki (116)/ <br> Abderi (55) | 236-234 | 247-243 | 214-210 | 237-237 | 210-210 | 263-241 | 155-135-133 | 193-189 | 249-249 | 170-165 | 222-208 | 171-169 |
| Chasselas | 236-228 | 247-239 | 214-210 | 271-221 | 216-212 | 241-241 | 143-133 | 205-195 | 259-251 | 198-192 | 218-204 | 167-167 |
| Nebbiolo | 236-232 | 249-247 | 214-210 | 271-239 | 212-212 | 263-241 | 155-133 | 201-195 | 251-243 | 192-170 | 218-218 | 167-167 |
| Pinot Noir | 238-228 | 243-239 | 218-216 | 239-221 | 216-216 | 273-241 | 151-137 | 195-189 | 245-239 | 198-170 | 204-204 | 167-157 |
| Syrah | 232-226 | 239-239 | 216-210 | 231-221 | 216-212 | 273-241 | 133-133 | 195-189 | 251-245 | 198-195 | 218-218 | 171-167 |

Genetic characterization and relationships of grape cultivars

Table 3. Number of alleles $(N)$, observed heterozygosity $\left(\mathrm{Ho}_{\mathrm{o}}\right)$ and probability of identity (PI) of 89 cultivars analysed at 12 microsatellite markers

| Loci |  | N | Ho |
| :--- | ---: | :---: | :---: |
| PI |  |  |  |
| VVMD5 | 9 | 0.8347 | 0.087 |
| VVMD7 | 13 | 0.8123 | 0.109 |
| VVMD24 | 6 | 0.5979 | 0.231 |
| VVMD28 | 16 | 0.8118 | 0.114 |
| VVMD31 | 9 | 0.7801 | 0.147 |
| VVMD32 | 16 | 0.8531 | 0.067 |
| VVS2 | 14 | 0.8607 | 0.061 |
| VrZAG62 | 11 | 0.8246 | 0.099 |
| VrZAG79 | 14 | 0.8587 | 0.723 |
| VMC2C3 | 10 | 0.6955 | 0.204 |
| VMC2H4 | 12 | 0.8732 | 0.054 |
| VMC5A1 | 8 | 0.7457 | 0.161 |
| Total | 138 |  |  |
| Mean | 11.5 | 0.7957 |  |
| Cumulative |  |  | $1.666 \mathrm{e}-12$ |

paper. The true-to-type 'Gemre Siyah' should be the one with two accessions (75 and 76). Similarly, the true-totype 'Şıralık Beyaz' should be the one with three accessions (107, 108 and 109). The genotypes of 'Erik Kara', 'Hatun Parmağı' and 'Kabarcık' in Benjak et al. (2005) were difficult to harmonize with our data since they did not use any standard cultivar. However, the cultivar 'Bogdanuša' from Croatia was common to both Benjak et al. (2005) and Sefc et al. (2000), so that we were able to adjust the allele sizes of the Turkish accession. None

Table 4. Identical genotypes of grape cultivars from Georgia and Turkey uncovered by the analysis of 12 microsatellite markers: identical pairs with different colour and/or use are highlighted in italic

| Identical pairs |  |
| :---: | :---: |
| Georgia |  |
| Dondglabi (23) | Gorula (26) |
|  | Kapistoni Imeretinskii (28) |
|  | Tavkara (49) |
| Khounalige (31) | Chinuri (19) |
| Saperavi | Saperavi Pachkha (47) |
| Mrgvalmarzvala (46) |  |
| Chkovra (22) | Dondglabi Tetri (24) |
|  | Krakhuna (35) |
| Turkey |  |
| İri Daneli Ak Üzüm (85) | Italia (IASMA) |
| Parmak (102) | Jerusalem Bleu (UCD) |
| Mor Üzüm (96) | Tsaousi (GVD) |
| Aşeri (56) | Hasandede Beyazı (79) |
| Boğazkere (62 and 63) | Şaraplık Siyah (104) |
| Ekssi Kara (70) | Dökülgen (68) |
| Kayseri Karası (90) | İri Beyaz (84) |
| Muhammediye (98) | Kabarcık (89) |
| Öküzgözü (100) | Erik Kara (74) |
| Vilki (116) | Abderi (55) |

Table 5. Homonyms of grape cultivars from Armenia and Turkey uncovered by the analysis of 12 microsatellite markers

| Homonym pairs |  |
| :---: | :---: |
| Armenia |  |
| Areni Chernyi (2) | Areni Chernyi (3) |
| Turkey |  |
| Burdur Dimriti (64) | Burdur Dimriti (65) |
| Dökülgen (67 and 69) | Dökülgen (68) |
| Ekşi kara (70) | Ekşi kara (71) |
| Gemre Siyah (75 and 76) | Gemre Siyah (77) |
| Şıralık (107, 108 and 109) | Şıralık (105) |

matched our corresponding accessions at the five markers in common. In particular, Benjak et al. (2005) found that their 'Hatun Parmağı' was identical to 'Kişmiş'. According to Galet (2000), this is a synonym of 'Sultanina' (also called 'Thompson Seedless'). However, 'Kişmiş' genotype did not match any 'Sultanina'/'Thompson Seedless' accession (e.g. Sanchez-Escribano et al., 1998; Sefc et al., 1998b; Crespan et al., 1999). In Benjak et al. (2005), 'Kabarcik' appeared to be a clonal mutation of 'Kissmiss', but our 'Kabarcik' (two independent accessions) had a different genotype identical to 'Muhammediye'. As a consequence, there is obviously a need to determine the true-to-type accessions for many cultivars in Turkey, Georgia or Armenia, by verifying ampelographic descriptions and/or by searching for additional accessions.

## Genetic relationships

Since ‘İri Daneli Ak Üzüm' turned out to be identical to the cultivar 'Italia' which certainly belongs to other germplasms, it was discarded from the genetic analysis. However, the four standard cultivars ('Chasselas', 'Nebbiolo', 'Pinot Noir' and 'Syrah') were kept as outgroups. For chimeric genotypes (Table 2), we have discarded the alleles with the weakest signal. The cladogram in Fig. 1 represents the genetic relationships of the Anatolian and Transcaucasian cultivars. Five major groups of cultivars were detected.

## Group 1

Group 1 appeared as predominantly Turkish (19 cultivars), with only one cultivar from Armenia ('Areni Chernyi' 2) and one from Georgia ('Chitistvala Bodburi'). As mentioned above, 'Areni Chernyi' 2 is not likely to be true-to-type, and this accession could represent a misnamed introduction from Turkish germplasm. 'Chitistvala Bodburi' is supposed to be exclusively cultivated in Georgia, but it may also represent an introduction from

Turkey. Interestingly, the closely related pair 'Ekşi Kara' 71 and 'İri Kara' clustered with the homonym 'Ekşi Kara' 70/'Dökülgen' 68. We investigated their possible relationships and found that the genotype of 'Ekşi Kara' 71 was consistent with being the progeny of 'İri Kara’ and 'Ekşi Kara' 70/'Dökülgen' 68. However, 12 microsatellite markers are not enough for parentage analysis, and according to other studies (Sefc et al., 1998c; Bowers et al., 1999a; Vouillamoz et al., 2003) we would suggest analysing a minimum of ca 30 markers to verify this parentage. Similarly, our data suggested a possible parent-progeny relationship between 'Sungurlu' and 'Asseri'/‘Hasandede Beyazı' (Table 2), two cultivars that clustered together in Fig. 1. Interestingly, the Vitis International Variety Catalogue (VIVC) lists 'Sungurlu' as a synonym of 'Hasandede Beyazl'. This synonymy is not supported by our data (Table 2), but we suggest that they are closely related.

## Group 2

Group 2 was the largest and contained accessions from all three countries. It was subdivided into three distinct sub-groups (2.1-2.3). Sub-group 2.1 was predominantly Armenian, with the exception of 'Hatun Parmağı' from South-Eastern Anatolia, not far from the Armenian border. The true-to-typeness of 'Hatun Parmağı' still has to be established, since our accession did not match the homonym in Benjak et al. (2005). However, the name similarity between 'Hatun Parmağ' and 'Khatun Khardjzhi' from Armenia and their clustering together let us suggest that our accession of 'Hatun Parmağı' could be an Armenian introduction to Turkey, even if their colours and use are different (Table 1). Moreover, 'Hatun Parmağ1' is a local name that does not exist in VIVC or in Galet (2000). 'Areni Chernyi' 3 clustered with other Armenian cultivars: this supports our hypothesis that 'Areni Chernyi' 3 is true-to-type and 'Areni Chernyi' 2 is another cultivar, probably introduced from Turkey. Sub-group 2.2 can be separated into two distinct clusters, one comprised of Armenian and Georgian grapes and one made of Turkish grapes. All these grapes might have a common origin. Their genetic (and perhaps phenotypic) similarity probably explains why the 'Şıralık Beyaz' homonyms clustered together and share the same name. Sub-group 2.3 can be separated into four distinct clusters. Cluster 2.3.1 almost exclusively consisted of Georgian grapes with the exception of 'Kachet' from Armenia. However, as suggested by its etymology, 'Kachet' is supposed to have been introduced from Khakhetia in Georgia (G. Melyian, personal communication). Our results supported this hypothesis. According to VIVC, 'Kisi' is an artificial cross 'Mtsvane' $\times$ ' Rkatsiteli', and 'Mtsvane' is a synonym of 'Kundza' (VIVC; Galet, 2000). This parentage hypothesis was not
supported here (Table 2). Our data suggested a possible parent-progeny relationship between 'Kvira' and 'Meskhuri Shavi' (Table 2) that clustered together in Fig. 1. Cluster 2.3.2 contained grapes from Georgia and Turkey that might have a common origin. All these Georgian cultivars are cultivated in western Georgia (Table 1). In particular, the Georgian 'Khupishizh' clustered with Turkish cultivars probably because it is cultivated in Abkhazia, a region bordering Turkey. The closely related pair 'Boğazkere' (identical to 'Şaraplık Siyah') and 'Morek' had genotypes consistent with a possible parent-progeny relationship (Table 2). Clusters 2.3.3 and 2.3.4 were mostly from Turkey and rather isolated. They might have different origins.

## Group 3

Group 3 was comprised almost entirely of Turkish cultivars, with the exception of 'Mskhali' from Armenia and 'Odzhaleshi' from Georgia. 'Mskhali' is mainly used for brandy in Armenia. Since brandy production came later than wine production in Armenia, it is reasonable to suggest that 'Mskhali' was introduced from Turkey. 'Odzhaleshi', meaning 'grape to put on a tree', is one of the best red wine varieties and is mainly cultivated in Mingrelia in Western Georgia, near the Turkish border. Although it is considered an ancient Georgian variety, its position in Fig. 1 suggests an introduction from Turkey.

## Group 4

Group 4 consisted of miscellaneous cultivars from all studied areas and was considerably separated from the other groups. Among those from Armenia, 'Ak-Kaltak' is mainly cultivated in Uzbekistan, 'Karmir Kakhani' is a dioecious traditional table grape and 'Vardagujn Jerevani' is a seedless cultivar probably obtained from a deliberate cross (G. Melyian, personal communication). These particularities could explain why they did not cluster with other Armenian varieties (sub-groups 2.1 and 2.2). 'Saperavi Mrgvalmarzvala/Pachkha' is the most praised red wine variety in Georgia. However, it was genetically isolated from many other Georgian varieties (sub-groups 2.2 and 2.3 and group 5). The Turkish 'Mor Üzüm' clustered with several Georgian grapes. We found (Table 4) that 'Mor Üzüm' was identical to the Greek 'Tsaousi', and Galet (2000) suggested it is one and the same as 'Chaouch'. 'Chaouch' is a table grape widespread all over the Near and Middle East. It is known in Turkey as 'Çavuş Chaouch', a name encompassing several distinct types. According to our data, this variety might originate from Georgia.

## Group 5

Group 5 was also considerably separated from the other groups. Interestingly, it exclusively consisted of Georgian
cultivars that clustered with all four Western European standards ('Chasselas', 'Nebbiolo', 'Pinot' and 'Syrah'). This suggests that these Western European cultivars, or more likely some of their ancestors, initially originated from Georgia. These four Western European cultivars are supposed to be quite divergent from each other, but they most likely clustered together because they are more related to each other than to any Eastern cultivars. In particular, close clustering between 'Pinot' and 'Syrah' can be explained by their recently proven genetic relationship (J. F. Vouillamoz and M. S. Grando, unpublished data). However, the scale of the cladogram is not linear, so that close clustering between e.g. 'Chasselas' and 'Nebbiolo' does not represent the same genetic similarity as e.g. 'Ekşi Kara' T71 and its putative parents.

On the lower part of the cladogram (Fig. 1), most of the Armenian, Georgian and Turkish germplasms were generally well separated. This suggests that very few recent grape exchanges occurred between these areas, with the exception of some clusters with cultivars from all three areas. On the whole, Armenian cultivars were usually closer to Georgian than Turkish cultivars. On the upper part of the cladogram (Fig. 1), groups 1 and 2 clustered together, suggesting some common ancestors, and they could represent recent evolution of these cultivars in Trancaucasia and Anatolia. Groups 3, 4 and 5 were very distinct and could represent three separate ancient origins. One of them (the Georgian group 5) might have common ancestors with Western European cultivars.

## Conclusion

In the present work on Anatolian and Transcaucasian cultivars, we detected 17 identical genotypes and six homonymy cases among 116 accessions corresponding to 89 distinct grape cultivars according to our analyses, thus helping to improve the management of these germplasms. However, they need to be more deeply investigated, since we only genotyped here a fraction of the existing autochthonous cultivars. Special care should be taken to conserve these cultivars in ampelographic collections, especially in areas where old vineyards are rapidly disappearing (Gasparyan and Melyan, 2003). Additional studies on ampelography, origin, etymology and distribution of these cultivars should help in solving the true-to-typeness issues mentioned in this paper. Similarly, we suggested a few possible parentages within our sampling, but additional microsatellite markers would be undoubtedly necessary to test these hypotheses. The cladogram topology suggested that most of the Armenian, Georgian and Turkish germplasms were well separated and could have multiple origins, although they are likely to have common ancestors. A few examples of varieties having possibly been exchanged between these
countries were discussed above. Since the four varieties from Western Europe were closely related to a group of Georgian cultivars, we propose that they could have some ancient Georgian ancestors. As a next objective, we will investigate the genetic relationships between the cultivars genotyped in this study and additional Western European cultivars as well as wild grapes from Transcaucasia and Anatolia in order to determine the origins of Western European cultivars and locate putative sites of primary domestication. For the present study, it has not been possible to expand the sampling area to neighbouring countries, but we plan to include Azerbaijan, Iran and Lebanon in the near future. Hopefully, an expansion of the database with the analyses of additional varieties will further elucidate the origins of grape cultivars.

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## References

Ağaoğlu YS and Celik H (1986) Conservation of Vitis vinifera L. germplasm in Turkey. Vignevini 13(Suppl. 12): 40-42.
Alleweldt G (1997) Genetics of grapevine breeding. Progress in Botany 58: 441-454.
Aradhya MK, Dangl GS, Prins BH, Boursiquot JM, Walker MA, Meredith CP and Simon CJ (2003) Genetic structure and differentiation in cultivated grape, Vitis vinifera L. Genetical Research 81(3): 179-192.
Belkhir K, Borsa P, Chikhi L, Raufaste N and Bonhomme F (1996-2002) GENETIX 4.04, logiciel sous Windows TM pour la génétique des populations. Montpellier: Laboratoire Génome, Populations, Interactions, CNRS UMR 5000, Université de Montpellier II.
Benjak A, Ercisli S, Vokurka A, Maletic E and Pejic I (2005) Genetic relationships among grapevine cultivars native to Croatia, Greece and Turkey. Vitis 44(2): 73-78.
Bowers JE, Dangl GS, Vignani R and Meredith CP (1996) Isolation and characterization of new polymorphic simple sequence repeat loci in grape (Vitis vinifera L.). Genome 39: 628-633.
Bowers JE, Boursiquot JM, This P, Chu K, Johansson H and Meredith CP (1999a) Historical genetics: the parentage of Chardonnay, Gamay, and other wine grapes of Northeastern France. Science 285: 1562-1565.
Bowers JE, Dangl GS and Meredith CP (1999b) Development and characterization of additional microsatellite DNA markers for grape. American Journal of Enology \& Viticulture 50: 243-246.
Chkhartishvili N (2003) IPGRI project on 'conservation and sustainable use of grapevine genetic resources in the Caucasus and Northern Black sea region'-implementation in Georgia. First meeting of the ECP/GR working group on Vitis, Paliæ, Serbia and Montenegro, 12-14 June.
Chkhartishvili N and Tsertsvadze NV (2003) Status of grapevine genetic resources (Vitis vinifera) in Georgia. Report at the international conference 'Retention and the Use of Genetic Resources of the Grapevine of the Caucasus and North Black Sea Area', Tbilisi, Georgia, 16 October.
Costantini L, Monaco A, Vouillamoz JF, Forlani M and Grando MS (2005) Genetic relationships among local Vitis vinifera cultivars from Campania (Italy). Vitis 44(1): 25-34.
Crespan M (2003) The parentage of Muscat of Hamburg. Vitis 42(4): 193-197.
Crespan M, Botta R and Milani N (1999) Molecular characterization of twenty seeded and seedless table grape cultivars (Vitis vinifera L.). Vitis 38: 87-92.
Ergül A, Marasali B and Ağaoğlu YS (2002) Molecular discrimination and identification of some Turkish grape cultivars (Vitis vinifera L.) by RAPD markers. Vitis 41(3): 159-160.
Franks T, Botta R and Thomas MR (2002) Chimerism in grapevines: implications for cultivar identity, ancestry and
genetic improvement. Theoretical and Applied Genetics 104: 192-199.
Galet P (2000) Dictionnaire encyclopédique des cépages. Paris: Hachette.
Gasparyan S and Melyan G (2003) Condition and prospects preservation of genetic resources of grapes on Armenia. Report at the international conference 'Retention and the Use of Genetic Resources of the Grapevine of the Caucasus and North Black Sea Area', Tbilisi, Georgia, 15 October.
Grando MS and Frisinghelli C (1998) Grape microsatellite markers-sizing of DNA alleles and genotype analysis of some grapevine cultivars. Vitis 37(2): 79-82.
Ketskhoveli N, Ramishvili M and Tabidze D (1960) Ampelography of Georgia. Tbilisi: Publishing House of the Academy of Sciences.
Lamboy WF and Alpha CG (1998) Using simple sequence repeats (SSRs) for DNA fingerprinting germplasm accessions of grape (Vitis L.) species. Journal of the American Society for Horticultural Science 123: 182-188.
Langella O (2002) POPULATIONS, version 1.2.28. Gif sur Yvette, Paris: CNRS UPR9034.
Lefort F and Roubelakis-Angelakis KA (2001) Genetic comparison of Greek cultivars of Vitis vinifera L. by nuclear microsatellite profiling. American Journal of Enology \& Viticulture 52: 101-108.
Levadoux L (1956) Les populations sauvages et cultivées de Vitis vinifera L. Annales d'Amelioration des Plantes 1: 59-118.
Lodhi MA, Ye GN, Weeden NF and Reisch BI (1994) A simple and efficient method for DNA extraction from grapevine cultivars and Vitis species. Plant Molecular Biology Reporter 12: 6-13.
Lopes MS, Sefc KM, Dias EE, Steinkellner H, Machado MLD and Machado AD (1999) The use of microsatellites for germplasm management in a Portuguese grapevine collection. Theoretical and Applied Genetics 99: 733-739.
Maghradze D (2003) Status of Vitis collections in Georgia. First meeting of the ECP/GR working group on Vitis, Paliæ, Serbia and Montenegro, 12-14 June.
Maletic E, Sefc KM, Steinkellner H, Kontic JK and Pejic I (1999) Genetic characterization of Croatian grapevine cultivars and detection of synonymous cultivars in neighboring regions. Vitis 38(2): 79-83.
McGovern PE (2003) Ancient Wine: The Search for the Origins of Viniculture. Princeton, NJ: Princeton University Press.
Meredith CP, Bowers JE, Riaz S, Handley V, Bandman EB and Dangl GS (1999) The identity and parentage of the variety known in California as Petite Sirah. American Journal of Enology \& Viticulture 50: 236-242.
Minch E, Ruiz-Linares A, Goldstein DB, Feldman M and CavalliSforza LL (1995) Microsat (Version 1.4d): A Computer Program for Calculating Various Statistics on Microsatellite Allele Data. Stanford, CA: University of Stanford.
Musayev MK (2003) Grapevine genetic resources in Azerbaijan. First meeting of the ECP/GR working group on Vitis, Paliæ, Serbia and Montenegro, 12-14 June.
Negrul AM (1938) Evolution of cultivated forms of grapes. Comptes Rendus (Doklady) Académie Sciences USSR 18: 585-588.
Nei M (1972) Genetic distance between populations. American Naturalist 106: 283-291.
Nei M (1973) The theory and estimation of genetic distance. In: Morton NE (ed.) Genetic Structure of Populations. Honolulu: University Press of Hawaii, pp. 45-54.
Nei M, Tajima F and Tateno Y (1983) Accuracy of estimated phylogenetic trees from molecular data. Journal of Molecular Evolution 19: 153-170.

Olmo HP (1995) The origin and domestication of the Vinifera grape. In: McGovern PE, Fleming SJ and Katz SH (eds) The Origins and Ancient History of Wine. Amsterdam: Gordon and Breach Science Publishers, pp. 31-43.
Page RD (1996) TREEVIEW: an application to display phylogenetic trees on personal computers. Computer Applications for Bioscience 12: 357-358.
Raymond M and Rousset F (1995) GENEPOP (version 1.2): population genetics software for exact tests and ecumenicism. Journal of Heredity 86: 248-249.
Riaz S, Garrison KE, Dangl GS, Boursiquot JM and Meredith CP (2002) Genetic divergence and chimerism within ancient asexually propagated winegrape cultivars. Journal of the American Society for Horticultural Science 127: 508-514.
Rogers JS (1972) Measures of genetic similarity and genetic distance. Studies in Genetics VII. Publication 7213. Austin: University of Texas, pp. 145-153.
Sanchez-Escribano EM, Ortiz JM and Cenis JL (1998) Varietal identification of table grape cultivars (Vitis vinifera L.) by the isoenzymes from the woody stems. Genetic Resources and Crop Evolution 45: 173-179.
Sanchez-Escribano EM, Martin JR, Carreno J and Cenis JL (1999) Use of sequence-tagged microsatellite site markers for characterizing table grape cultivars. Genome 42: 87-93.
Sefc KM, Regner F, Glossl J and Steinkellner H (1998a) Genotyping of grapevine and rootstock cultivars using microsatellite markers. Vitis 37(1): 15-20.
Sefc KM, Guggenberger S, Regner F, Lexer C, Glossl J and Steinkellner H (1998b) Genetic analysis of grape berries and raisins using microsatellite markers. Vitis 37(3): 123-125.
Sefc KM, Steinkellner H, Gloessl J, Kampfer S and Regner F (1998c) Reconstruction of a grapevine pedigree by microsatellite analysis. Theoretical and Applied Genetics 97: 227-231.

Sefc KM, Regner F, Turetschek E, Glossl J and Steinkellner H (1999) Identification of microsatellite sequences in Vitis riparia and their applicability for genotyping of different Vitis species. Genome 42: 367-373.
Sefc KM, Lopes MS, Lefort F, Botta R, Roubelakis-Angelakis KA, Ibanez J, Pejic I, Wagner HW, Glössl J and Steinkellner H (2000) Microsatellite variability in grapevine cultivars from different European regions and evaluation of assignment testing to assess the geographic origin of cultivars. Theoretical and Applied Genetics 100: 498-505.
Takezaki N and Nei M (1996) Genetic distances and reconstruction of phylogenetic trees from microsatellite DNA. Genetics 144(6): 189-399.
This P, Jung A, Boccacci P, Borrego J, Botta R, Costantini L, Crespan M, Dangl GS, Eisenheld C, Ferreira Monteiro F, Grando MS, Ibanez J, Lacombe T, Laucou V, Magalhaes N, Meredith CP, Milani N, Peterlunger E, Regner F, Zulini L and Maul E (2004) Development of a standard set of microsatellite reference alleles for identification of grape cultivars. Theoretical and Applied Genetics 109: 1448-1458.
Thomas MR and Scott NS (1993) Microsatellite repeats in grapevine reveal DNA polymorphisms when analysed as sequence-tagged sites (STSs). Theoretical and Applied Genetics 86: 985-990.
Vavilov NI (1926) Studies on the Origin of Cultivated Plants. Leningrad: Institute of Applied Botanical Plant Breeding.
Vouillamoz J, Maigre D and Meredith CP (2003) Microsatellite analysis of ancient alpine grape cultivars: pedigree reconstruction of Vitis vinifera L. 'Cornalin du Valais'. Theoretical and Applied Genetics 107: 448-454.
Wagner HW and Sefc KM (1999) IDENTITY 1.0. Vienna: Centre for Applied Genetics, University of Agricultural Sciences.
Zohary D and Hopf M (2000) Domestication of Plants in the Old World. New York: Oxford University Press.


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[^1]:    Where possible, names are spelled according to the Vitis International Variety Catalogue (http://www.genres. de/idb/vitis/vitis.htm). The colour of the berries is indicated as $n$ (noir, blue berries), $b$ (blanc, white berries) or $r$ (rose, pink berries), according to international usage. Grapes can be used for table ( t ), wine ( w ) or raisin ( r ). Accessions were sampled from grape collections or vineyards. Cultivation area gives an idea of the grape's distribution.
    ${ }^{\text {a }}$ Names not found in the Vitis International Variety Catalogue are marked with an asterisk (*).
    ${ }^{\mathrm{b}}$ Accessions marked with a dagger $\left({ }^{+}\right)$were sampled from vineyards. Abbreviations for ampelographic collections are: YR, Yeras Rahun collection, Scientific Centre of Agrochemistry and Farmer, Armenia; N, Nalbaldian collection, Armavir region, Armenia; TVI, Tbilisi Viticulture Institute collection; AUKVRES, Ankara University, Kalecik Viticultural Research and Experiment Station; TNGRV, Turkey National Germplasm Repository Vineyard.
    ${ }^{\text {c }}$ Particular areas within the country or areas other than sampling area, based on Galet (2000) or various personal communications.

