

Alma Mater Studiorum Università di Bologna Archivio istituzionale della ricerca

Massarilactones D and H, phytotoxins produced by Kalmusia variispora, associated with grapevine trunk diseases (GTDs) in Iran

This is the final peer-reviewed author's accepted manuscript (postprint) of the following publication:

Published Version:

Cimmino A., Bahmani Z., Masi M., Di Lecce R., Amini J., Abdollahzadeh J., et al. (2021). Massarilactones D and H, phytotoxins produced by Kalmusia variispora, associated with grapevine trunk diseases (GTDs) in Iran. NATURAL PRODUCT RESEARCH, 35(23), 5192-5198 [10.1080/14786419.2020.1791116].

Availability:

This version is available at: https://hdl.handle.net/11585/958234 since: 2024-06-17

Published:

DOI: http://doi.org/10.1080/14786419.2020.1791116

Terms of use:

Some rights reserved. The terms and conditions for the reuse of this version of the manuscript are specified in the publishing policy. For all terms of use and more information see the publisher's website.

This item was downloaded from IRIS Università di Bologna (https://cris.unibo.it/). When citing, please refer to the published version.

(Article begins on next page)

Massarilactones D and H, phytotoxins produced by *Kalmusia variispora*, associated with grapevine trunk diseases (GTDs) in Iran

Alessio Cimmino^{a,*}, Zeinab Bahmani^b Marco Masi^a, Roberta Di Lecce^a, Jahanshir Amini^b, Jafar Abdollahzadeh^{b,*} Angela Tuzi^a, and Antonio Evidente^a

^aDipartimento di Scienze Chimiche, Università di Napoli Federico II, Complesso Universitario Monte S. Angelo, Via Cintia 4, 80126 Napoli, Italy; <u>alessio.cimmino@unina.it</u> (A.C), <u>marco.masi@unina.it</u> (M.M.), <u>evidente@unina.it</u> (A.E.)

^bDepartment of Plant Protection, Agriculture Faculty, University of Kurdistan, P.O. Box 416, Sanandaj, Iran. <u>z_bahmani65@yahoo.com</u> (Z.B.), <u>jamini@uok.ac.ir</u> (J.A.), <u>j.abdollahzadeh@uok.ac.ir</u> (J.A.),

CONTACT A. Cimmino ⊠<u>alessio.cimmino@unina.it; j.abdollahzadeh@uok.ac.ir</u>

Supplemental data for this article can be accessed at https://doi.org/10.1080/14786419.2020.1791116.

ABSTRACT

A strain of *Kalmusia variispora* associated with grapevine trunk diseases (GTDs) was identified in Iran and induced disease symptoms on the host in greenhouse conditions. The grapevine pathogens are able to produce a plethora of toxic metabolites belonging to different classes of naturally occurring compounds. Two homogeneous compounds were isolated from the organic extract of *K. variispora* culture filtrates. They were identified by physic (specific optical rotation), and spectroscopic (essentially 1D ¹H and ¹³C NMR and HR ESIMS) methods as the fungal polyketides massarilactones D and H (1 and 2). The unassigned absolute configuration of massarilactone D was unambiguously determined by X-ray diffractometric analysis. Massarilactones D and H showed phytotoxic activity on *Vitis vinifera* L. at two concentrations used and depending from the days of inoculation. Phytotoxicity is also increased when the 3,4,7-O,O',O''-triacetyl derivative of massarilactone D (**3**) was assayed on the host plant. This is the first report on the investigation of phytotoxic metabolites produced by *K. variispora* isolated from infected grapevine in Iran and they seem to be involved in the development of disease symptoms.

KEYWORDS

Grapevine trunk diseases (GTDs); *Kalmusia variispora;* phytotoxins; massarilactone D and H; absolute configuration of massarilactone D

1. Introduction

Grapevine is now cultivated and distributed all around the world and its economic impact is essentially due to the vine production. Thus, the management of abiotic and biotic stresses that strongly affected the specific organoleptic feature and vine production industry has received increased attention. Among biotic stresses the phytopathogenic fungi play a significant role for the negative effects on the quantity and quality of harvested grapevine. Grapevine trunk diseases (GTDs) are among the most important fungal diseases of grape. Botryosphaeria dieback, esca and eutypa dieback are well-known and the most significant grapevine trunk diseases which lead to grape decline and threaten the grape cultivation and wine industry (Bertsch et al., 2013). Esca disease complex the most important grapevine truck disease with five syndromes including brown wood streaking, Petri disease, young esca or grapevine leaf stripe disease (GLSD), esca and esca proper is associated with the tracheomycotic fungal species belong to ascomycetes genera (i.e. Phaeomoniella and Phaeoacremonium) and basidiomycetes fungi such as Fomitiporia and other genera (Mugnai et al., 1999; Wagschal et al., 2008; Andolfi et al., 2011; Bertsch et al. 2013; Fisher et al., 2019). Eutypa lata and its relatives are associated with eutypa dieback (Fallot et al., 1997; Jiménez-Teja et al., 2006; Wagschal et al., 2008; Andolfi et al., 2011; Bertsch et al. 2013), and several species of *Botryosphaeriaceae* are involved in botryosphaeria dieback (Andolfi et al. 2011; Bertsch et al. 2013; Fontaine et al., 2016; Billones-Baaijens and Savocchia 2019). Apart from these well-known and common fungi several different fungal species have been reported in association with grapevine trunk diseases (Jayawardena et al. 2018; Abed-Ashtiani et al. 2019).

These fungi produce a plethora of phytotoxic secondary metabolites the play an important role in the induction of the disease symptoms (Bruno et al., 2007; Evidente et al., 2010; Andolfi et al., 2011; Bertsch et al. 2013). They belong to several different class of natural compounds such as chromanones, cyclohexen epoxide, dihydrofuranes, jasmonic acid ester, naphthalenones, and related compounds and a recent perspective review has been reported on the advanced results on this topic (Masi et al., 2018). During August to November 2016 in a survey on grapevine trunk diseases in vineyards located in Ilam, Kermanshah, Lorestan and Markazi provinces 19 fungal species belong to different group of fungi were isolated from grapes showing general decline symptoms and vascular discoloration and necrosis related to grapevine trunk diseases. Of which, *Kalmusia variispora* (syn. = *Dendrothyrium variisporum*) isolated from infected wood tissues (Figure 1A) was identified based on ITS phylogeny using maximum parsimony analysis and confirmed as pathogen on 2-year old grapevines in greenhouse condition (Figure 1B). Thus far, this species has been reported from Iran and Syria on *Vitis vinifera*, *Vitis* sp., *Rosa hybrid* and *Pinus eldarica* (Farr and Rossman, 2020), but no studies were carried out on the phytotoxic metabolites produced in vitro by *K. variispora*.

This manuscript reports the isolation and the identification of the two main phytotoxic metabolites produced *in vitro* by a strain of *K. variispora* isolated from infected grapevine in Iran. The phytotoxic activity of these compounds on host plant is also reported for the first time as well as the unambiguously assignment of absolute configuration of massarilactone D.

2. Results and discussion

The organic extract of the culture filtrates of *K. variispora* was purified, by combination of column and TLC as detailed described in the Experimental section, to yield two homogeneous compounds identified as massarilactones D and H (**1** and **2**, Figure 2). In particular, **1** was identified by its optical (specific optical rotation) and spectroscopic properties (essentially ¹H NMR and HR ESIMS, see Figures S6 and S7 and Tables S1). However, the absolute configuration (AC) of **1** remained undetermined considering that Kock et al. (2007) assigned to **1** the same AC of massarilactone B (**4**, Figure 1) only on the basis of their specific optical rotation similarity. The AC of **4** was previously assigned by X-ray of its *bis*-(4-bromobenzoate) ester (**5**, Figure 1) when it was isolated together to massarilactone A as an antibiotic metabolite from the culture of the freshwater aquatic fungus *Massarina tunicate* (Oh et al. 2001). **1** gave optical active single crystal by slow evaporation from *i*-PrOH-H₂O (99:1) solution at room temperature, and its absolute configuration was unambiguously assigned by X-ray diffractometric analysis. Massarilactone D (**1**) crystallizes in the monoclinic P $2_12_12_1$ space group with one molecule of **1** and one H₂O solvent molecule contained in the asymmetric unit. Bond lengths and angles are in the normal range and clearly indicate the presence of sp² hybrid carbon atoms at C5, C4a and C7a (Tables S2–S4 of Supplementary Materials). The molecule consists of one six- and one five-membered rings condensed through a double bond at C4a-C7a junction and correspond to a α,β unsaturated lactone. The ring core assumes an almost planar conformation with a degree of flexibility in the six-membered ring that adopts a half-chair conformation with C2 and C3 atoms up and down the mean plane C4/C4a/C7a/O1 (Figures S2 and S3). The hydroxy groups occupy axial positions and are involved in a 3D hydrogen bonding pattern that includes the solvent H₂O molecules in the crystal packing (Figure S4). Four chiral centres are present at C2,C3,C4,C7 and the 2*S*,3*R*,4*S*,7*S* AC was assigned by X-ray analysis accordingly to the literature methods reported to assign the AC in light-atom structures when MoK α radiation is used (Escudero-Adan et al. 2014; Parsons et al. 2013; Parsons 2017). The crystal data are reported in Tables S2–S4.

Massarilactone D (1) was also converted into the 3,4,7-O,O',O''-triacetyl derivative (3) by usual acetylation carried out with pyridine and acetic anhydride. The ¹H NMR of **3** (Table S1, Figures S10 and S11) differed from that of **2** for the downfield shifts ($\Delta\delta$ 0.84 and 0.87) of H-4 and H-3, resonating as two broad singlets at δ 5.17 and 4.69, respectively, and for the presence of a broad singlet at δ 2.1 due to the overlapped signals of the three acetyl groups. Its ESI MS showed the sodiated dimer [2M + Na]⁺ and sodiated [M + Na]⁺ adducts at m/z 759 and 391 and the significant fragmentation ions due to successive losses of two acetic acid moieties [M + H - AcOH]⁺ and [M + H - 2xAcOH]⁺ and a ketene residue [M + H - 2xAcOH - CH₂CO]⁺ at m/z 309, 249, 207.

Massarilactone H (2), obtained as an amorphous solid was identified by comparing its optical (specific optical rotation) and spectroscopic (¹H NMR and ESI MS, see Figures S8 and S9) data with those reported by Zhang et al. (2012).

Massarilactone D (1) was previously isolated from the endophyitic fungus *Coniothyrium* sp. obtained from *Carpobrotus edulis*, together with massarilactone C, massarigenin E, coniothyrenol, graphislactone A and massarilactone A (Kock et al. 2007). When tested at 5 mg/mL for antifungal, antibacterial and antialgal activities, 1 as well as all the other metabolites were inactive against the gram-positive bacterium, *Bacillus megaterium*, the fungus, *Microbotryum violaceum*, and the green alga, *Chlorella fusca* (Kock et al. 2007).

Massarilactone H (2) was previously isolated, together with the new polyketide arthropsadiol C and the already known massarilactones B, E and G, massarigenins C and D, and enalin A, from the culture filtrates of the marine-derived fungus *Phoma herbarum*. Its structure was determined by

single-crystal X-ray analysis of its 3,4-*O*,*O*'-dibenzoate derivative. **2** showed moderate neuraminidase inhibitory activity compared to the positive control oseltamivir (Zhang et al. 2012).

Successively, both massarilactones D and H, were isolated together with the two new furanone derivatives (5*S*)-*cis*-gregatin B and graminin D from the culture filtrates extract of *Dendrothyrium variisporum*, a fungus isolated from the roots of the Algerian plant *Globularia alypum*. Massarilactone D, three new and two known anthranilic acid analogues, and three cyclopeptides were biosynthesized from the same fungus when fermented in large scale (Teponno et al. 2017). Massarilactones D and H when tested for antimicrobial and cytotoxic activities against various bacteria, fungi, and two mammalian cell lines did not showed any biological activity (Teponno et al. 2017). Some pyranfuranones close related to massarilacones D and H, named isoaigialones A, B, and C were recently isolated from the organic extract of a *Phaeoacremonium* sp. culture filtrates, an endophytic fungus obtained from the same organic extract and showed antifungal activity against the phytopathogenic fungi *Cladosporium cladosporioides* and *C. sphaerospermum* (Silva et al. 2017).

Assayed on *Vitis vinifera* L. (Table S5, Figure S5), massarilactones D and H showed phytotoxic effects at two concentrations used $(10^{-3} \text{ and } 10^{-4} \text{ M})$ and depending from the days of inoculation. Phytotoxicity is also increased when the 3,4,7-*O*,*O*',*O*"-triacetyl derivative of massarilactone D (**3**) was assayed on the same host plant. In particular, 72 h after inoculation **3** induced severe necrosis and shriveling of the leaves at both concentrations while a marked shriveling and moderate wilting was observed for both massarilactones D and H depending from concentration. The highest phytotoxicity of **3** in respect to **1** and **2** could be explained by the known lethal metabolism (Hassal 1990). In fact, due to its higher liphofilicity, **3** could more easily translocated across cell membrane and then at physiological pH be hydrolyzed in massarilactone D (**1**).

3. Experimental

The general experimental procedures, the identification data and the ¹H NMR and HRESI MS spectra of compounds **1** and **2**, the preparation of compound **3** and its spectroscopic data, X-ray crystallographic analysis of **1** and the phytotoxic bioassays of **1**–**3** are available as supplementary materials.

3.1. Fungal isolates and culture conditions

The strain of *K. variispora* (BMM-K1) used in this study was obtained from grapevine shoot of a vineyard located in Ostorinan, Lorestan Province, Iran showing decline symptoms and vascular discoloration and necrosis related to grapevine trunk diseases. DNA extraction and identification of the strain BMN-K1 based on ITS sequence data were carried out as described by Abdollahzadeh et al. (2009). The strain BMN-K1 was stored on potato dextrose agar (PDA) at 4 °C in fungal collection of the Department of Plant Protection, University of Kurdistan, Iran. For phytotoxin production it was inoculated and grown in stationary culture of potato dextrose broth (PDB) for 21 days at 25 °C in the dark. The mycelium was removed, and the liquid cultures were lyophilized prior to the extraction procedure.

3.2. Extraction and purification of K. variispora phytotoxins

The culture filtrates (4 L) of *K. variispora* were dissolved in 1/10 of the initial volume (pH 6) and extracted with EtOAc (3 x 400 mL). The organic extracts were combined, dried (Na₂SO₄), and evaporated under reduced pressure giving a corresponding oily residue (370 mg). The resultant aqueous phase was acidified (pH 2.5) with 1M formic acid and extracted as reported above yielding a corresponding oily residue (330 mg). The neutral organic extract was purified by silica gel column chromatography, eluted with CHCl₃-*i*-PrOH (9:1) to (7:3) yielding seven homogeneous fraction groups. The residue of fraction 2 (70 mg) was further purified by TLC eluted with CHCl₃-*i*-PrOH (98:2), resulting in one amorphous solid identified as massarilactone H (**2**, Figure 1, 50 mg). The residue of fraction 3, crystallized from slow evaporation of a *i*-PrOH-H₂O (99:1) solution at room temperature and the crystals were identified by as massarilactone D (**1**, Figure 1, 200 mg). The acidic organic extract was purified by silica gel column chromatography, as reported above yielding a further amount of massarilactone D (90 mg).

4. Conclusions

This manuscript reports for the first time the isolation and identification of the known fungal polyketides massarilactones D and H as the two main phytotoxic metabolites produced *in vitro* by a strain of *K. variispora* isolated from infected grapevine tissues in Iran. Furthermore, the unambiguously assignment of absolute configuration of massarilactone D, determined by X-ray diffractometric analysis, is reported. This is the first report on the investigation of the phytotoxic secondary metabolites produced by *K. variispora* isolated from infected grapevine in Iran.

Acknowledgements

Prof. Evidente is associated to Istituto di Chimica Biomolecolare del CNR, Pozzuoli, Italy. Zeinab Bahmani was supported by University of Kurdistan, Sanandaj and received a grant from Ministry of Science, Research and Technology, Teheran, Iran.

Disclosure statement

No potential conflict of interest was reported by the authors.

ORCID

Alessio Cimmino: 0000-0002-1551-4237 Marco Masi: 0000-0003-0609-8920 Jahanshir Amini: 0000-0003-4777-4073 Jafar Abdollahzadeh: 0000-0001-8642-1401 Angela Tuzi: 0000-0002-2523-5039 Antonio Evidente: 0000-0001-9110-1656

References

- Abdollahzadeh J, Goltapeh EM, Javadi A, Shams-Bakhsh M, Zare R, Phillips A. 2009. *Barriopsis iraniana* and *Phaeobotryon cupressi*: two new species of the Botryosphaeriaceae from trees in Iran. Persoonia. 23:1–8.
- Abed-Ashtiani F, Narmani A, Arzanlou M. 2019. Analysis of Kalmusia variispora associated with grapevine decline in Iran. Eur J Plant Pathol. 154(3):787–799.
- Andolfi A, Mugnai L, Luque J, Surico G, Cimmino A, Evidente A. 2011. Phytotoxins produced by fungi associated with grapevine trunk diseases. Toxins (Basel)). 3(12):1569–1605.
- Bertsch C, Ramírez-Suero M, Magnin-Robert M, Larignon P, Chong J, Abou-Mansour E, Spagnolo A, Clément C, Fontaine F. 2013. Grapevine trunk diseases: complex and still poorly understood. Plant Pathol. 62(2):243–265.
- Billones-Baaijens R, Savocchia S. 2019. A review of Botryosphaeriaceae species associated with grapevine trunk diseases in Australia and New Zealand. Aust Plant Pathol. 48(1):3–18.
- Bruno G, Sparapano L, Graniti A. 2007. Effects of three esca-associated fungi on *Vitis vinifera* L.: IV. Diffusion through the xylem of metabolites produced by two tracheiphilous fungi in the woody tissue of grapevine leads to esca-like symptoms on leaves and berries. Physiol Mol Plant Pathol. 71(1-3):106–124.
- Escudero-Adan EC, Benet-Buchholz J, Ballester P. 2014. The use of Mo Kα radiation in the assignment of the absolute configuration of light-atom molecules; the importance of high-resolution data. Acta Crystallogr B Struct Sci Cryst Eng Mater. 70(Pt 4):660–668.
- Evidente A, Peduto F, Andolfi A, Marchi G, Mugnai L, Surico G. 2010. Fattori di virulenza dei funghi dell'esca. Fitotossine di *Phaeomoniella chlamydospora*, *Phaeocremonium aleophilum* e *Fomitiporia mediterranea*: Produzione e attività biologica. In: Surico G, Mugnai L, editors. Il mal dell'esca della vite: Interventi di ricerca e sperimentazione per il contenimento della malattia. Firenze (Italy): Arsia Regione Toscana; p. 241–260.

- Fallot J, Deswarte C, Dalmayrac S, Colrat S, Roustan JP. 1997. L'Eutypiose de la vigne: isolement d'une molécule synthétisée par *Eutypa lata* et toxique pour la vigne. Compt Rendus Acad Sci III Scie Vie. 320(2):149–158.
- Farr DF, Rossman AY. 2020. Fungal databases, systematic mycology and microbiology laboratory, ARS, USDA. [accessed 2020 May]. nt.ars-grin.gov/fungaldatabases/
- Fischer M, Peighami-Ashnaei S. 2019. Grapevine, esca complex, and environment: the disease triangle. Phytopathol Mediterr. 58:17–37.
- Fontaine F, Pinto C, Vallet J, Clément C, Gomes AC, Spagnolo A. 2016. The effects of grapevine trunk diseases (GTDs) on vine physiology. Eur J Plant Pathol. 144(4):707–721.
- Hassal KA. 1990. Biochemistry and use of pesticides. Weinheim, (Germany): Verlag Chemie; p. 58, 72, 304, 429, and 497.
- Jayawardena RS, Purahong W, Zhang W, Wubet T, Li X, Liu M, Zhao W, Hyde KD, Liu J, Yan J. 2018. Biodiversity of fungi on *Vitis vinifera* L. revealed by traditional and high-resolution culture- independent approaches. Fungal Divers. 90(1):1–84.
- Jiménez-Teja D, Hernández-Galán R, Collado IG. 2006. Metabolites from Eutypa species that are pathogens on grapes. Nat Prod Rep. 23(1):108–116.
- Kock I, Krohn K, Egold H, Draeger S, Schulz B, Rheinheimer J. 2007. New massarilactones, massarigenin E, and coniothyrenol, isolated from the endophytic fungus *Coniothyrium* sp. From *Carpobrotus edulis*. Eur J Org Chem. 2007(13):2186–2190.
- Masi M, Cimmino A, Reveglia P, Mugnai L, Surico G, Evidente A. 2018. Advances on fungal phytotoxins and their role in grapevine trunk diseases. J Agric Food Chem. 66(24):5948–5958.
- Mugnai L, Graniti A, Surico G. 1999. Esca (black measles) and brown wood-streaking: two old and elusive diseases of grapevines. Plant Dis. 83(5):404–418.

- Oh H, Swenson DC, Gloer JB, Shearer CA. 2001. Massarilactones A and B: novel secondary metabolites from the freshwater aquatic fungus *Massarina tunicata*. Tetrahedron Lett. 42(6): 975–977.
- Parsons S. 2017. Determination of absolute configuration using X-ray diffraction. Tetrahedron Asymmetry. 28(10):1304–1313.
- Parsons S, Flack HD, Wagner T. 2013. Use of intensity quotients and differences in absolute structure refinement. Acta Crystallogr B Struct Sci Cryst Eng Mater. 69(Pt 3):249–259.
- Silva GH, Zeraik ML, De Oliveira CM, Teles HL, Trevisan HC, Pfenning LH, Nicolli CPMCM, Young MCM, Mascarenhas YP, Abreu LM, Saraiva AC, et al. 2017. Lactone derivatives produced by a *Phaeoacremonium* sp., an endophytic fungus from *Senna spectabilis*. J Nat Prod. 80(5): 1674–1678.
- Teponno RB, Noumeur SR, Helaly SE, H€uttel S, Harzallah D, Stadler M. 2017. Furanones and anthranilic acid derivatives from the endophytic fungus *Dendrothyrium variisporum*. Molecules. 22(10):1674.
- Wahschal I, Abou-Mansour E, Petit A-N, Clement C, Fontaine F. 2008. Wood diseases of grapevine: A review on eutypa dieback and esca. In: Ait Barka E, Clément C, editors. Plantmicrobe interactions. Kerala (India): Research Signpost, Chapter 16; p. 367–391.
- Zhang GF, Han WB, Cui JT, Ng SW, Guo ZK, Tan RX, Ge HM. 2012. Neuraminidase inhibitory polyketides from the marine-derived fungus *Phoma herbarum*. Planta Med. 78(1):76–78.







2

5, R=H 6, R=**p-**Br-PhCO

Figure 1. The structure of compounds 1–5.

1, R=H 3, R=*p-*Br-PhCO 4, R=Ac

.