

THE SOFTWARE FOR AN ‘UNIVERSAL GRAPEVINE DATABASE’

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Abstract

A software for an “universal grapevine database” has been developed to support the activity of characterization of grapevine biodiversity. This database allows a decentralized data management: the registered users have the possibility to submit and to manage his own data at any time. The uploaded data have three different levels of visualisation: private level (the data are visible only to his submitter), middle level (the data are visible to all other submitters), public level (the data are visible to all public users). Only the data approved by a specific scientific committee can be elevated to public level. When a submitter introduces the microsatellite data in the database the application allows a specific standardization procedure based on some specific selected accessions called ‘system accessions’. The main classes of data represented in the database are the ‘grapevine variety’, the ‘ampelographic-ampelometric and phenological-productive descriptors’ (as reported in the second edition of the OIV descriptor list for grape varieties and *Vitis* species), the ‘microsatellite profile’, the polyphenols and aroma profiles. Several search options have been implemented: a search by variety and other general parameter, a search by ampelographic and ampelometric parameters, a search by microsatellite profile. About microsatellite profile it is also possible a search by range and by a particular standardization procedure. The application has been implemented using the most recent database software and languages, so it is flexible and dynamics especially as concern the addition of other classes of data, like new type of descriptors and molecular markers. The application is on the web at <http://www.vitisdb.it> and at present it is adopted for the ‘Italian Vitis Database’ managed by the “Vitis Database Working Group”.

INTRODUCTION

In viticulture the identification of variety is extremely important for commercialization of planting materials, for recovery, characterization and valorization of local germplasm, and therefore for preservation of biodiversity. But, the high number of varieties, their rapid diffusion, their morphological and phenological variability, and consequently the presence of many synonyms and homonyms, get grapevine variety’s identification particularly complex.

The development of DNA molecular markers during the last decades, and in particular of ‘microsatellite’ molecular markers, provides the opportunity, as never before, to identify with certainty each grapevine variety.

In this contest, the Department of Fruit Science and Plant Protection, University of Pisa, created the application for a 'Universal Grapevine Database' that supports some important innovations.

In 2004, Dr. Claudio D'Onofrio, made the logical and conceptual scheme of an universal grapevine database as his activity of thesis in the master of bioinformatics at the University of Turin (D'Onofrio, 2004). This project was presented during the Italian national symposium on autochthonous grapevine varieties in 2006 ('I vitigni autoctoni minori: aspetti tecnici, normativi e commerciali', 30 November – 1 December 2006, Turin). In February 2007, in a meeting among the members of the majority of the Italian viticulture research Institutions, it was proposed the physical implementation of the above application and the adoption for a possible Italian grapevine database. Thanks to the financial support of the agriculture society 'ColleMassari s.p.a.', it was achieved the physical implementation of the proposed application. The application was presented in a following meeting among the members of the Italian viticulture Institutions in May 2008. The participants of this meeting definitively decided for the adoption to the presented application for the 'Italian Vitis Database' and established the "Vitis Database Working Group" (Fig. 1).

The application was also presented in Gödöllő, Hungary, during the "Second Annual Workshop for the EU-project GrapeGen06" (8th – 10th July, 2008), while the 'Italian Vitis Database' and the 'Vitis Database Working Group' have been officially presented during the second Italian national symposium of viticulture (Marsala, Sicily, 14-19 July 2008. The public level of the 'Italian Vitis Database' (www.vitisedb.it) has been opened during the third Italian national symposium of viticulture (S.Michele all' Adige, Trento, 5–9 July 2010)..

MATERIALS AND METHODS

The logical and conceptual scheme of the application for a universal vitis database was mainly prepared in 2004 by Dr. D'Onofrio as his thesis of the master of bioinformatics of University of Turin, while the physical implementation has started in 2007 and it is still in progress.

The logical and conceptual scheme, and consequently the definition of the static (data) and dynamic (data management) aspects, followed the relational model and it resulted from the analysis of literature and plant databases, and from the specificity of viticulture and of possible requirements of the potential users of the application.

As concern the physical implementation, it has been identified and studied the main functions of the application and consequently it has been chosen the more appropriate informatics technologies. Then, it has been decided the informative aspect of the web site and the relationships of the different levels with the intend to obtain an homogeneous, well structured and easily navigable system. Thereafter, the structure and colours of the web site pages and the main masks for the management, visualization and refreshment of data have been designed in conformity to logical and conceptual scheme. Subsequently, the masks of management of all classes of data, of general utility and administration have been created

RESULTS AND DISCUSSION

The application for the 'Universal Grapevine Database' allows a decentralized data management: the registered users have the possibility to submit and to manage his own data at any time.

Level of visualization

In this database the data have three different levels of visualisation: private level, middle level and public level. The registered user (submitter) has the access to his private level and to middle level.

1. Private level. The data in the private level are visible only to his submitter. This level represents a powerful tool for the description, characterization and management of grapevine accession. All the private data can be compared with the data in the middle level. The submitter can elevate to middle level each single accession with all the data associated.

2. Middle level. The data in the middle level are visible to all other submitters. The middle level allows to share the data among all the submitter of the database and consequently it allows the identification of synonyms and homonyms, of duplications and errors before the elevation of the data in the public level.

3. Public level. The data in the public level are visible to all public users. The elevation of data from private or middle level to public level is requested by his submitter and must be approved by a specific Scientific Committee. So, while the private and middle levels represent a useful tool for the characterization of grapevine varieties and accessions, the data of the public level represents a reference for all the whole scientific community and all public users.

Users

The application has three types of users: public users, registered users (submitters), and the administrator.

1. Public users. All the generic people. They can navigate in the public level of the database without authentication.

2. Submitters. The submitters are registered users that with a specific username and password can have the access to the private and middle levels.

3. Administrators. Responsible for database's functionality. The administrator is also responsible for the management and modification of the application's structure upon suggestion of the Scientific Committee.

Scientific Committee

The Scientific Committee has the responsibility to approve the data for the public level, to approve the registration of new submitters, to make decisions about the management of the database and about the modification of the software's structure. Actually the Scientific Committee is composed by the database administrator, member of Department of Fruit Science and Plant Protection (University of Pisa) that promoted the project, expert in viticulture, expert in ampelography, expert in molecular biology, expert in analysis of grape secondary metabolites.

Classes of data

The main classes of data in the database are: 'grapevine variety', 'grapevine accession', 'microsatellite profile', 'ampelographic-ampelometric and phenological-productive descriptors', berry polyphenols and aroma profiles (Fig. 2).

1. Variety. In the 'variety' main page (Fig. 3) some general information are reported: botanical information, official and documented synonyms, wrong

denominations, released clones, a short history of the variety and information on distribution and variation, agronomical characteristics, technological use and bibliography. The general information of a variety are managed by a single submitter (reported on the variety's page) with the permission of the Scientific Committee. In the variety's page are also showed the microsatellite profile, ampelographic-ampelometric descriptors and the photos of the 'variety main accession'. The 'variety main accession' must be approved by the Scientific Committee, and it is an accession that has at least the minimum microsatellite profile, the minimum ampelographic descriptors and a photo of shoot, adult leaf and bunch. The page of a variety can appear in the public level only if the main accession has been approved for public level. A variety groups all the accessions having a microsatellite profile compatible with that one of the 'variety main accession'.

2. Accession. The upload of an accession in the database needs at least the minimum microsatellite profile. The other classes of accession's data are: ampelographic-ampelometric descriptors, phenological-productive descriptors, berry polyphenols, berry aroma, berry sensorial profile, sanitary status, true-to-type information, photos of all the main ampelographic organs and bibliography.

3. Microsatellite profile. The microsatellite profile of an accession can include all the possible characterized microsatellite loci. Actually the minimum microsatellite profile includes the 6 loci suggested in the EU-project GENRES CT96 No 81 and it will be extended to the 9 loci identified in the current European GrapeGen06 project. The application allows a specific standardization's procedure of microsatellite profile based on some specific selected accessions called 'system accessions' and on the submitter microsatellite profile of reference (submitter microsatellite profile). Both no-standardized data and standardized microsatellite data are maintained in the database (the no-standardized data are visible only in the middle and private levels). Only the accessions with standardized microsatellite profile can be elevated to the middle and public levels.

4. Morphological, phenological and productive descriptors. In the database it is possible to upload all the ampelographic, ampelometric, phenological and productive descriptors as reported in the second edition of the OIV descriptor list for grape varieties and *Vitis* species (OIV, 2009). For the variety's main accessions there is a minimum list of OIV ampelographic descriptors.

5. Berry profiles. Berry polyphenol, aroma and sensorial profiles include all the main parameters reported in the literature, separately for skin, pulp and seed. The system automatically calculates some specific ratios among polyphenols and aroma that are useful for the variety characterization as reported in the literature.

6. Sanitary status. In the sanitary status section are reported the established information decided at international and national level.

7. True-to-type. About the true-to-type, for each accession the information as suggested by Schneider Anna are reported

8. Photos.

The photos of all the main ampelographic organs can be introduced in the database: shoot, bud, leaf, petiol sinus, flower, berry, seed and bunch.

Language

The application can support all the possible languages and the language can be changed at any time during the navigation. Actually the English and Italian language have been implemented.

Search options

Several search options have been implemented: by general parameters, by ampelographic and ampelometric descriptors, by microsatellite profile.

The search by microsatellites is possible by ranges, by a three steps standardization procedure (Fig. 4), that calculate the similarity index and the cumulative similarity index and also indicate the possibility of direct parentage. Moreover, the graphical representation of results indicates by colours the similarity of each locus compared among the query and the subject: red, both alleles equals; yellow, one allele equal; blue, different alleles; gray, locus non compared. In the public level the search by microsatellite is performed on varieties' microsatellite profile of reference (microsatellite profile of the variety main accession), while in the middle and private level the research is performed both in standardized and non-standardized microsatellite profile of each accession.

Agreement and registration

For uploading data in the database or just for the access to the middle level it is necessary to be a registered member. For registration it is necessary to sign a specific agreement. The registered users also became members of the 'Vitis Database Working Group' and consequently can take part in the management and improvement of the application.

CONCLUSIONS

The application has been implemented using the most recent database software and languages, so it is flexible and dynamic especially as concern the addition of other classes of data, like new type of descriptors and molecular markers. Actually the application is adopted for the 'Italian Vitis Database' but it could be adopted for all other national databases and eventually for a 'Universal Grapevine Database'.

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Figures



Fig. 1. Home page of the 'Italian Vitis Database' (www.vitisdb.it) managed by the 'Vitis Database Working Group'.



Fig. 2. Descriptors' page.

Italian Vitis Data Base

donofrio logout language: English ▾

Variety *Sangiovese* (vitisdb-var_1)

view | edit

general info | microsatellite profile | ampelography | ampelometry | accessions

general information managed by
 Dipartimento di Coltivazione e Difesa delle Specie Legnose "G Scaramuzzi" - Università di Pisa

botanical information

name:	Sangiovese	subspecies:	sativa
species:	Vitis vinifera	parent 2:	Calabrese di montenuovo
parent 1:	Cilegiolo	variety for:	wine
variety group:	not available		

synonyms (23)

official synonyms (1)

- > "Sangiovese" ()

documented synonyms (18)

- > "Morellino" (Scansano, IT), "Nielluccio" (Corse, FR), "Prugnolo" (), "Prugnolo gentile" (), "Sangiochetto" (), "Sangiovese di lamole" (), "Sangiovese di romagna" (), "Sangiovese dolce" (), "Sangiovese gentile" (), "Sangiovese grosso" (), "Sangiovese romagnolo" (), "Sangiovese toscano" (), "Sangiovese" (), "San Gioveeto grosso" (), "Sangiovese montanino" (), "Sanzoveto" (), "San zoveto" (), "Uvetta" ()

wrong denominations (5)

- > "Cilegiolo" (), "Montepulciano" (), "Moscato nero" (), "Moscato rosso" (), "Uva Tosca" ()

released clones (87)

- > "I - 9 ISV - CSV", "I - Agri 3", "I - Agri 45", "I - AGRI 6", "I - Ampelos TEA 10D", "I - Ampelos TEA 15", "I - Ampelos TEA 7", "I - Ampelos TEA 8", "I - Ampelos TEA 9", "I - AP-SG 1", "I - AP-SG 2", "I - B-BS-11", "I - BF 10", "I - BF 30", "I - CCL 2000/5", "I - CCL 2000/6", "I - CCL 2000/7", "I - C-FUTURO 1", "I - C-FUTURO 2", "I - C-FUTURO 3", "I - C-FUTURO 4", "I - CHIANTI CLASSICO 2000/1", "I - CHIANTI CLASSICO 2000/2", "I - CHIANTI CLASSICO 2000/3", "I - CHIANTI CLASSICO 2000/4", "I - CSV-AP-SGS", "I - CSV - SG 4", "I - CSV - SG 6", "I - Fedit 20-CH", "I - Fedit 21-CH", "I - Fedit 22-CH", "I - Fedit 2 ESAVE", "I - Fedit 30 ESAVE", "I - Fedit 38 ESAVE", "I - ISV 2", "I - ISV RC 1", "I - JANUS-10", "I - JANUS-20", "I - JANUS 50", "I - MI-BF-50", "I - MI-TIN-20", "I - MI-TIN-30", "I - MI-TIN-40", "I - Montalcino 42", "I - Peccioli 1", "I - Rauscedo 10 (Grosso Lamole)", "I - Rauscedo 24 (Medio Predappio)", "I - SA - PA - 9", "I - SG 12 T", "I - SG 2 T", "I - SG 4 T", "I - SG-CDO-4", "I - SG CDO 6", "I - SG P.A 1", "I - SG P.A 0", "I - SS F9 A5 40", "I - TCA 6", "I - TIN 10", "I - TIN 50", "I - UBA 63/F", "I - UBA 63/L", "I - UBA 74/C", "I - UBA 79/C", "I - UNIMI-VITIS SANG VV 101", "I - UNIMI-VITIS SANG VV 110", "I - U.S. FI-PI 172", "I - U.S. FI-PI 3", "I - VCR 1", "I - VCR 102", "I - VCR 103", "I - VCR 105", "I - VCR 106", "I - VCR 108", "I - VCR 109", "I - VCR 116", "I - VCR 16", "I - VCR 19", "I - VCR 207", "I - VCR 209", "I - VCR 214", "I - VCR 218", "I - VCR 23", "I - VCR 237", "I - VCR 30", "I - VCR 4", "I - VCR 5", "I - VCR 6"

images

shoot	shoot upper side	shoot lower side	bud
leaf	leaf upper side	leaf lower side	petiol sinus
bunch	berry	seed	

short history


The lack of historical evidence before the sixteenth century makes it very difficult to trace the origins of Sangiovese in Middle Italy viticulture and the leading hypothesis explains the great interest to find out the origin of its name. The name Sangiovese over which "Toscana" and "Emilia Romagna" have been quarrelling for many years is linked back to myth which recalled blood, one of the symbols associated with Jove- (sanguis Jovis). The name's semantics refers to yoke, "giogo" (jugum), "giove-vese", that is blood of hill yokes, or of a wine "giovevole al sangue".

Other connections among the Etruscan language, the religious aspects of the name are hypothesized. In a phrase not completely interpreted written in an Etruscan calendar used to wrap an Egyptian mummy of the first century AD, the word "Sangiovese" that has an excellent assonance to the words used to define Sangiovese are other assonances related to the ritual sphere with Sangiovese such as "Sangiovese" someone who does a rite, "thezin-eis" (offer to the God) or "sani-sva", verb "Sangiovese" that has the meaning of father or ancestor standing for father (Sangiovese 2001).

Tracing back the origins of the Sangiovese grapevine to the Etruscan culture is very fascinating, but the latest findings

Completato

Fig. 3. Variety's page.



Italian Vitis Data Base

Home

About VitisDB

Search

Descriptors

Varieties

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Participants

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Links

login

Variety search by standardized microsatellite profile

generic |
 by microsatellite profile |
 by standardized microsatellite profile

Step 3 - risultati della ricerca effettuata

Variety selected for comparison **Sangiovese**

SSR locus:	VVS2	VVMD5	VVMD7	VVMD27	VrZAG62	VrZAG79	VVMD25	VVMD28	VVMD32									
allele:	A1 A2	A1 A2	A1 A2	A1 A2	A1 A2	A1 A2	A1 A2	A1 A2	A1 A2									
size:	131 131	225 235	238 262	177 183	192 194	243 259	242 242	235 245	253 257									
Your SSR locus size for Sangiovese:	133 133	225 235	240 264	180 186	196 198	243 259	240 240	237 247	253 257									
deviation of allele lengths in base pairs	-2 -2	0 0	-2 -2	-3 -3	-4 -4	0 0	2 2	-2 -2	0 0									
profile of the variety to be identified:	133 135	225 227	240 250	180 194	188 198	249 255	248 254	239 249	261 263									
standardized profile to be identified:	131 133	225 227	238 248	177 191	184 194	249 255	250 256	237 247	261 263									
Error range (+/-)	2																	
range considered equals	129 133	131 135	223 227	225 229	236 240	246 250	175 179	189 193	182 186	192 196	247 251	253 257	248 252	254 258	235 239	245 249	259 263	261 265

order by: SI Sij compared locus

search

similarity graph

Legend

both alleles equals
one allele equal
different alleles
not compared

SSR locus:	VVS2	VVMD5	VVMD7	VVMD27	VrZAG62	VrZAG79	VVMD25	VVMD28	VVMD32	SI	Sij	compared locus	equals allele	may be parent
Aleatico										1.0	9.0	9	18	true
Teroldego										0.5	4.5	9	9	false
Famoso										0.5	4.0	8	8	false
Sangiovese										0.4444	4.0	9	8	false
Vermentino										0.3889	3.5	9	7	false
Griano										0.4167	2.5	6	5	false
Ruggine										0.4167	2.5	6	5	false
Arneis										0.3333	2.0	6	4	false

Search Results

Aleatico

SSR locus: VVS2 VVMD5 VVMD7 VVMD27 VrZAG62 VrZAG79 VVMD25 VVMD28 VVMD32

allele: A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2

query: 131 133 225 227 238 248 177 191 184 194 249 255 250 256 237 247 261 263

131 133 225 227 238 248 177 191 184 194 249 255 250 256 237 247 261 263

Teroldego

SSR locus: VVS2 VVMD5 VVMD7 VVMD27 VrZAG62 VrZAG79 VVMD25 VVMD28 VVMD32

allele: A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2

query: 131 133 225 227 238 248 177 191 184 194 249 255 250 256 237 247 261 263

135 153 225 227 238 246 175 181 192 192 243 255 240 242 229 237 241 265

Famoso

SSR locus: VVS2 VVMD5 VVMD7 VVMD27 VrZAG62 VrZAG79 VVMD25 VVMD28 VVMD32

allele: A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2 A1 A2

query: 131 133 225 227 238 248 177 191 184 194 249 255 250 256 237 247 261 263

131 131 225 227 238 246 177 187 186 202 241 245 242 242 229 261

Fig. 4. Search by standardized microsatellite profile.