

MYCOTOXIN-PRODUCING PENICILLIUM AND OTHER FUNGI ISOLATED FROM SLOVAK WINE GRAPES AT VARIOUS STAGES OF MATURATION

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ARTICLE INFO	ABSTRACT
Received 6. 4. 2016 Revised 4. 5. 2016 Accepted 12. 5. 2016 Published 1. 8. 2016 Regular article	The aim of the study was to monitor the mycobiota in grape samples at 3 maturation stages (pea berry, early veraison and ripe berry) and to test the ability of selected <i>Penicillium</i> strains to produce mycotoxins in year 2014. Grapes came from Nitra wine growing region, Šintava subregion and were analyzed by plating methods. From Devín grape variety the 339 strains were detected and identified. The most abundant genera found by descending order were <i>Alternaria</i> (42.8%), <i>Cladosporium</i> (24.2%) and <i>Epicoccum</i> (10.0%). <i>Penicillium expansum</i> was isolated from all 3 maturation stages but in low average relative density (1.2%). From Frankovka modrá (Frankish) grape variety the 562 strains were detected and identified. The most abundant genera found by descending order were <i>Alternaria</i> (30.6%), <i>Cladosporium</i> (21.3%) and <i>Penicillium</i> (20.3%). Totally were identified 8 species of the genus <i>Penicillium</i> . The most important species, on the basis of the isolation from all maturation stages and relative density, was <i>P. expansum</i> . Selected isolates of <i>Penicillium</i> species were tested for their toxigenity, according to thin layer chromatography method. In total 14 <i>Penicillium</i> strains representing 3 potentially toxigenic species were tested for their toxigenic ability, namely <i>P. expansum</i> , <i>P. griseofulvum</i> and <i>P. chrysogenum</i> . Out of 14 strains, 64% produced et least one mycotoxin as revealed by the method used here.

Keywords: Penicillium, wine grapes, maturation stages of grapes, TLC method, mycotoxins

INTRODUCTION

The grape microbial ecosystem is composed of highly diverse microorganisms, including yeasts, bacteria and fungi (**Barata** *et al.*, **2012**). Molds are ubiquitous with various genera commonly found on grapes (**Rosa** *et al.*, **2002**). Mold growth plays an important role in spoilage of grapes, of their pathogenicity to the harvested products, in the physical and chemical stability as well as the sensory properties of the future wine. Fungal spoilage of grapes causes substantial financial losses to growers and processors and may pose a health threat to the consumer if the contaminating fungi produce mycotoxins.

Contamination of grapes by different moulds occurs during preharvesting, harversting and grape processing. The fungal growth begins in grapes if temperature and humidity are suitable. Rotting and spoilage of grape berries before harvest can be caused by a variety of fungal species such as *Alternaria* spp., *Aspergillus* spp., *Botrytis cinerea*, *Cladosporium* spp., *Eurotium* spp., *Penicillium* spp. and *Rhizopus* spp. These genus are regarded as the main natural contaminants of grapes (**Valero** *et al.*, **2005**).

Changes in the total fungal population over the growing season have been observed. At berry set, fungal populations were large in all vineyards, regardless of the grape variety, but the predominant species differed (**Bau** *et al.*, 2005; **Bellí** *et al.*, 2006; **Gomez** *et al.*, 2006; **Melki Ben Fredj** *et al.*, 2007). Harvest was considered to be a critical period in the growing season for fungal development. A decrease in acidity, sugar accumulation, the berry cuticle becoming brittle, damage to berries and an adaptation of mycotoxin-producing fungi to hot and humid environments may account for the abundance of fungi and the high levels of mycotoxins (Clouvel *et al.*, 2008; Lasram *et al.*, 2012).

It is important to identify fungal contaminants in grapes because some moulds can grow and produce mycotoxins while certain yeasts and moulds can cause infections or allergies (**Tournas and Katsoudas, 2005**). The mycotoxin production is characteristic of the species and therefore by identifying the species one can predict potential mycotoxin hazards. The major mycotoxigenic fungi that attack harvested fruits and vegetables are *Aspergillus*, *Penicillium* and *Alternaria* species – and the mycotoxins produced by them in the host tissues, e.g., aflatoxins, ochratoxin A, patulin and alternaria toxins (**Barkai-Golan, 2001**).

Penicillium species are filamentous fungi that are widely distributed in nature and can spoil food and food products (**Samson and Frisvad**, **2004**). Most *Penicillium* spp. are ubiquitous soil fungi that are associated with organic matter in nature, however, their presence as epiphytes on grapes has also been reported. Some species of *Penicillium* are important plant pathogens that cause considerable economic losses in apples, citrus, grapes, and other unrelated crops in different regions of the world (**Donso and Lattore, 2006**). For instance, *P. expansum* Link is an aggressive, destructive postharvest pathogen and a mycotoxigenic species that has been frequently reported as the cause of blue mold in stored apples and pears and causes a large part of the economic losses that occur during storage and shipment (**Barkai-Golan, 2001**). The species of *Penicillium* have gained attention as grapevine pathogens that cause blue mold decay at harvest (**Donso and Latore, 2006**).

The objective of the study was to identify the natural mycobiota in wine grapes collected from the Nitra wine region, with focus on genera *Penicillium*. In the vineyard, samples were taken at the pea-sized berry stage, at the beginning of berry ripening (veraison), and at harvest stage. Special emphasis was laid on the ability of some potentially toxigenic penicillia to produce some selected mycotoxins by thin layer chromatography method (TLC).

MATERIAL AND METHODS

Study area and samples

Slovak republic has 6 distinct wine-growing zones (the Small Carpathians, the Southern Slovak, the Nitra, the Central Slovak, the Eastern Slovak and the Tokaj wine regions). They spread from the west to the east of the country along its southern and south-western borders. We had samples from the Nitra wine region, Šintava subregion, village Báb. Two samples -1 of white grape variety Devín and

1 of blue grape variety Frankovka modrá at 3 maturation stages were mycologically analyzed. One sample of a wine grape variety was represented by three subsamples of wine grapes, which were sampled in left, middle and right part of the vineyard. Samples were collected from July 2014 to October 2014, in the maturation stages corresponding to pea berry (early July), early veraison (middle August) and harvest (middle October). The berries from the vineyards sampled were generally in good condition without visible damage. The samples of wine grapes were taken to sterile plastic bags and transported to a mycological laboratory for immediate processing.

Mycological analysis of grapes

The detection of fungi in grape samples was made by plating methods without disinfection. total 50 surface А of berries (6 - 7 healthy berries per bunch) from each sample were randomly selected and plated on Dichloran Rose Bengal Chloramphenicol agar medium (DRBC, MERCK, Germany), and incubated at 25 °C in the dark for one week. The sporeproducing filamentous fungi detected were identified according to the manuals of Samson et al. (2002a, 2010). Penicillium strains were isolated and cultivated in MEA (Malt extract agar, Samson et al., 2010), CYA (Czapek yeast agar, Samson et al., 2010), CREA (Creatine-Sucrose agar, Samson et al., 2010) and YES (Yeast Extract agar, Samson et al., 2010) to obtain pure cultures and identify further species. Genus Penicillium was identified to species level based on morphological characters according to special mycological literature of Pitt and Hocking (1997), Samson and Frisvad (2004) and Samson et al. (2002a, 2010).

Results evaluation

The obtained results were evaluated and expressed according to relative density (RD). The relative density (%) is defined as the percentage of isolates of the species or genus, occurring in the analyzed sample (**Guatam et al., 2009**). These values were calculated according to **González et al.** (1999) as follows:

RD (%) = (ni / Ni) x 100

ni - number of isolates of a species or genus; Ni - total number of isolated fungi.

Toxinogenity analysis

Toxinogenity of selected isolates was screened in *in vitro* conditions by means of thin layer chromatography (TLC) according to **Samson et al. (2002b)**, modified by **Labuda and Tančinová (2006).** Extracellular metabolites – citrinin, patulin and griseofulvin were carried out on YES agar and intracellural roquefortin C and cyclopiazonic acid on CYA agar. A few pieces of mycelium with approximate size 5 x 5 mm were cut from colonies and placed in an Eppendorf tube with 500 µL of chloroform: methanol – 2:1 (Reachem, Slovak Republic). The content of the tubes was stirred for 5 min. by Vortex Genie ® 2 (MO BIO Laboratories, Inc. – Carlsbad, CA, USA). The volume 30 µL of liquid phase of extracts along with 10 µL standards (Sigma, Germany) was applied on TLC plate (Alugram ® SIL G, Macherey – Nagel, Germany). The plate was put into TEF solvent (toluene: ethyl acetate: formic acid – 5:4:1, toluene – Mikrochem, Slovak Republic; ethyl acetate and formic acid – Slavus, Slovak Republic). After elution the plate was air-dried. Identification of the metabolites was done by comparison with

metabolite standards. Roquefortin C was visible after spraying with Ce(SO₄)₂ x 4 H₂O as an orange spot. Cyclopiazonic acid was visible directly in daylight after spraying with the Ehrlich reagent as a violet-tailed spot. Patulin by spraying with 0,5% methylbenzothiazolone hydrochloride (MBTH, Merck, Germany) in methanol and heating at 130 °C for 8 min and then detectable as a yellow-orange spot. Directly under UV light with a wavelength of 365 nm was visualized citrinin as a yellow-green-tailed spot and griseofulvin as a blue spot.

RESULTS AND DISCUSSION

The filamentous fungi identified in white grape variety Devin from July 2014 to September 2014 by the direct plating method are indicated in Table 1. Without surface disinfection, a total of 339 strains belonging to 7 genera and *Mycelia sterilia* were identified. The three most abundant genera found by descending order were *Alternaria*, *Cladosporium* and *Epicoccum*. *Trichoderma* and *Penicillium expansum* were detected in more than 1% of the berries analyzed. The remaining genera were detected in less than 1% of the berries. *Penicillium expansum* can cause rot in grapes, but does not usually attack grapes before harvest (**Serra** *et al.*, 2005). *Penicillium expansum* was isolated from all 3 maturation stages but in low frequency. Also other studies have identified *P. expansum* as the species most frequently isolated from Portuguese (Abrunhosa *et al.*, 2001) and French (La Guerche et al., 2004; Bejaoui et al., 2006) vineyards.

The mycobiota changed with maturation stage. The most frequent genera by descending order at pea berry were *Alternaria, Cladosporium* and *Epicoccum*. These constitute 79% of the fungi identified. The highest incidence of *Cladosporium* in grape was observed at this stage. *Rhizopus* occurred only at this stage. The fungal genera most frequently isolated at setting were *Alternaria* (**Bau** *et al.*, 2005; Bellí *et al.*, 2006) and *Aspergillus* (Gomez *et al.*, 2006; Melki Ben Fredj *et al.*, 2005; Bellí *et al.*, 2006; Gomez *et al.*, 2006; Melki Ben Fredj *et al.*, 2007).

The changes occurred in the incidence of *Alternaria, Cladosporium, Epicoccum* and *Penicillium expansum* at early veraison. *Alternaria, Epicoccum* and *Penicillium expansum* increased, while the *Cladosporium* incidence decreased. *Mucor* was detected only at this stage. The most frequent genera by descending order were *Alternaria, Cladosporium* and *Epicoccum*, representing 80.5% of the fungi identified at early veraison.

The changes occurred in the incidence of *Alternaria, Cladosporium* and *Penicillium expansum*, which decreased from early veraison to ripe berry. Our results corresponding with **Bau et al. (2005)**, **Bellí et al. (2006)**, **Gomez et al. (2006)**, **Melki Ben Fredj et al. (2007)**. There the percentage of grapes contaminated with *Alternaria* spp. and *Cladosporium* spp. also clearly decreased with the ripening of the grape berries. The most frequent genera by descending order were *Alternaria, Cladosporium* and *Epicoccum*, representing 71.3% of the fungi identified at this stage. *Trichoderma* occurred only at this maturation stage.

Table 1 Fungi identified in white grape variety Devín from July 2014 to September 2014 by the direct plating method

Funci	Number of colonized berries							
Fuligi	Pea berry	RD (%)	Early veraison	RD (%)	Harvest	RD (%)	Total	RD (%)
Alternaria	41	37	58	48.3	46	42.6	145	42.8
Cladosporium	37	33.3	26	21.6	19	17.6	82	24.2
Epicoccum	10	9	12	10.0	12	11.1	34	10.0
Mucor	0	-	3	2.5	0	-	3	< 1
Penicillium expansum	1	< 1	2	1.7	1	< 1	4	1.2
Rhizopus	1	< 1	0	-	0	-	1	< 1
Trichoderma	0	-	0	-	5	4.6	5	1.5
Mycelia sterilia	21	18.9	19	15.8	25	23.1	65	19.2
Total identified fungi	111		120		108		339	

Legend: RD – Relative density

The filamentous fungi identified from blue grape variety Frankovka modrá by the direct plating method are indicated in Table 2. Without surface disinfection, a total of 562 strains belonging to 11 genera and *Mycelia sterilia* were identified. The 4 most abundant genera found by descending order were *Alternaria, Cladosporium, Penicillium* and *Epicoccum. Botrytis cinerea, Fusarium* and *Trichoderma* were detected in more than 1 % of the berries analyzed. *Arthrnium, Geotrichum, Mucor* and *Rhizopus* were detected in less than 1% of the berries. **Abrunhosa et al. (2001)** reported that *Alternaria* and *Cladosporium* were more frequently isolated from red grape varieties than from white grape varieties, regardless of the vineyard in Portugal considered. A total of 114 *Penicillium* isolates were obtained from healthy red cultivares Frankovka modrá, 12 were

obtained at pea berry stage, 23 at early veraison and 79 at ripe berry stage. Based on the results obtained, increasing *Penicillium* populations were found near harvest in agreement with studies **Latorre** *et al.* (2011) and **Díaz** *et al.* (2011), however, the overall *Penicillium* population was considerably less abundant than the populations of *Cladosporium* spp., which are epiphyte fungi commonly found on grapevines (**Latorre** *et al.*, 2011).

A total of 181 fungal isolates, representing 7 genera and *Mycelia sterilia* were identified from pea berry samples. The most frequent genera by descending order at pea berry were *Alternaria* and *Cladosporium*, which constitute 69% of the fungi identified. The highest incidence of *Fusarium* in grape, which was not detected in white grape variety Devín, was observed at this stage. Healthy cluster

of Cabernet Sauvignon grapes obtained *Penicillium* population low in Alhué and Alto Jahuel in Chile (0.006 CFU.cm⁻² berry) at the pea-sized berry stage. They increased to 0.009 and 0.12 CFU.cm⁻² berry at the beginning of berry ripening stage and to 0.016 and 0.013 CFU.cm⁻² of berry at harvest in Alhué and Alto Jahuel, respectively (**Díaz et al., 2011**) what corresponding with our results.

The highest number of isolates (220) represented 7 genera and *Mycelia sterilia* was detected at early veraison. The most frequent genera by descending order were *Alternaria, Cladosporium, Epicoccum* and *Penicillium*, representing 83.6% of the fungi identified at this stage. All of these fungi increased from pea berry to early veraison. The highest incidence of *Alternaria, Cladosporium, Epicoccum, Botrytis cinerea and Trichoderma* was observed at this stage.

The most frequent genera by descending order were *Penicillium*, *Epicoccum*, *Alternaria* and *Cladosporium* at harvest stage, representing 85.8% of the fungi identified. *Penicillium* increased, while *Alternaria*, *Epicoccum* and *Cladosporium* decreased from early veraison to ripe berry. The highest incidence of *Penicillium* was observed at this stage (49.1%). *Arthrinium*, *Geotrichum*, *Mucor* and *Rhizopus* occurred only at maturation stage. At harvest, the most prevalent mycobiota in four French vineyards (Alsace, Beaujolais, Bordeaux and Burgundy) were found to be *Penicillium* species (**Diguta** *et al.*, **2011**). By contrast, in southern vineyards, black aspergilli have been identified as the

predominant mycobiota on grapes at harvest time, although these fungi may be found on the surface of healthy grapes at all stages, regardless of the vineyard considered (Cabañes et al., 2002; Da Rocha Rosa et al., 2002; Sage et al., 2002; Serra et al., 2003; Bellí et al., 2006; Bau et al., 2005; Gomez et al., 2006; Guzev et al., 2006; Leong et al., 2006). The genus Aspergillus is a component of the epiphytic flora of grapes and may be present on grape berries. The surface of infected berries is covered with a black mold that occurs particularly on grapevines growing in warm climates. The symptoms first become evident when lesions are observed on the skin of the berry, they are generally most evident within the grapes after veraison and increase in intensity with berry damage and ripeness (Cabañes et al., 2002; Da Rocha Rosa et al., 2002; Sage et al., 2002; Serra et al., 2003; Bellí et al., 2006; Bau et al., 2005; Gomez et al., 2006; Guzev et al., 2006; Leong et al., 2006; Somma et al., 2012). In the study the genus Aspergillus was not detected in any stages of the grapes growth. The genus Penicillium seems to be more frequent in temperate and cold climates, such as those in northern Europe, whereas Aspergillus is more frequently associated with warmer and wetter regions (Pitt and Hocking, 1997; Battilani et al., 2006; Bejaoui et al., 2006; Serra et al., 2006).

Table 2 Fungi identified in blue grape variety	Frankovka modrá from Jul	y 2014 to September	r 2014 by the direc	t plating method
Mounter	-f 1			

Euroi	Number of color	lized berries						
Fungi	Pea berry	RD (%)	Early veraison	RD (%)	Harvest	RD (%)	Total	RD (%)
Alternaria	75	41.4	80	36.4	17	10.6	172	30.6
Arthrinium	0	-	0	-	1	< 1	1	< 1
Botrytis cinerea	15	8.3	17	7.7	4	2.5	36	6.4
Cladosporium	50	27.6	53	24.1	17	10.6	120	21.3
Epicoccum	7	3.9	28	12.7	25	15.5	60	10.7
Fusarium	9	5.0	5	2.3	3	1.9	17	3.0
Geotrichum	0	-	0	-	1	< 1	1	< 1
Mucor	0	-	0	-	1	< 1	1	< 1
Penicillium	12	6.6	23	10.4	79	49.1	114	20.3
Rhizopus	0	-	0	-	1	< 1	1	< 1
Trichoderma	4	2.2	6	2.7	4	2.5	14	2.5
Mycelia sterilia	9	5.0	8	3.6	8	5.0	25	4.4
Total identified fungi	181		220		161		562	

Legend: RD – Relative density

Penicillium represents 20.3% from microfungi in blue grape variety Frankovka modrá in the year 2014. The spectrum of isolated penicillia consisted of 8 species (Table 3). The most presented species of all isolates (114) was *P. expansum* (73). From this point of view, the relative density was the highest (64%).

Four species were detected at pea berry, namely *P. aurantiogriseum*, *P. brevicompactum*, *P. expansum* and *P. glabrum*. The most presented species of all isolates (12) was *P. expansum* (58.3%).

Six species were present in the veraison time, namely *P. aurantiogriseum*, *P. brevicompactum*, *P. corylophilum*, *P. expansum*, *P. glabrum* and *P. polonicum*. The largest number of isolates belonged to *P. expansum* (10, 43.5% RD), *P. brevicompactum* (5, 21.7% RD) and *P. corylophilum* (3, 13.0% RD). Another isolates were present under 10%.

The genera *Penicillium* was predominant in harvest time (49.1%), and it was represented by *P. aurantiogriseum*, *P. brevicompactum*, *P. corylophilum*, *P. expansum*, *P. glabrum*, *P. griseofulvum* and *P. chrysogenum*. During the harvest time, a number of *Penicillium* strains belonging to species with ability to produce mycotoxins (*P. expansum*, *P. griseofulvum* and *P. chrysogenum*). They were present with a 73.5% relative

density of 79 *Penicillium* strains. *Penicillium verrucosum* and *P. nordicum*, the only confirmed *Penicillium* species that are able to produce ochratoxin A, were not isolated.

Fifty-nine different species of *Penicillium* have been isolated from grapes in vineyards around the world (**Rousseaux** *et al.*, 2014). The predominant species of *Penicillium* isolated from grapes differs between vineyards and vintages. For example, *P. chrysogenum* is the species most frequently isolated in Argentina (**Magnoli** *et al.*, 2003). *Penicillium brevicompactum* has been identified as the *Penicillium* species most frequently isolated from French and Portuguese vineyards (**Sage** *et al.*, 2002, 2004; **Serra** *et al.*, 2005, 2006). **Diguta** *et al.* (2011) identified *P. spinolusum* as the most frequently isolated species of *Penicillium*, followed by *P. expansum* and *P. minioluteum*, for the 2008 vintage, in Burgundy. **La Guerche** *et al.* (2004) identified *P. expansum* as the predominant species, which may generate organoleptic defects, depends on both vineyard and vintage. It is therefore likely to be difficult to generalize the management of filamentous fungus control (**Rousseaux** *et al.*, 2014).

Table 3 Penicillium spo	ecies identified in blue gra	pe variety Frankovka	modrá from Jul	y 2014 to Se	ptember 2014
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Daniaillium anagias	Number of color	nized berries						
Penicilium species	Pea berry	RD (%)	Early veraison	RD (%)	Harvest	RD (%)	Total	RD (%)
P. aurantiogriseum	2	16.7	1	4.3	3	3.8	6	5.3
P. brevicompactum	2	16.7	5	21.7	7	8.9	14	12.3
P. corylophilum	0	-	3	13.0	7	8.9	10	8.8
P. expansum	7	58.3	10	43.5	56	70.9	73	64.0
P. glabrum	1	8.3	2	8.7	4	5.1	7	6.1
P. griseofulvum	0	-	0	-	1	1.3	1	<1
P. chrysogenum	0	-	0	-	1	1.3	1	<1
P. polonicum	0	-	2	8.7	0	-	2	1.7
Total	12		23		79		114	

Legend: RD – Relative density

In total 14 Penicillium strains representing 3 potentially toxigenic species were tested for their toxigenic ability, namely P. expansum, P. griseofulvum and P.

chrysogenum (Table 4). Of all potentially toxigenic strains P. expansum from white grape variety Devín and P. chrysogenum from blue grape variety

Frankovka modrá were positive on tested mycotoxins. A 50% of P. expansum from Frankovka modrá were positive for citrinin, a higher percentage (80%) of isolates were positive for patulin and all were positive for roquefortin C. A small percentage (20%) of P. expansum isolates were positive for patulin on YES agar, whereas they were all positive for citrinin, reported Abrunhosa et al. (2001) from the wine-producing regions of Portugal. Penicillium griseofulvum from Frankovka modrá was positive for griseofulvin, cyclopiazonic acid and roquefortin C but not for patulin. Out of 14 strains, 64% produced et least one mycotoxin as revealed by the method used here. Patulin, produced primarily by P. expansum is a thermal resistant, causes gastrointestinal problems, skin rashes, and is known to be mutagenic, immunologic, and neurotoxic mycotoxin (Abrunhosa et al., 2001). Patulin contaminates apples and apple derivates (Frisvad and Flitenborg, 1989). However, patulin has also been reported in grapes (Moake et al., 2005), processed grape juice and fermenting wine (Majerus et al., 2008). Patulin inhibits the fermenting yeast Saccharomyces cerevisiae in the must. It is partially degraded by the addition of sulfur dioxide and completely degraded during alcoholic fermentation (Ough and Corison, 1980; Díaz et al., 2011). It is therefore unlikely to be present in wine. Citrinin is a nephrotoxic mycotoxin produced by several species of the genera Aspergillus, Penicillium and Monascus. Penicillium citrinum, the main producer of citrinin, was not isolat from our grapes. Citrinin is not degraded during alcoholic fermentation and may be present in very small amounts in wine. However, wine contamination is unlikely, due to the low abundance of citrinin producing species on grapes.

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Penicillium species	Isolated from	С	Р	G	CPA	RC
P. expansum	Devín	2*/2**	2/2			2/2
P. expansum	Frankovka modrá	5/10	8/10			10/10
P. griseofulvum	Frankovka modrá		0/1	1/1	1/1	1/1
P. chrysogenum	Frankovka modrá					1/1

Legend: C – citrinin; P – patulin; G – griseofulvin; CPA – cyclopiazonic acid; RC – roquefortin C; * positive isolates; ** number of tested isolates

CONCLUSION

Two grape varieties Devín and Frankovka modrá (Blue Frankish) were collected in the maturation stages corresponding to pea berry, early veraison and harvest from the Nitra wine region. Samples were mycological analysed with focus on genera Penicillium. The mycobiota changed with maturation stage. The most presented genera by descending order at pea berry and early veraison were Alternaria, Cladosporium and Epicoccum and at ripe berry Alternaria, Cladosporium and Penicillium expansum. Penicillium contamination in all stages types of grapes Devín was lower than in Frankovka modrá grape variety, where 79 isolates of 7 Penicillium species were found. Moulds commonly isolated from grape variety Frankovka modrá were Alternaria, Cladosporium and Penicillium. Results indicate that Penicillium expansum is the major species of Penicillium found on grapevins that can potentionally contaminate grapes with patulin. However, the average relative density of patulin-producing strains of P. expansum in grapes variety Devín was relatively low (1.2%) but in blue grapes higher (20.3%). Three potentially toxigenic species isolated from exogenous mycobiota were tested for their toxigenic ability by thin layer chromatography method, namely P. expansum, P. griseofulvum and P. chrysogenum. Out of 14 strains, 64% produced at least one mycotoxin. In the research, ochratoxigenic microfungi were not found in grape samples. The use of good quality raw materials is essential for mycotoxin control in food products.

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