



Article

First Report of *Nigrospora* Species Causing Leaf Spot on Olive (*Olea europaea* L.)

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Abstract: Leaf spot symptoms were spotted in two olive orchards in Istria and in Kvarner Gulf, Croatia. Fungal species from three representative isolates (P13 LECIII, R18 BI, JA20 NP) have been morphologically characterized based on the colony and conidial characteristics. Several techniques were performed for inducing the sporulation of the JA20 NP isolate. Only PDA + banana medium was successful. PCR was conducted for ITS, TUB, and EF1 α gene regions. Phylogenetic analyses were performed using internal transcribed spacer, beta-tubulin, and translation elongation factor 1-alpha sequence data. Three types of tests were conducted: a pathogenicity test on detached leaves, on detached and scratched leaves, and on olive seedlings. Ultimately, from the morphological characterizations, DNA sequence analysis of ITS, TUB, and EF1 α gene regions, and phylogenetic analysis, these species were identified as *Nigrospora gorlenkoana* Novobr., *Nigrospora osmanthi* Mei Wang & L. Cai, and *Nigrospora philosophiae-doctoris* M. Raza, Qian Chen & L. Cai. This is the first report of *Nigrospora* species causing leaf spot on olive trees and the first report of *Nigrospora philosophiae-doctoris* as a plant pathogen. Fungal leaf diseases in conditions that are favorable for infection and disease development can lead to a decrease in the yield and olive oil quality. Therefore, it is necessary to conduct further research and the monitoring of fungal leaf diseases.

Keywords: fungal disease; isolate; olive tree; pathogenicity



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1. Introduction

The olive tree, *Olea europaea* L., is among the world's most important crops. At present, the approximate production levels per year are 23.0 million tons of olives and 3.0 million tons of oil [1]. The crop is indigenous to the Mediterranean region with a mild, rainy winter and a hot, dry summer [2]. In the Republic of Croatia, the production area includes from the north of the Istrian peninsula, the Kvarner Gulf, and the Dalmatia coastal belt with the islands to the south. In recent years, interest in olive oil production has been rising due to certain socio-economic factors, such as a higher demand for olive oil, the possibility of achieving higher prices, and tourism development.

Among all fungal pathogens affecting olives, *Venturia oleaginea* (Castagne) Rossman & Crous and *Pseudocercospora cladosporioides* (Sacc.) U. Braun are two of the most important pathogens causing leaf spots: peacock spot disease (syn. bird's-eye spot, olive leaf scab, olive leaf spot) and cercosporiosis (syn. cercospora leaf spot) [3]. Olive leaf spots caused by *V. oleaginea* and *P. cladosporioides* result in the defoliation of leaves and weakness or death of branches, a reduced fruit set, and a decrease in the oil yield in the following years [4–8].

Other fungal causal agents of leaf spot symptoms on olives are *Alternaria alternata* (Fr.) Keissl. [9], *Colletotrichum acutatum* J.H. Simmonds and *C. gloeosporioides* (Penz.) Penz. & Sacc. [10], *Neofabraea kienholzii* (Seifert, Spotts & Levesque) Spotts, Levesque & Seifert, and *Phylctema vagabunda* Desmazières [11].

Leaf spot usually manifests as small, circular-to-elliptical spots on the leaves. Spots are initially green or yellow-green but gradually turn brown or black as the disease progresses. They often have a dark border and may have a yellow halo around them. In severe cases, the spots can merge, leading to significant defoliation. Leaf spot fungi usually thrive in warm and humid conditions. Rain or overhead irrigation can facilitate the spread of the disease by splashing fungal spores onto healthy leaves. The disease exhibits its highest activity during the wetter months of the year. While leaf spot primarily affects the leaves, a severe infection can weaken the tree and reduce its overall vitality. In severe cases, defoliation can expose fruit to direct sunlight, leading to sunburn and reduced fruit quality. Leaf spot diseases also weaken trees by interrupting the process of photosynthesis. This means the tree produces less energy, which can lead to stunted growth and smaller and less flavorful olives. This can be problematic for olive growers who rely on high-quality fruit for oil production or table olives. Leaf spot diseases can also affect the appearance of olive trees, which may be a concern for growers who want their orchard to be visually appealing. Management practices, such as maintaining good tree spacing and air circulation, pruning, avoiding excessive moisture on the leaves, proper disposal of infected leaves and pruning debris, planting resistant cultivars, etc., can help to maintain the health and productivity of olive orchards.

This study focuses on fungal species from the *Nigrospora* genus, new causal agents of a leaf spot symptom on olives. The *Nigrospora* genus has been introduced in 1902. for *N. panici*, which was isolated as an endophyte from leaves of *Panicum amphibium* in Java, Indonesia [12]. Based on its conidial characteristics, *Nigrospora* was placed in *Dermateaceae* (*Moniliales*) by Barnett and Hunter [13]. Kirk et al. [14] assigned *Nigrospora* and its *Khuskia* sexual morph to *Trichosphaeriaceae* (*Trichosphaeriales*). Wang et al. [15] placed the *Nigrospora* species in the family *Apiosporaceae* based on the phylogenetic analyses of combined ITS, TUB, and EF1- α sequence data of 165 strains from China and Europe [16].

Currently, there are 45 records of *Nigrospora* species in the MycoBank database, namely *Nigrospora aerophila*, *N. arundinacea*, *N. aurantiaca*, *N. bambusae*, *N. brasiliensis*, *N. camelliae-sinensis*, *N. canescens*, *N. chinensis*, *N. cooperae*, *N. covidalis*, *N. endophytica*, *N. falsivesicularis*, *N. gallarum*, *N. globosa*, *N. globospora*, *N. gorlenkoana*, *N. gorlenkoanum*, *N. gossypii*, *N. guangdongensis*, *N. guilinensis*, *N. hainanensis*, *N. javanica*, *N. lacticolonia*, *N. macarangae*, *N. magnoliae*, *N. manihoticola*, *N. maydis*, *N. musae*, *N. oryzae*, *N. osmanthi*, *N. padwickii*, *N. panici*, *N. pernam-bucoensis*, *N. philosophiae-doctoris*, *N. pyriformis*, *N. rubi*, *N. sacchari*, *N. sacchari-officinarum*, *N. saccharicola*, *N. singularis*, *N. sphaerica*, *N. vesicularifera*, *N. vesicularis*, *N. vietnamensis*, and *N. zimmermanii* [17].

Until now, only *Nigrospora oryzae* was isolated from olive trees [18,19], but it was not described as a pathogen on olives.

The aims of this research were to determine the causal agent of leaf spot symptoms on olive trees in Croatia, to morphologically characterize the fungal species from representative isolates, to molecularly identify the isolates of phytopathogenic fungal species using PCR and DNA sequence analysis of ITS, TUB, and EF1 α gene regions, and to determine isolate pathogenicity in pathogenicity tests conducted in the laboratory and in a greenhouse experiment.

2. Materials and Methods

2.1. Collection of Plant Materials and Fungal Isolations

During the summer and autumn of 2021, leaf spot symptoms were spotted on olive trees on the coastal belt in Jadranovo, Kvarner Gulf (45°13'46" N, 14°36'40" E), and in olive orchards in Vabriga (45°17'14" N, 13°36'41" E) and Španidiga (45°03'02.2" N, 13°42'43.9" E) in Istria, Croatia (Figure 1).

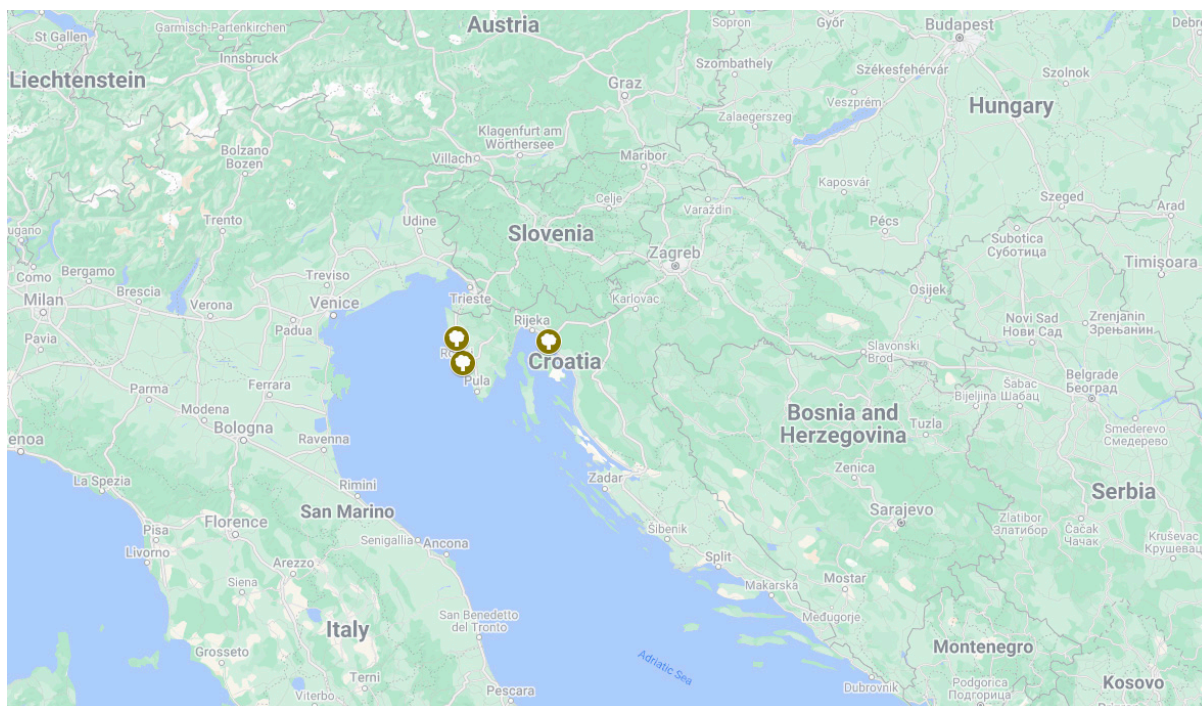


Figure 1. Locations of collected samples: Vabriga, and Špainidiga in Istria, and Jadranovo in Kvarner Gulf.

The symptoms were yellow and brown spots on leaves and defoliation. The samples from symptomatic trees were collected (30 leaves from each tree) in a sterile plastic bag, placed in a portable refrigerator at +4 °C, and immediately brought to the Laboratory for Plant Protection at the Institute of Agriculture and Tourism in Poreč (Croatia) for analysis. Olive varieties on which samples were collected in Istria were ‘Buža’ and ‘Leccino’. The olive variety on which samples were collected in Kvarner Gulfs remains unknown.

Fresh olive leaves were used to isolate the causal agent of yellow and brown spots on leaves. For surface sterilization, leaves were rinsed under tap water for 1 min and transferred to aseptic conditions. Leaves were submerged in 70% ethanol for two minutes, rinsed in sterile distilled water, and placed on a sterile paper sheet in a laminar flow cabinet to surface-dry. Entire leaves were plated on potato dextrose agar (PDA) supplemented with 35 mg/L of penicillin and incubated at 25 °C under dark conditions. After seven days of incubation, the isolates were transferred into the fresh PDA medium for pure culture.

2.2. Morphological Characterization

After 7 and 30 days of incubation at 28 °C in dark conditions, pure fungal cultures were taken for examination. Fungal species, from three representative isolates (P13 LECIII, R18 BI, JA20 NP), have been characterized based on the colony characteristics (color, form, elevation, margin, surface, and opacity) and conidial characteristics (color, shape, presence or absence of septum, dimensions). A Boeco BM-2000 microscope, Boeco BCAM10 camera, and B-View software (Boeckel + Co (GmbH + Co), Hamburg, Germany) were used to capture conidia and hyphae. Isolate JA20 NP did not sporulate on PDA. Several techniques were performed for inducing the sporulation of the JA20 NP isolate. Pine needle medium was prepared according to Su et al. [20]. The banana peel technique was performed based on Kindo et al. [21]. The PDA + banana medium was prepared based on a technique for the pine needle medium described in Su et al. [20], by putting 100 g of fresh banana peel (instead of pine needle) and 20 g of potatoes into 1 L of distilled water, boiling for 30 min, filtrating, and keeping the volume at 1 L by adding distilled water. After filtration, it was amended with 20 g of agar and autoclaved for 20 min at 121 °C.

2.3. DNA Extraction, Amplification, and Sequencing

Fresh fungal mycelia of fungal isolates grown on PDA for 5 days, at 28 °C, under dark conditions, were scraped with a sterile laboratory needle from the colony margins and used for genomic DNA (deoxyribonucleic acid) extraction. Total DNA from the isolate was extracted using the Extract-N-Amp™ Plant PCR kit (Sigma-Aldrich, Merck, Saint Louis, MO, USA) according to manufacturer's protocol. The PCR (polymerase chain reaction) amplification process was performed using ITS1/ITS4 [22], ITS5/ITS4 [23], Btub2Fd/Btub4Rd [24], and EF1-728F/EF1-986R [25] pairs of primers (Table 1). The PCR reaction mixture was composed of 12.5 µL of EmeraldAmp® GT PCR Master Mix, 0.5 µL (10 µM) of each primer, 6.5 µL of nuclease-free water, and 5 µL (4.5 ng/µL) of genomic DNA. The PCR was conducted in a SureCycler 8800 Thermal Cycler (Agilent Technologies, Santa Clara, CA, USA) using different PCR conditions for ITS, TUB, and EF1α gene regions (Table 2). In the case of the R18 BI isolate, amplification of the EF1-alpha region was unsuccessful. Therefore, a second PCR was conducted using 1 µL of the initial PCR amplification as the template. Electrophoresis was performed using 1% agarose gel amended with two drops of GelRed® Nucleic Acid Gel Stain (Biotium, Fremont, CA, USA) at 110 V for 30 min in 1x TAE buffer with a BIO-RAD Power Pac 300 electrophoresis power supply (Agilent, Santa Clara, CA, USA). After electrophoresis, the PCR products were visualized using an iBright CL1000 Imaging System (Invitrogen, Thermo Fisher Scientific, Waltham, MA, USA). Purification of PCR products was performed with the GenElute™ PCR Clean-Up Kit (Sigma-Aldrich®, Burlington, MA, USA).

Table 1. List of the primers used for PCR and sequencing.

Locus	Primer	Sequence (5'-3')
Internal transcribed spacer (ITS)	ITS1	5' TCCGTAGGTGAACCTGCGG 3'
	ITS4	5' TCCTCCGCTTATTGATATGC 3'
	ITS5	5' GGAAGTAAAAGTCGTAACAAGG 3'
Beta-tubulin	Btub2Fd	5' AACATGCGTGAGATTGTAAGT 3'
	Btub4Rd	5' TAGTGACCCTTGGCCAGTTG 3'
Translation elongation factor 1-alpha	EF1-728F	5' CATCGAGAAGTTCGAGAAGG 3'
	EF1-986R	5' TACTTGAAGGAACCCTTACC 3'

Table 2. PCR amplification program for ITS and EF1α region sets, according to White et al. [22], and for ITS and TUB region sets, according to Hao et al. [26].

ITS1/ITS4 and EF-728F/EF-986R						
HOT START 95 °C	Start Cycle	Denaturation 95 °C	Annealing 58 °C	Elongation 72 °C	End Cycle	Elongation 72 °C
3 min	34 times	30 s	30 s	1 min		10 min
ITS5/ITS4, and Btub2Fd/Btub4Rd						
HOT START 95 °C	Start Cycle	Denaturation 95 °C	Annealing 58 °C	Elongation 72 °C	End Cycle	Elongation 72 °C
3 min	34 times	30 s	30 s	1 min		10 min

2.4. DNA Sequence Assembly and Phylogenetic Analysis

Sequencing of the PCR products was performed by MacroGen Europe (Amsterdam, The Netherlands). Sequences were edited in Sequencher® (Gene Codes Corporation, Ann Arbor, MI, USA) and compared with sequences from GenBank®.

The phylogenetic trees were constructed and the evolutionary history of the isolated fungi was concluded based on the Neighbor-joining method [27]. Phylogenetic analysis was

performed using ITS, TUB, and EF1 α sequence data from isolates and relevant sequence data of *Nigrospora* and *Botryosphaeria dothidea* (outgroup) isolates from GenBank[®] (a list of isolates used is presented in Table 3). The sequences were aligned using ClustalX2 (UCD Dublin, Dublin, Ireland) software, and a phylogenetic tree was made using MEGA11 software (Pennsylvania State University, State College, PA, USA).

Table 3. Genbank accession numbers of isolates used for phylogenetic analysis based on research carried out by Chen et al. [28].

Species	GenBank Accession Number			References
	ITS	TUB	TEF1 α	
<i>Botryosphaeria dothidea</i>	AY236949	AY236927	AY236898	[29]
<i>Nigrospora aurantiaca</i>	KX986064	KY019465	KY019295	[15]
	MN215771	MN329935	MN264010	[30]
<i>N. bambusae</i>	KY385307	KY385319	KY385313	[15]
	KY385306	KY385320	KY385314	[15]
<i>N. brasiliensis</i>	KY569629	MK720816	MK753271	[31]
	KY569630	MK720817	MK753272	[31]
<i>N. cameliae-sinensis</i>	KX985986	KY019460	KY019293	[15]
	MN215775	MN329939	MN264014	[30]
<i>N. chinensis</i>	KX986023	KY019462	KY019422	[15]
	KX986026	KY019548	KY019445	[15]
<i>N. covidalis</i>	OK335209	OK431479	OK431485	[28]
	OK335210	OK431480	OK431486	[28]
<i>N. falsivesicularis</i>	MN215778	MN329942	MN264017	[30]
	MN215779	MN329943	MN264018	[30]
<i>N. globospora</i>	OK335211	OK431481	OK431487	[28]
	OK335212	OK431482	OK431488	[28]
<i>N. gorlenkoana</i>	KX986048	KY019456	KY019420	[15]
<i>N. guilinensis</i>	KX985983	KY019459	KY019292	[15]
	KX986063	KY019608	KY019404	[15]
<i>N. hainanensis</i>	KX986091	/	KY019415	[15]
	MN215780	MN329944	MN264019	[30]
<i>N. lacticolonina</i>	KX985978	KY019458	KY019291	[15]
	/	MN329948	MN264023	[30]
<i>N. musae</i>	KX986076	KY019455	KY019419	[15]
	KX986042	KY019567	KY019371	[15]
<i>N. oryzae</i>	KX985931	KY019601	KY019396	[15]
	KX985954	KY019481	KY019307	[15]
<i>N. osmanthi</i>	KX986010	KY019461	KY019421	[15]
	KX986017	KY019540	KY019438	[15]
<i>N. philosophiae-doctoris</i>	OK335213	OK431483	OK431489	[28]
	OK335214	OK431484	OK431490	[28]

Table 3. Cont.

Species	GenBank Accession Number			References
	ITS	TUB	TEF1 α	
<i>N. pyriformis</i>	KX985940	KY019457	KY019290	[15]
	MN215787	MN329988	MN264026	[30]
<i>N. rubi</i>	KX985948	KY019475	KY019302	[15]
<i>N. sacchari-officinarum</i>	MN215791	MN329954	MN264030	[30]
	MN215792	MN329955	MN264031	[30]
<i>N. saccharicola</i>	MN21578	/	MN264027	[30]
	MN215789	MN329952	MN264028	[30]
<i>N. singularis</i>	MN215793	MN329956	MN264032	[30]
	MN215794	MN329957	MN264033	[30]
<i>N. sphaerica</i>	KX985965	KY019492	KY019318	[15]
	MN215811	MN329974	MN264050	[30]
<i>N. vesicularifera</i>	MN215812	MN329975	MN264051	[30]
	MN215814	MN329977	MN264053	[30]
<i>N. vesicularis</i>	KX986088	KY019463	KY019294	[15]
	KX985939	KY019467	/	[15]
<i>N. zimmermanii</i>	KY385309	KY385317	KY385311	[15]
	MN215824	MN329987	MN264063	[30]

2.5. Pathogenicity Test

Three pathogenicity tests were conducted to determine the isolates' pathogenicity: pathogenicity test on detached leaves, pathogenicity test on detached and scratched leaves, and pathogenicity test on olive seedlings.

2.5.1. Pathogenicity Test on Detached Leaves and Scratched Detached Leaves

In May 2022, healthy olive leaves of the cultivars Leccino and Buža were collected from a collection orchard at the Institute of Agriculture and Tourism in Poreč (Istria, Croatia). Leaves were washed with tap water, surface-sterilized in 1% sodium hypochlorite solution for three minutes, rinsed with sterilized distilled water for one minute, and placed in a laminar flow cabinet, on sterile paper, until dry. After air-drying, ten leaves of each cultivar (per isolate), scratched with a needle, and ten unscratched leaves of each cultivar (per isolate), were inoculated by placing a 5 mm-diameter mycelium plug, taken from the margins of a five-day-old PDA culture of the isolates. The same numbers of scratched and unscratched leaves inoculated using pure PDA agar plugs were used as controls. Leaves were placed on a sterile filter paper, sprayed with sterile distilled water, in a Petri dish, and protected with parafilm. Two replicates were performed. Leaves were incubated at 28 °C, in dark conditions. After nine days, samples were analyzed.

2.5.2. Pathogenicity Test on Olive Seedlings

For the whole plant assay, the fungus was grown on PDA at 28 °C for five days, and mycelia were collected. Approximately one gram of mycelia was grinded in 10 mL of sterilized distilled water. The grinded mycelia were homogenized by mixing with a vortex mixer. In the greenhouse of the Institute of Agriculture and Tourism, the grinded mycelia were injected into the petiole of the leaves of three-year-old olive seedlings of cultivar Rosinjola. Thirty leaves (ten per isolate) were inoculated with a grinded mycelium. An equal number of control leaves were inoculated with sterile distilled water to serve as a

negative control. Two replicates were performed. The inoculated plants were kept in a greenhouse for two weeks. After the incubation period, samples from all tests were collected and, in an effort to adhere to Koch's postulate, small sections of necrotic tissue from the periphery of lesions were excised and placed on PDA to isolate the inoculated fungus.

3. Results

3.1. Field Survey and Disease Symptoms

Symptoms of leaf spot were observed on mature 10- to 38-year-old trees in Istria, Croatia. Leaves were dry and yellowish to chocolate-brown in color (Figure 2). Some of them had brown spots. Defoliation was observed on leaves with expanded spots, and symptoms affected only part of the trees.



Figure 2. Symptoms on olive leaves, (a) *Nigrospora gorlenkoana*, (b) *Nigrospora osmanthi*, (c,d) *Nigrospora philosophiae-doctoris*.

3.2. Molecular Phylogenetic Identification

For molecular identification, consensus sequences of isolates were produced and registered in GenBank®. GenBank® accession numbers for each isolate and its genome region are represent in Table 4.

Table 4. GenBank accession numbers of the sequences.

SPECIES	ISOLATE	COLLECTION DATE	Varieties	Genbank Accession Number		
				ITS	TUB	Ef1 α
<i>Nigrospora gorlenkoana</i>	P13 LECIII	24 September 2021	Leccino	OP999642	OQ286068	OQ286069
<i>Nigrospora osmanthi</i>	JA20 NP	31 October 2021	Unknown	OP999639	OQ275027	OQ275028
<i>Nigrospora philosophiae-doctoris</i>	R18 BI	14 October 2021	Buža	OP999644	OQ286067	OQ286066

BLAST analysis of the sequences from the P13 LECIII isolate showed 100% similarity for ITS and TUB and 99.27% similarity for the EF1 α gene region to *N. gorlenkoana*. BLAST

analysis of the sequences from the JA20 NP isolate showed 100% similarity for ITS, TUB, and EF1 α gene regions to *N. osmanthi*. BLAST analysis of the sequences from the R18 BI isolate showed 100% similarity for ITS and TUB and 98.64% similarity for the EF1 α gene region to *N. philosophiae-doctoris*. Phylogenetic trees were constructed by aligning ITS, TUB, and EF1 α sequences, and the evolutionary history was inferred using the Neighbor-Joining method [27]. Finally, a multilocus tree was created from a combination of ITS, TUB, and EF1 α sequence alignments. The optimal trees are displayed in Figures 3–6. Beneath the branches are the percentages of replicate trees where related taxa clustered together in the bootstrap test based on 1000 replicates [32]. The Maximum Composite Likelihood method [33] was used to calculate evolutionary distances, expressed in the units of base substitutions per site. The *Botryosphaeria dothidea* isolate CMW8000 was used as the outgroup. Ambiguous positions were excluded via pairwise detection using MEGA11 software [34].

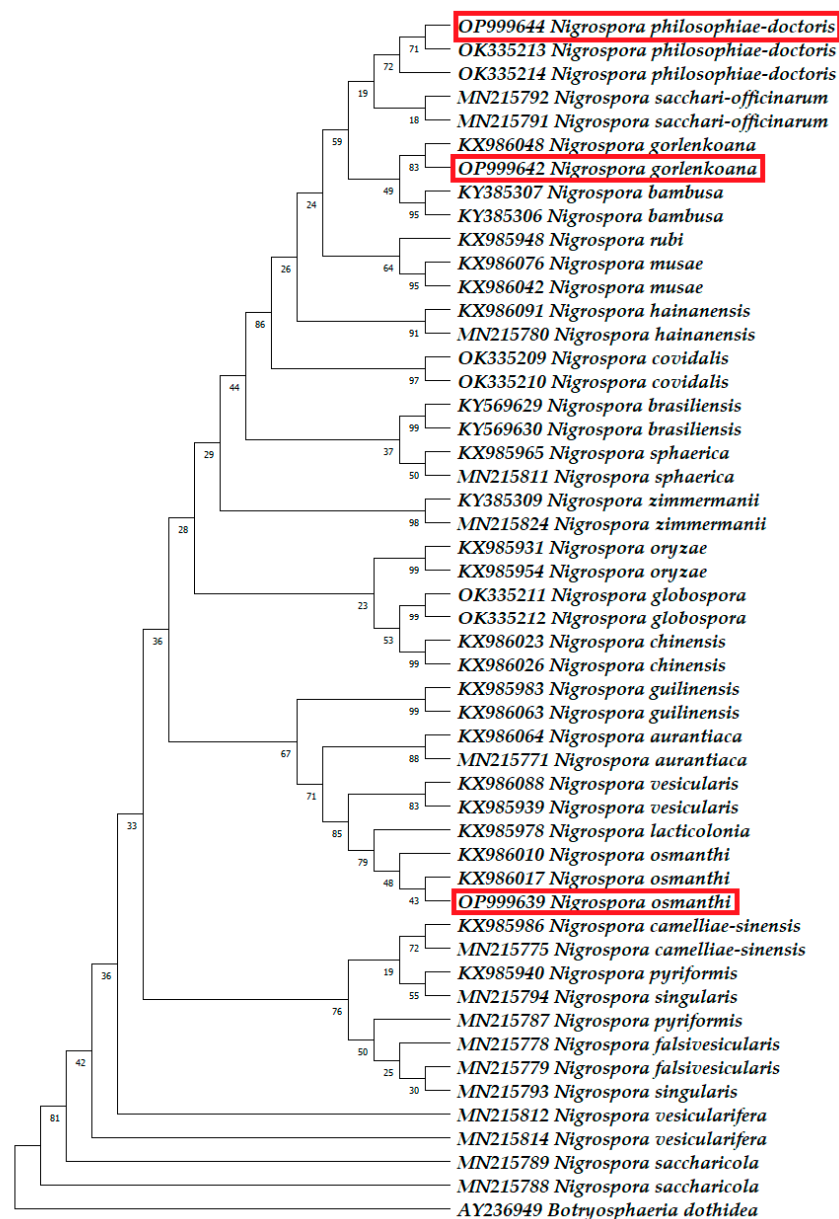


Figure 3. Phylogenetic tree based on internal transcribed spacer sequence alignment. Sequences identified in this research are highlighted with red rectangles. This analysis encompassed 51 nucleotide sequences, resulting in a final dataset comprising 595 positions.

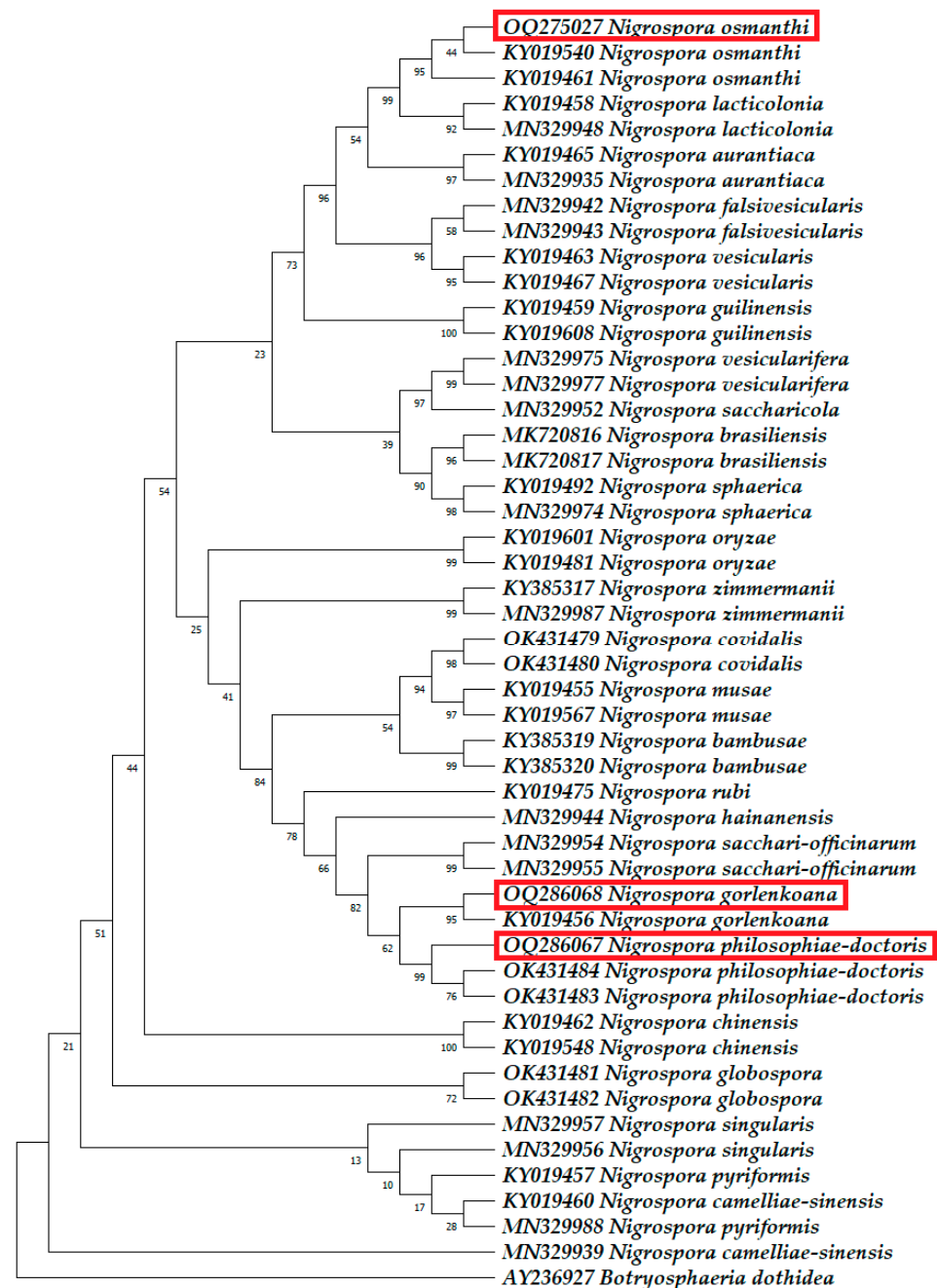


Figure 4. Phylogenetic tree based on beta-tubulin sequence alignment. Sequences identified in this research are highlighted with red rectangles. This analysis encompassed 50 nucleotide sequences, resulting in a final dataset comprising 808 positions.

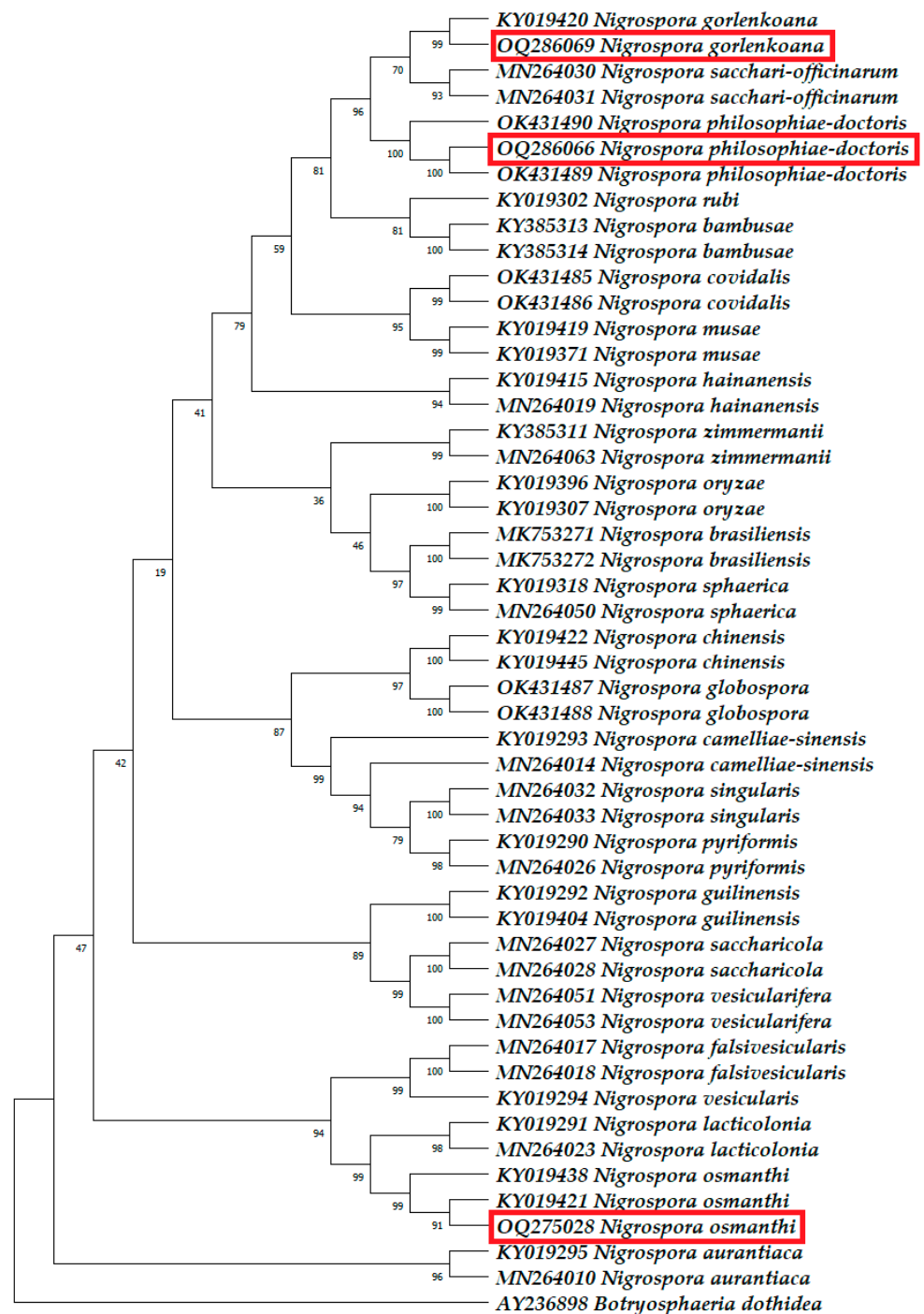


Figure 5. Phylogenetic tree based on translation elongation factor 1-alpha sequence alignment. Sequences identified in this research are highlighted with red rectangles. This analysis encompassed 51 nucleotide sequences, resulting in a final dataset comprising 567 positions.

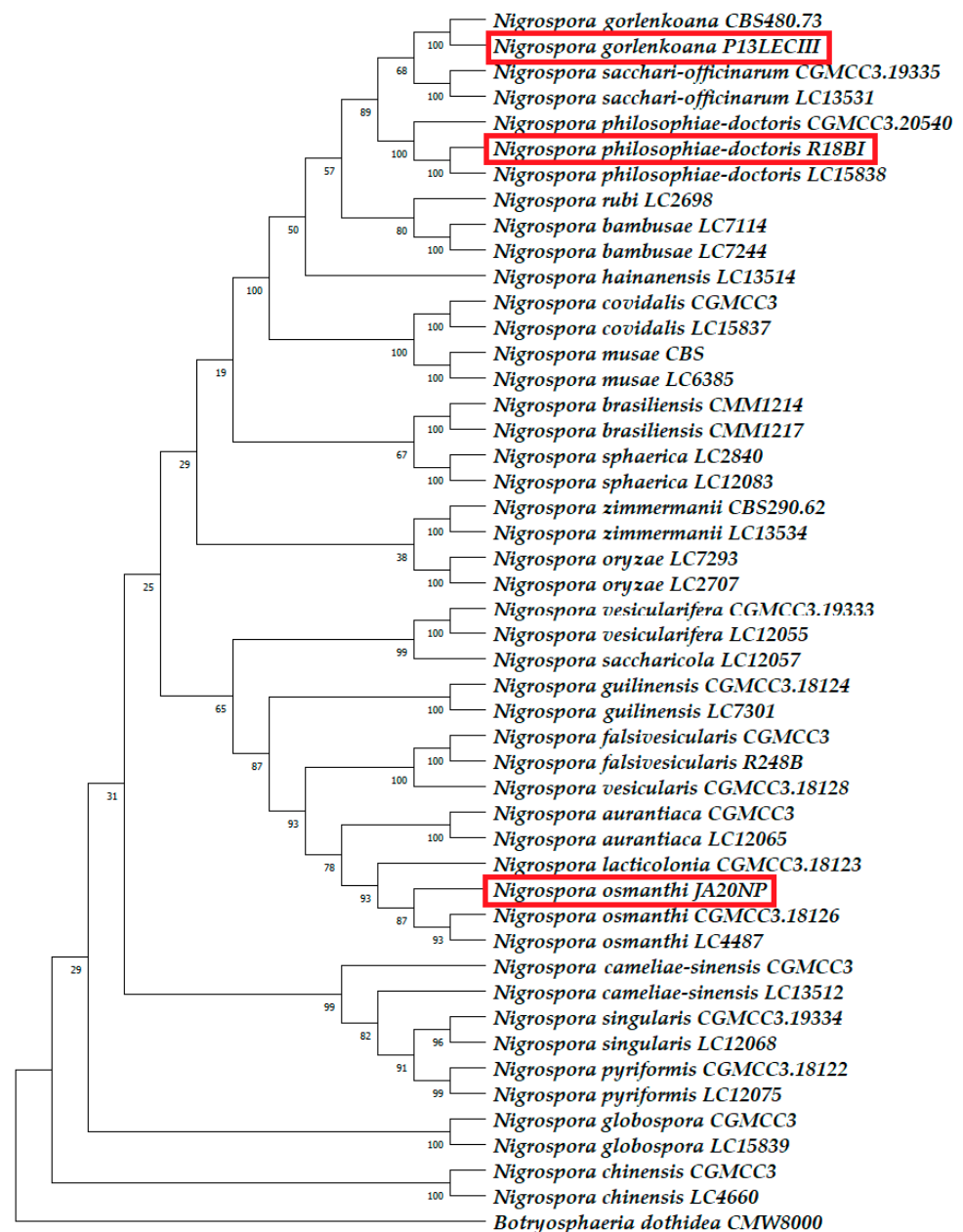


Figure 6. Multilocus tree based on internal transcribed spacer, beta-tubulin, and translation elongation factor 1-alpha sequence alignment. Sequences identified in this research are highlighted with red rectangles. This analysis encompassed 48 nucleotide sequences, resulting in a final dataset comprising 1581 positions.

Ultimately, from the DNA sequence analysis of ITS, TUB, and EF1 α gene regions and the phylogenetic analysis, these species were identified as *Nigrospora gorlenkoana* Novobr., *Nigrospora osmanthi* Mei Wang & L. Cai, and *Nigrospora philosophiae-doctoris* M. Raza, Qian Chen & L. Cai.

3.3. Morphological Characterization and Fungal Incidence

Regarding sporulation, successful sporulation of the JA20 NP isolate was observed exclusively when it was cultured on a PDA + banana medium, as shown in Table 5.

Table 5. List of techniques used for inducing sporulation of the JA20 NP isolate and results.

TECHNIQUES									
PDA Temperatures: 22 °C, 25 °C, 28 °C, 30 °C	1/2 strength PDA medium	WA Temperatures: 22 °C, 25 °C, 28 °C, 30 °C	Pine needle extracts + WA	MEA Temperatures: 22 °C, 25 °C, 28 °C, 30 °C	Host tissue	Slide culture	Exposure to near- ultraviolet light (12 h day/12 h night)	Banana peel	PDA + banana medium
x	x	x	x	x	x	x	x	x	✓

x—sporulation not determined, ✓—sporulation recorded, WA—water agar, MEA—malt extract agar.

3.3.1. *Nigrospora gorlenkoana*

Colonies on PDA had reached a nine-centimeter diameter after two days at 28 °C and on WA after five days and sporulated after three days of incubation on PDA medium. Colonies of *N. gorlenkoana* developing on PDA (Figure 7a,b) were circular-shaped with aerial, woolly mycelium, entire-margined, opaque, floccose, raised with fuzzy edges, and growing rapidly; at first, they were white, becoming light grey when they matured and the reverse initially white, becoming darker grey when they matured. Conidia of *N. gorlenkoana* were round-shaped, light-brown-to-black-colored and aseptated, solitary, smooth, and 10.5–13.8 × 13.5–17.3 µm in diameter (\bar{x} = 11.9 × 15.1 µm, n = 30). Hyphae were smooth, septate, hyaline, and yellowish. On WA, colonies were white and growing poorly.

3.3.2. *Nigrospora osmanthi*

Colonies on PDA had reached a nine-centimeter diameter after three days at 28 °C and on WA after seven days and sporulated after five days of incubation on PDA + banana medium. Colonies of *N. osmanthi* developing on PDA (Figure 7c,d) were circular-shaped with an aerial, slightly woolly mycelium, entire-margined, opaque, raised a little, filiform, and growing rapidly; they were creamy-white-colored up and the reverse becoming greyish when mature. Conidia of *N. osmanthi* were round-shaped, black-colored, and aseptated, solitary, smooth, and 11.2–12.8 × 12.7–15.4 in diameter (\bar{x} = 12.5 × 15.3 µm, n = 30). Hyphae were smooth, septate, hyaline, and yellowish. On WA, colonies were white and growing poorly.

3.3.3. *Nigrospora philosophiae-doctoris*

Colonies on PDA had reached a nine-centimeter diameter after three days at 28 °C and on WA after nine days and sporulated after four days of incubation on PDA medium. Colonies of *N. philosophiae-doctoris* developing on PDA (Figure 7e,f) were circular-shaped with aerial, woolly mycelium, entire-margined, opaque, floccose, raised with fuzzy edges, and growing rapidly; they were white-to-greyish-colored and reverse initially white, becoming creamy white when mature. Conidia of *N. philosophiae-doctoris* were round-shaped, brown-to-black-colored, aseptated, solitary, smooth, and 10.8–15.6 × 8.4–14.4 in diameter (\bar{x} = 11.6 × 13.2, n = 30). Hyphae were smooth, septate, and hyaline. On WA, colonies were white to greyish and growing poorly.

3.4. Pathogenicity Tests

3.4.1. Pathogenicity Test on Detached Leaves

The symptoms of the disease on olive leaves tested in the laboratory showed similar symptoms as the leaf samples collected from the field survey. All inoculated leaves had yellowish-to-chocolate-brown spots (Figure 8). No symptoms were spotted on the control leaves. The re-isolated fungus from the diseased leaves was identical to the *Nigrospora* species.

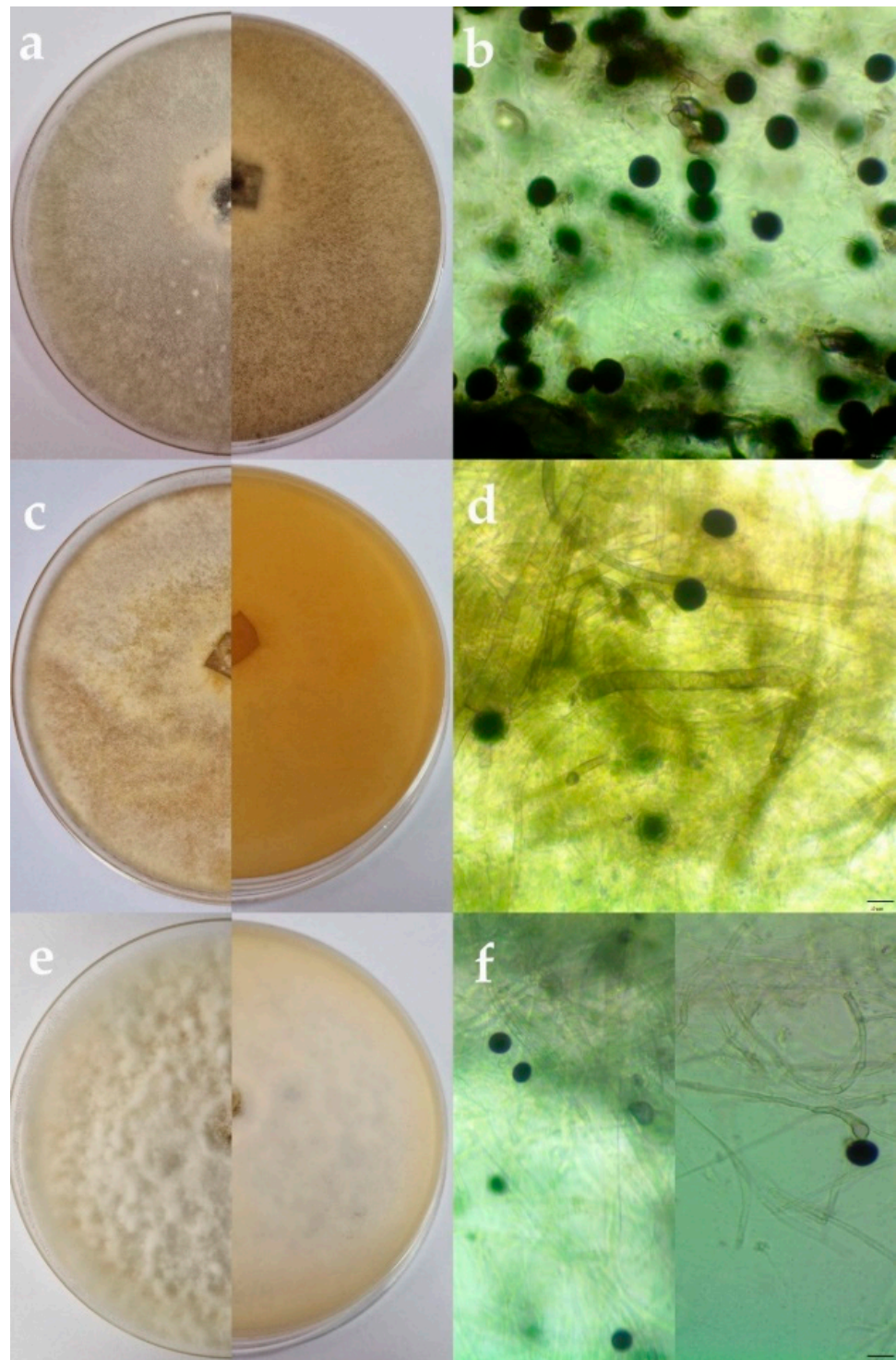


Figure 7. Left: Upper surface and reverse overview of cultures five days after incubation at 28 °C on PDA medium. Right: micrographs of isolates under the microscope with conidia. Scale bar = 10 µm, (a,b) *Nigrospora gorlenkoana*, (c,d) *N. osmanthi*, (e,f) *N. philosophiae-doctoris*.

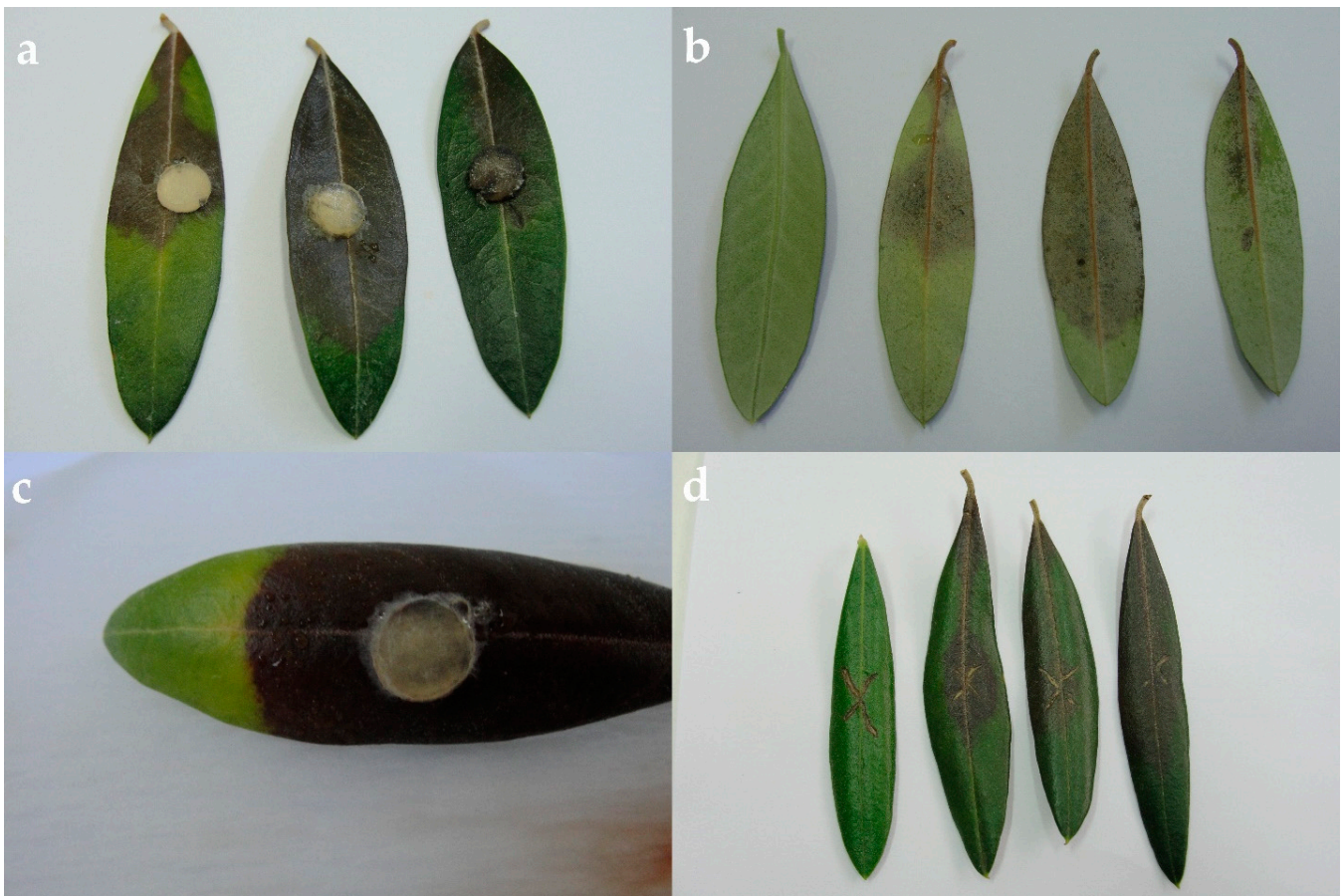


Figure 8. (a) Symptoms on unwounded leaves, from left to right: *N. philosophiae-doctoris*, *N. gorlenkoana*, *N. osmanthi*. (b) Symptoms on the bottoms of unwounded leaves, top left: control, from control to right: *N. philosophiae-doctoris*, *N. gorlenkoana*, *N. osmanthi*. (c) Symptoms on unwounded leaf: *N. philosophiae-doctoris*. (d) Symptoms on wounded leaves, from left to right: control, *N. philosophiae-doctoris*, *N. gorlenkoana*, *N. osmanthi*.

3.4.2. Pathogenicity Test on Olive Seedlings

The first symptoms were observed three days after inoculation. All inoculated leaves had chocolate-brown spots, similar to those observed in the field. The symptoms progressed until the entire leaf surface was covered (Figure 9). No symptoms were observed on the control leaves. The re-isolated fungus from the diseased leaves was identical to the *Nigrospora* species.



Figure 9. Symptoms on olive seedling leaves (cv. Rosinjola) in the pathogenicity test in the greenhouse, (a) *Nigrospora gorlenkoana*, (b) *N. osmanthi*, (c) *N. philosophiae-doctoris* after 9 days, (d) *N. philosophiae-doctoris* after 15 days.

4. Discussion

In this study, three different species of the *Nigrospora* genus, i.e., *N. gorlenkoana*, *N. osmanthi*, and *N. philosophiae-doctoris*, were detected on olive trees in Croatia. These species were the causal agents of leaf spots on olive trees in Istria and in Kvarner Gulf (Croatia).

Nigrospora species are cosmopolitan, filamentous, dematiaceous taxa distributed on various hosts, including crops with economic importance [15,16]. These species were isolated from different hosts around the world, such as *Cirsium setosum*, *Nelumbo* sp., *Oryza sativa*, *Vitis vinifera*, etc. [26]. They are known as plant pathogens, endophytes, and saprobes [35,36]. *Nigrospora* species known as plant pathogens cause many diseases, but

the most common disease is leaf spot. A list of reported *Nigrospora* pathogens, diseases, and distributions is represented in Table 6. *Nigrospora* sp. are also known as contaminants on farm-stored maize [37] and stored wheat [38] and are the causative agent of the post-harvest rot of ginger rhizomes [39] and post-harvest black rot of kiwifruit [40,41]. There are records of *N. oryzae* infecting the roots of 21 plant species [42] and *N. sphaerica* isolated from diseased grapevines [43], but there are no data on the pathogenicity of this fungus on the mentioned plant species.

Based on GenBank® data, *N. philosophiae-doctoris* was first isolated from the plant species *Disporum sessile* (Thunb.) D. Don ex Schult. & Schult. f., but there are no reports of this fungus as a plant pathogen. *N. philosophiae-doctoris* clustered in a well-supported clade closely related to *N. sacchari-officinarum* and *N. gorlenkoana*. *N. philosophiae-doctoris* produces smaller conidiogenous cells, when compared to those in *N. sacchari-officinarum* and *N. gorlenkoana*, and smaller conidia than those of *N. sacchari-officinarum* [28]. The conidial sizes frequently overlap among morphologically similar, but phylogenetically distinct, species of *Nigrospora*, and identification based on molecular and phylogenetic data for this fungal species is crucial [15]. In this research for molecular identification, consensus sequences were made of ITS1, ITS5, and ITS4 sequence data for the ITS gene region, Btub2Fd, and Btub4Rd sequence data for the TUB gene region and EF-728F and EF-986R sequence data for the EF1 α gene region. In our research, *Nigrospora* species were distinguished based on morphological and molecular data, four phylogenetic trees were made, and they support three separate species.

Sporulation in fungi usually occurs when the growth rate is reduced and is hampered under conditions that favor rapid mycelial growth [44]. Many techniques are used for inducing the sporulation of fungi, such as slide culture [45], low-nutrient media [46], sporulation on host tissue, etc. In our survey, several techniques were performed for inducing the sporulation of the JA20 NP isolate (Table 5) in order to carry out morphological research. Banana peel culture is used for inducing the sporulation of *Nigrospora sphaerica* [21] but was not effective for the *Nigrospora osmanthi* JA20 NP isolate. Interestingly, this isolate sporulates only on PDA + banana medium. Spore dispersal in *Nigrospora* is aided by the wind, rain splashes, and insect vectors [47], so it can be easily spread around the orchard and create defoliation and economical losses. In some cases, an irregular string of a mucilaginous substance (small hyaline drop) was found to be attached to the spore [48]. It has been hypothesized that this substance facilitates adherence to the host substrate or to a vector as a successful spore-dispersal mechanism [26]. The moth *Sitotroga cerealella* transports the spores of *Nigrospora oryzae*, which adhere to its body [49]. Alfaro [50] has described how the non-gravid females of the mite *Pediculopsis graminum* transport the conidia of *Nigrospora oryzae* in their abdominal sacks. Webster [48] states that the spores can be transmitted while adhered to the body of the mite. A study of airborne fungal spores carried out at nine locations in Nigeria showed that the numbers of *Nigrospora* spores significantly correlate with the relative humidity, light intensity, and temperature [51]. Research conducted on bananas has shown that with an increase in the temperature, the rotting of bananas caused by *Nigrospora* species speeds up, with maximum infection recorded at 30 °C [52]. In our survey, out of the four temperatures at which all three isolates were kept on PDA (represented in Table 5), the fastest growth was recorded at 28 °C. Therefore, this temperature was chosen as the incubation temperature for morphological analysis of the isolates.

Regarding protection measures, for *N. oryzae*, Azoxystrobin (EC₅₀ = 0.0001 mg/L) had the most significant fungal-controlling effect, followed by Prochloraz (copper salt), 15% Difenconazole + 15% Propiconazole, Difenconazole, Pyraclostrobin, and Myclobutanil [53]. Lignans, isopicropodophyllone, and dehydropodophyllotoxin, isolated from the leaves of *Podophyllum hexandrum* (Royle) T. S. Ying of Pakistani origin showed strong antifungal activity against *N. oryzae* [54]. *Bacillus thuringiensis* var. *israelensis* has shown inhibitory effects against *Nigrospora* sp. [55]. The mushroom species *Coprinellus disseminatus* (isolate 12b), *Marasmiellus palmivorus* (isolate 42b), *Trametes maxima* (isolate 56e), and *Lenti-*

nus sajan-caju (isolate 60a) have potential antagonistic effects on *Nigrospora* species via the production of secondary metabolites and mycoparasitic interactions [56]. Unfortunately, there are no data about protection measures for these three species identified in this study.

Some *Nigrospora* species can act as an antagonist against fungal and bacterial plant pathogens [57,58]. Phomalactone from *Nigrospora sphaerica* exhibits a broad spectrum of antimicrobial activity against human and phytopathogenic bacteria and fungi [59], such as *Phytophthora infestans* (Mont.) de Bary, which causes tomato late blight [60]. An ethyl acetate extract of *Nigrospora sphaerica* affects the cell wall in growing methicillin-resistant *Staphylococcus aureus* Rosenbach and *Klebsiella pneumonia* (Schroeter 1886) Trevisan [61].

Table 6. A list of reported *Nigrospora* pathogens, diseases, and distributions.

SPECIES	HOST (Common Name)	TAXONOMY ID	DISEASE SYMPTOMS	DISTRIBUTION	REFERENCES
<i>Nigrospora aurantiaca</i>	Chinese chestnut	<i>Castanea mollissima</i> Blume	Leaf spot	China	[62]
	Pandan rampoh	<i>Pandanus amaryllifolius</i> Roxb.	Leaf spot	Malaysia	[63]
	Sugarcane	<i>Saccharum officinarum</i> L.	Leaf spot	China	[30]
	Tobacco	<i>Nicotiana tabacum</i> L.	Leaf spot	China	[64]
<i>Nigrospora brasiliensis</i>	Cochineal cactus	<i>Nopalea cochenillifera</i> (L.) Salm-Dyck	Brown leaf spot	Brazil	[31]
<i>Nigrospora camelliae-sinensis</i>	Black tea	<i>Camellia sinensis</i> L.	Leaf blight	China	[65]
	Sugarcane	<i>Saccharum officinarum</i> L.	Leaf spot	China	[30]
<i>Nigrospora chiensis</i>	Tea-oil plant	<i>Camellia oleifera</i> C. Abel	Leaf blight	China	[66]
<i>Nigrospora falsivesicularis</i>	Sugarcane	<i>Saccharum officinarum</i> L.	Leaf spot	China	[30]
<i>Nigrospora guiliensis</i>	Chinese corktree	<i>Phellodendron chinense</i> C. K. Schneid.	Leaf spot	China	[67]
<i>Nigrospora hainanensis</i>	Cochineal cactus	<i>Nopalea cochenillifera</i> (L.) Salm-Dyck	Brown spot	Brazil	[68]
	Pink wood sorrel	<i>Oxalis corymbosa</i> DC.	Leaf spot	China	[69]
	Sugarcane	<i>Saccharum officinarum</i> L.	Leaf spot	China	[30]
<i>Nigrospora lacticolonia</i>	Date palm	<i>Phoenix dactylifera</i> L.	Leaf spot	Oman	[70]
	Dragon fruit	<i>Hylocereus polyrhizus</i> (F.A.C.Weber) Britton & Rose	Reddish-brown spot	Malaysia	[71]
	Great Bougainvillea	<i>Bougainvillea spectabilis</i> Raeusch. Willd.	Leaf spot	China	[72]
	Sugarcane	<i>Aloe vera</i> var. <i>chinensis</i> (Haw.) A.Berger	Leaf spot	China	[73]
			Leaf spot	Pakistan	[74]
Leaf spot			Bangladesh	[75]	
Asiatic dayflower	<i>Commelina communis</i> L.	Leaf spot	China	[76]	
<i>Nigrospora oryzae</i>	Bayberry	<i>Morella rubra</i> Lour.	Twig blight	China	[53]
	Blueberry	<i>Vaccinium corymbosum</i> L.	Leaf spot	China	[77]
	Chinese photinia	<i>Photinia serratifolia</i> (Desf.) Kalkman (syn. <i>Photinia serrulata</i> Lindl.)	Leaf spot	China	[78]
	Cotton	<i>Gossypium hirsutum</i> L.	Leaf spot	China	[79]

Table 6. Cont.

SPECIES	HOST (Common Name)	TAXONOMY ID	DISEASE SYMPTOMS	DISTRIBUTION	REFERENCES
	Cotton-rose	<i>Hibiscus mutabilis</i> Mill.	Black leaf spot	China	[80]
	Crepe-ginger	<i>Hellenia speciosa</i> (J.Koenig) Govaerts (syn. <i>Costus speciosus</i> (J.Koenig) Sm.)	Leaf spot	China	[36]
	Dendrobium (Shi Hu)	<i>Dendrobium candidum</i> Wall. ex Lindl.	Leaf spot	China	[81]
	Dove tree	<i>Davidia involucrata</i> Baill.	Leaf blight	China	[82]
	Dryland winter wheat	<i>Triticum</i> L.	Crown and rot root	Azerbaijan	[83]
	Giant red	<i>Arundo donax</i> L.	Foliar and cane rot	Europe	[84]
	Ginger	<i>Zingiber officinale</i> Roscoe	Leaf spot	China	[85]
	Indian lotus	<i>Nelumbo nucifera</i> Gaertn.	Leaf spot	China	[86]
	Indian mustard	<i>Brassica juncea</i> (L.) Czern.	Stem blight	India	[87]
	Kentucky bluegrass	<i>Poa pratensis</i> L.	Leaf spot	Ontario	[88]
	Kidney bean	<i>Phaseolus vulgaris</i> L.	Leaf spot	China	[89]
	Kiwifruit	<i>Actinidia deliciosa</i> (A.Chev.) C.F.Liang & A.R.Ferguson	Brown/black spot	China	[90]
	Million bells	<i>Calibrachoa</i> hybrid cultivar	Leaf spot	Argentina	[91]
	Pearl millet	<i>Cenchrus americanus</i> (L.) Morrone (syn. <i>Pennisetum americanum</i> (L.) Leeke)	Leaf spot	Iran	[92]
	Peppermint	<i>Mentha spicata</i> L.	Brown leaf spot	Iran	[93]
	Poplar	<i>Populus alba</i> L. × <i>P. berolinensis</i> Dipp. (hybrid poplar)	Leaf blight	China	[94]
	Rice	<i>Oryza sativa</i> L.	Sheaths and grains of sheath rot	Bangladesh	[95]
	Tobacco	<i>Nicotiana tabacum</i> L.	Leaf spot	China	[96]
	Watermelon	<i>Citrullus lanatus</i> (Thunb.) Matsum. & Nakai	Leaf spot	China	[97]
	Wheat	<i>Triticum aestivum</i> Vill.	Dark brown to black lesions	Kazakhstan	[98]
			Crown and root rot	Kazakhstan	[99]
	Wild rice	<i>Oryza rufipogon</i> Griff.	Leaf spot	China	[100]
	Zebra leaf aloe	<i>Aloe zebrina</i> Baker	Flower malformation	Namibia	[101]
	Fiddle-leaf fig	<i>Ficus pandurata</i> Hance	Leaf blight	China	[102]
	Java tea	<i>Orthosiphon stamineus</i> Benth.	Leaf blight	Malaysia	[103]
<i>Nigrospora osmanthi</i>	St. Augustine grass	<i>Stenotaphrum secundatum</i> (Walter) Kuntze	Leaf blight	China	[104]
	Tartary buckwheat	<i>Fagopyrum tataricum</i> (L.) Gaertn.	Leaf spot	China	[105]
	Big marigold	<i>Tagetes erecta</i> L.	Leaf blight	Bangladesh	[106]
<i>Nigrospora panici</i>	French marigold	<i>Tagetes patula</i> L.	Leaf blight	Bangladesh	[106]
	Sugarcane	<i>Saccharum officinarum</i> L.	Leaf spot	China	[30]
<i>Nigrospora pyriformis</i>	White goosefoot	<i>Chenopodium album</i> L.	Leaf spot	China	[107]

Table 6. Cont.

SPECIES	HOST (Common Name)	TAXONOMY ID	DISEASE SYMPTOMS	DISTRIBUTION	REFERENCES	
<i>Nigrospora sacchari-officinarum</i>	Sugarcane	<i>Saccharum officinarum</i> L.	Leaf spot	China	[30]	
<i>Nigrospora saccharicola</i>	Sugarcane	<i>Saccharum officinarum</i> L.	Leaf spot	China	[30]	
<i>Nigrospora singularis</i>	Sugarcane	<i>Saccharum officinarum</i> L.	Leaf spot	China	[30]	
<i>Nigrospora</i> sp.	Black tea	<i>Camellia sinensis</i> L.	Blister blight lesions	India	[108]	
	Cochineal cactus	<i>Nopalea cochenillifera</i> (L.) Salm-Dyck	Brown spot	Brazil	[68]	
	Maize	<i>Zea mays</i> L.	Weight / discoloration / necrosis of grains	Brazil	[109]	
<i>Nigrospora sphaerica</i>	Balloon flower	<i>Platycodon grandiflorus</i> (Jacq.) A. DC.	Necrosis	China	[28]	
	Black tea	<i>Camellia sinensis</i> L.	Leaf blight	India	[110]	
				China	[111]	
	Blueberry	<i>Vaccinium corymbosum</i> L.	Leaf spot, twig and shoot blight	Argentina	[112]	
	Calabash	<i>Lagenaria siceraria</i> (Molina) Standl.	Leaf spot	Georgia	[113]	
	China fir	<i>Cunninghamia lanceolata</i> (Lamb.) Hook.	Leaf blight	China	[114]	
	Chinese Wisteria	<i>Wisteria sinensis</i> (Sims) DC., 1825	Leaf spot	Turkey	[115]	
	Cochineal cactus	<i>Nopalea cochenillifera</i> (L.) Salm-Dyck	Brown spot	Brazil	[68]	
	Corn mint	<i>Mentha canadensis</i> L.	Leaf blight	China	[116]	
	Cowpea	<i>Vigna unguiculata</i> (L.) Walp.	Leaf spot	India	[117]	
	Curcuma	<i>Curcuma wenyujin</i> Y.H.Chen & C.Ling	Leaf blight	China	[118]	
				Not applicable	Iraq	[119]
				Root disease	Oman	[120]
	Date palm	<i>Phoenix dactylifera</i> L.	Leaf spot	Pakistan	[121]	
	Devilpepper	<i>Rauwolfia serpentina</i> (L.) Benth. ex Kurz	Leaf spot and antracnose	Bangladesh	[122]	
	Dragon fruit	<i>Selenicereus monacanthus</i> (hort. ex Lem.) D.R.Hunt (syn. <i>Hylocereus polyrhizus</i> (F.A.C.Weber) Britton & Rose)	Reddish-brown spot	Malaysia	[71]	
	Dragon fruit (pitaya)	<i>Selenicereus undatus</i> (Haw.) D.R.Hunt (syn. <i>Hylocereus undatus</i> (Haw.) Britton & Rose)	Reddish-brown spot	Philippines	[123]	
			Reddish-brown spot	China	[124]	
	Elephant grass	<i>Cenchrus purpureus</i> (Schumach.) Morrone	Leaf blight	China	[125]	
European nettle tree	<i>Celtis australis</i> L., 1753	Leaf spot	India	[126]		
False Daisy	<i>Eclipta prostrata</i> (L.) L.	Leaf spot	China	[127]		
Kinnow mandarin	hybrid: <i>Citrus nobilis</i> Lour. × <i>Citrus deliciosa</i> Ten.	Leaf spot	Pakistan	[128]		
Kiwifruit	<i>Actinidia deliciosa</i> (A. Chev.) C.F.Liang & A.R.Ferguson	Leaf spot	China	[129]		
Liquorice	<i>Glycyrrhiza glabra</i> L.	Leaf spot	India	[130]		

Table 6. Cont.

SPECIES	HOST (Common Name)	TAXONOMY ID	DISEASE SYMPTOMS	DISTRIBUTION	REFERENCES
	Mango	<i>Mangifera indica</i> L.	Leaf spot	India	[131]
			Twig dieback and leaf spot	Egypt	[132]
	Moonlight cactus	<i>Selenicereus monacanthus</i> (hort. ex Lem.) D.R.Hunt (syn. <i>Hylocereus monacanthus</i> (hort. ex Lem.) Britton & Rose)	Reddish-brown spot	Philippines	[123]
	Mulberry	<i>Morus alba</i> Hort. ex Loudon L	Shot hole	China	[133]
				India	[134]
	Passion fruit	<i>Passiflora edulis</i> Sims	Leaf blight	China	[135]
	Peanut	<i>Arachis hypogaea</i> L.	Leaf blight	China	[136]
	Pitaya	<i>Selenicereus megalanthus</i> (K.Schum. ex Vaupel) Moran (syn. <i>Hylocereus megalanthus</i> (K.Schum. ex Vaupel) Ralf Bauer)	Reddish-brown spot	Philippines	[123]
	Purging nut	<i>Jatropha curcas</i> L.	Necrosis, chlorosis	India	[137]
	Qing qian liu	<i>Cyclocarya paliurus</i> (Batalin) Iljinsk.	Leaf blight	China	[138]
	Sesame	<i>Sesamum indicum</i> L.	Leaf blight	China	[139]
			Leaf blight	Pakistan	[140]
	Sugarcane	<i>Saccharum</i> spp.	Leaf blight	China	[141]
	Sugarcane	<i>Saccharum officinarum</i> L.	Leaf spot	China	[30]
	Three-leaf Akebia	<i>Akebia trifoliata</i> (Thunb.) Koidz.	Dried-shrink fruit	China	[142]
	Tea-oil plant	<i>Camellia oleifera</i> C. Abel	Leaf blight	China	[143]
	Watermelon (wild melon)	<i>Citrullus lanatus</i> (Thunb.) Matsum. & Nakai	Leaf spot	Malaysia	[144]
	White moho	<i>Helicarpus americanus</i> L.	Leaf spot	Brazil	[145]
<i>Nigrospora vesicularifera</i>	Sugarcane	<i>Saccharum officinarum</i> L.	Leaf spot	China	[30]
<i>Nigrospora zimmermani</i>	Sugarcane	<i>Saccharum officinarum</i> L.	Leaf spot	China	[30]

5. Conclusions

This study identified, described and characterized three fungal species that caused leaf spot symptoms on olive trees in Croatia. *Nigrospora* species can be economically significant as plant pathogens, causing crop losses in agriculture. Early detection can help prevent the spreading, so it is important to identify and manage leaf spot promptly to prevent it from causing damage to olive trees. Over time, repeated infections can weaken the olive tree. Weakened trees are more susceptible to other diseases and environmental stressors, which can lead to a decline in the tree's overall health and longevity. Also, fungi can spread to other parts of the tree and neighboring trees, which can lead to more widespread infections and increased management challenges for growers. Additionally, the cost of managing and treating leaf spot diseases can add to production expenses. In conclusion, leaf spot diseases on olive trees are important because they can negatively affect the tree's health, fruit quality, and overall productivity. Olive growers need to monitor for leaf spot diseases and implement effective management strategies to minimize their impact and ensure a healthy and productive orchard. It is also necessary to conduct further research

that will include monitoring these fungal diseases and studying the effectiveness of various substances or treatments in inhibiting the growth and reproduction of these fungi.

To our knowledge, this paper is the first report of *Nigrospora* species causing diseases on olives and the first report of *Nigrospora philosophiae-doctoris* causing plant disease.

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