# Grapevine cultivar Müller-Thurgau and its true to type descent

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## **Summary**

Ampelographic investigations supported by molecular marker analysis were used to reevaluate the progenitors of the grapevine cv. Müller-Thurgau. From these studies we conclude Müller-Thurgau to be a descent of the offspring of a Riesling and Madeleine Royale (syn. Königliche Magdalenentraube) hybridisation. The results reveal the importance of true to typeness of the grapevine varieties used for genotyping, parentage analyses, etc.

## Introduction

The grapevine cv. Müller-Thurgau is known to be crossed at Geisenheim at the "Königliche Lehranstalt für Obst- und Weinbau" by Prof. Dr. h.c. HERMANN MÜLLER-THURGAU in 1882 (BECKER 1982). In 1891 MÜLLER-THURGAU became director of the "Eidgenössische Lehrund Versuchsanstalt" at Wädenswil, Switzerland, and thus carried over cuttings of 150 preselected seedlings. One of these seedlings, breeding No. 58 (Riesling x Silvaner 1), was selected and propagated in 1897 for the first time. A few years later, in 1913, August Dern, a grapevine inspector in Franconia, brought cuttings of No. 58 which he named "Müller-Thurgau" back to Germany. Between 1920 and 1930 vines of Müller-Thurgau were grown on experimental plots in all German wine-growing regions. After World War II Germany needed an early ripening grape cultivar, easy to cultivate and with high yield stability. Elegant Müller-Thurgau wines were produced by reductive winemaking and the cultivar spread rapidly all over German vineyards. During the early seventies until the middle of the nineties the growing area of Müller-Thurgau exceeded that of Riesling. Müller-Thurgau still covers more than 20 % of today's vinegrowing area in Germany and is grown in several other countries, too (Becker 1982).

Concerning the descent of Müller-Thurgau as a Riesling x Silvaner cross doubts raised early: Müller-Thurgau himself replied in a letter to Dern that the cultivar brought to Franconia in 1913 was not the result of the first Riesling x Silvaner cross at Geisenheim (Eichelsbacher 1957). Therefore there has been much speculation about the true parents (Hillebrand *et al.* 1997). Many specialists suggested that Müller-Thurgau was a selfing of Riesling. Eichelsbacher (1957) indicated that Silvaner could not be a parent of Müller-Thurgau, comparing leaf and tip mor-

phology of Riesling, Silvaner, Müller-Thurgau and Rieslaner with those of selfings and backcrosses. In 1994, his hypothesis gained strong support by means of RAPD analysis, confirming (1) Riesling as one parent, (2) making the selfing hypothesis unlikely and (3) excluding Silvaner as the second parent (BÜSCHER et al. 1994). Analysis of microsatellite loci excluded that Riesling selfing generated Müller-Thurgau and confirmed that Silvaner is not a parent (THOMAS et al. 1994). In 1996, REGNER first proposed a cultivar of the Chasselas (Gutedel) family as the second gene donor. Microsatellites at 7 of 8 loci corresponded to the allele lengths of the Chasselas family (REGNER et al. 1996). A more detailed microsatellite analysis (Sefc et al. 1997) pointed to Admirable de Courtiller (syn. Chasselas de Courtiller), a table grape of minor importance, bred in the 19th century by Courtiller, director of the Botanical Garden in Saumur, France. In combination with the microsatellite pattern of Riesling the genotype of Müller-Thurgau could be explained at 24 different microsatellite loci (SEFC et al. 1997).

However, doubts about the proposed parentage still remained. According to the principles of heredity, characteristics of the parents are manifested in their offsprings. But from its phenotype Müller-Thurgau shows no similarities with Admirable de Courtiller though its genotype could be explained based on microsatellites. Since 5-10 % of the grape cultivars in grapevine collections are known to be not correctly annotated (Dettweller 1992), it might well be possible that a misnaming has occurred. This hypothesis has been addressed by ampelographic and by microsatellite analyses.

## Material and Methods

Herbarized leaves designated as Admirable de Courtiller were obtained from Klosterneuburg, Austria. For visual diagnosis and comparison leaves of true to type cultivars in the herbarium at Geilweilerhof were used and ampelographic literature with ampelographic descriptions and images of the cultivar typus was consulted (GALET 1964, 1990).

For molecular analysis young leaves of 5 cultivars, Madeleine Royale, Riesling White, Müller-Thurgau, Admirable de Courtiller and Silvaner White (all true to type) were collected from the Geilweilerhof collection. DNA samples of each cultivar were prepared according to the protocol of Thomas *et al.* (1993). Microsatellite analysis

was performed at 6 microsatellite loci VVS2 (Thomas and SCOTT 1993), VVMD5, VVMD7, VVMD27 (Bowers et al. 1996), and ssrVrZAG62, ssrVrZAG79 (Sefc et al. 1999). Three-step PCR was performed in 25 µl of reaction volumes using a Gene Amp PCR System 9600 (Perkin Elmer) following the protocol of CRESPAN et al. (1999). The PCR reaction mixtures contained 20-40 ng of template DNA, 1 U Taq DNA Polymerase (Roche Molecular Biochemicals, Mannheim, Germany), buffer (final concentration: 10 mM Tris-HCl, 1.5 mM MgCl, 50 mM KCl, pH 8.3 (20 °C)), 200 μM of each dNTP, 10 pmoles of each primer for the ssrVrZAG loci and 20 pmoles for the VVMD and VVS2 loci, respectively. For the electrophoresis, 2.0 µl of each PCR reaction were denatured at 94 °C for 3 min, loaded on a 8 % polyacrylamide/urea sequencing gel and visualized by silver staining according to Promega and modified according to ECHT et al. (1996). Allele sizes were determined using defined size markers.

## Results and Discussion

It is a well known phenomenon that plant genetic resource collections suffer to a certain degree from misnamings. Worldwide grapevine collections previously analyzed showed 90 –95 % of correct namings (Dettweller 1992). Therefore it is a continuous task to eliminate mistakes in order to maintain reliable collections of genetic resources.

During the course of reviewing herbarized grapevine leaves from various sources, leaves from Klosterneuburg designated as Admirable de Courtiller came across. Ampelographic studies on this material, however, showed no similarity with leaves of the true to type Admirable de Courtiller from Geilweilerhof (Fig. 1). According to its leaf characteristics (strong goffering of the blade around



Fig. 1: Herbarized leaf of the true to type accession of Admirable de Courtiller from Geilweilerhof.

the petiole sinus, 5-lobed, overlapping of the petiole sinus opening, overlapping of the upper leaf sinus, angular tooth shape) it became evident that the designated Admirable de Courtiller was in fact Madeleine Royale (syn. Königliche Magdalenentraube), a widespread table grape bred by a French, Moreau-Robert, in 1845 (Ambrosi *et al.* 1998). The result of the visual inspection according to ampelographic descriptions (Galet 1964, 1990) was supported by microsatellite analysis, using DNA of Madeleine Royale grown in the collection at Geilweilerhof. Thus, the plant

material obtained and designated as Admirable de Courtiller in fact turned out to be the table grape cultivar Madeleine Royale.

This finding results in a new description of the parentage of the grapevine cv. Müller-Thurgau. Despite the fact that the molecular analyses of REGNER et al. (1996) and SEFC et al. (1997) which have been confirmed by GRANDO et al. (1998) are correct, due to a misnaming Müller-Thurgau must be addressed as a cross between Riesling and Madeleine Royale (Fig. 2). This conclusion from ampelographic studies is supported by the allelic profile of Müller-Thurgau obtained at 6 loci (Table). The allele lengths found in Riesling and Madeleine Royale can explain all alleles of Müller-Thurgau, while the differences in both allele lengths at the loci VVS2, VVMD7, VVMD27 and ssrVrZAG79 definitely exclude the true to type Admirable de Courtiller as gene donor for Müller-Thurgau. Although the allele lengths of Madeleine Royale are not identical with the published allele lengths for the misnamed Admirable de Courtiller from Klosterneuburg (SEFC et al. 1997, GRANDO et al. 1998) the data presented are consistent within the experimental variation. It is obvious that the allelic profile of Madeleine Royale corresponds to that of Admirable de Courtiller from Klosterneuburg because only a shift of relative distances, caused by different methods of fragment length determination, is observed between the allelic patterns. This problem is frequently encountered in the interpretation of multiple bands (e.g. stuttering bands) and due to the varying techniques for allele sizing in different laboratories.

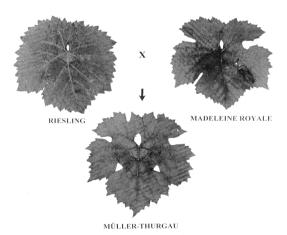


Fig. 2: Herbarized leaves of Müller-Thurgau and its parents, Riesling and Madeleine Royale.

Several phenotypical and phenological similarities between Müller-Thurgau and Madeleine Royale indicate their close relationship, the most prominent being: early ripening, moderate resistance to winter frost, high sensitivity to *Plasmopara viticola* and *Botrytis cinerea*, similarities of leaf anatomy (see above), roundish to slightly-elliptic berry shape. From our studies including molecular markers it can thus be stated that Riesling and Madeleine Royale are the parents of Müller-Thurgau. As a conclusion, there is no doubt that the confirmation of true to

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Parentage analysis at six microsatellite loci for Müller-Thurgau and its presumed parents

VVS2 VVMD5 VVMD7 VVMD27 SERV-ZAG-62

	VVS2	VVMD5	VVMD7	VVMD27	ssrVrZAG 62	ssrVrZAG79	Ref.
Müller-Thurgau	142:150	224:226	244:254		193:193	242:244	(1)
	139:148	222:224	245:255				(2)
	143:151	228:230	247:257	182:182	194:194	243:245	(3)
Riesling White	142:150	224:232	246:254		193:203	242:244	(1)
	139:148	222:230	247:255				(2)
	143:151	228:236	249:257	182:190	194:204	243:245	(3)
Madeleine Royale	151:155	230:238	243:247	182:190	188:194	245:259	(3)
Admirable de Courtiller	150:154	226:234	240:244		187:193	244:258	(1)
(Klosterneuburg)	148:152	224:232	241:245				(2)
Admirable de Courtiller							
(true to type at Geilweilerhof)	133:137	228:238	239:243	186:195	188:194	251:257	(3)
Silvaner White	150:152	224:230	240:244		187:203	248:250	(1)
	148:150	222:228	241:245				(2)
	151:153	228:234	243:247	190:195	188:204	249:251	(3)

- (1) Sefc et al. (1997): Analysis on ALF Express, 2-step PCR
- (2) Grando et al. (1998): Analysis on ABI 310, 3-step PCR
- (3) this publication: Analysis using silver staining, 3-step PCR

typeness of the cultivars in the international grapevine collections remains a serious problem and needs to be considered as a prerequisite for all marker analyses and parentage studies.

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