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Characterization of Mycoflora Associated with *Catharanthus roseus* Collected from Gardens in Kenitra City, Morocco

Najoua Mouden ^{1*} , Mohamed Elouardi ², Nouha Achajri ³, Jamal Mabrouki ², Rachid Benkirane ³, Allal Douira ³, Amina Ouazzani Touhami ³

¹ Laboratory of Molecular Chemistry, Materials and Environment, Multidisciplinary Faculty Nador (FPN), Mohammed 1st University, Oujda 62702, Morocco

² Laboratory of Spectroscopy, Molecular Modelling, Materials, Nanomaterials, Water and Environment, Faculty of Science, Mohammed V University, Rabat BP1014, Morocco

³ Laboratory of Plant, Animal and Agro-industry Productions, Ibn Tofail University, Kenitra 14000, Morocco

ABSTRACT

This paper reports on the composition of fungal communities occurring on diseased tissues of *Catharanthus roseus*, which differed between organs. In total, ten different filamentous fungi were isolated, and the percentage of isolation varied significantly among the organs. *Botrytis cinerea* was the most prevalent fungus found on the plant's aboveground parts, with a frequency exceeding 50%. On twigs, the occurrence rate was 95.6%. It was isolated from leaves with a frequency of 88%, followed by *Aspergillus niger* (71.66%), *Alternaria alternata* (67.33%), *Cladosporium herbarum* (61%), *Fusarium oxysporum* (50.66%), *Epicoccum nigrum* (57.66%), *Curvularia lunata* (49.66%), *Trichoderma harzianum* (40%), and *Penicillium* sp. (27%). Whereas, *Fusarium* genus was more represented and six species were recorded: *F. subglutinans* (26%), *F. chlamydosporium* (20%), *F. vertillioides* (15.66%), *F. solani* (10%), *F. oxysporum* and *F. nivale*. Results highlighted dissimilar distribution of *Fusarium* species was noted on *Catharanthus* tissues on which *F. subglutinans*, *F. chlamydosporium* and *F. oxysporum* coexist on leaves and roots while *F. solani* was retrieved from leaves against *F. nivale* from roots. The floral buds and pods harbored opportunist fungi such as *B. cinerea*, *Alternaria alternata* and *E.*

*CORRESPONDING AUTHOR:

Najoua Mouden, Laboratory of Molecular Chemistry, Materials and Environment, Multidisciplinary Faculty Nador (FPN), Mohammed 1st University, Oujda 62702, Morocco; Email: n.mouden@ump.ac.ma

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nigrum. On roots, *Aspergillus*, *Penicillium* and *Fusarium* were the main genera occurring with the frequencies of 26% (*F. chlamydosporium*), 40% (*F. nivale*), 72.33% (*A. niger*), 47.66% (*A. flavus*) and 37.66% (*A. fumigatus*). But no fungal species were detected on seeds of *Catharanthus roseus*. This is the first study to describe and enumerate the fungal complex associated with various symptoms on the aerial parts of *Catharanthus roseus*.

Keywords: Isolation; Identification; *Catharanthus roseus*; Mycoflora; Morocco

1. Introduction

Catharanthus roseus known as Madagascar periwinkle, is a perennial plant belonging to the Apocynaceae family, widely distributed ornamental plant and holds significant medicinal value^[1]. This species thrives in dry, warm areas and can grow in full sun or partial shade. There are two varieties of this plant which can be distinguished by the color of the flower viz, the rosea with pink flowers and the alba with white flowers^[2]

Catharanthus roseus is a valued commercial plant due to many essential alkaloids used to treat high blood pressure and circulatory disorders^[3, 4]. It also possesses anti-cancer, anti-diabetic, anti-Alzheimer's properties in addition to be effective for wound-treatment^[5, 6]. Periwinkle is also known to yield a variety of terpenoid indole alkaloids in the form of vincristine, vinblastine, catharanthine, ajmalicin, vindoline, etc., which are of great significance in the medical industry^[7]. The potential medicinal benefits include antiviral, antidiabetic, antihypertensive, antibacterial, antifungal and antioxidant activities^[8, 9]. But, its value as an ornamental^[2] also promises an increasing interest for use.

Catharanthus roseus (L.) G. is the host of hundreds of endophytic bacteria and fungal organisms that inhabit different areas of plants^[10]. The latter exhibit different types of association with their host organisms, such as a combination of symbiosis, pathogenesis, antagonistic relationships, etc., beside being able to promote plant growth, counteract plant stress conditions, and even elicit plant defense against any pathogenic attack^[11, 12]. *C. roseus* endophytic fungi have also been reported to produce certain compounds with potential cytotoxic activities^[13] and antioxidant activity^[14]. Dhayanithy^[15] have shown that *Chaetomium nigricolor* found among twenty endophytic isolates obtained from different parts of the plant, had significant cytotoxic, apoptotic and antioxidant potential. Various species of the

Nigrospora genus, such as *Nigrospora zimmermanii* and *N. sphaerica*, isolated from *Catharanthus* plants, have the capacity to produce the anticancer agent vincristine and hydrolytic enzymes^[16, 17].

On the other hand, there are several pathogens documented worldwide causing many different forms of disease in *C. roseus*^[18–21]. Diseases include root rot and seedling damping-off, caused by a wide range of fungi, such as *Rhizoctonia solani*, *Fusarium oxysporum*, *F. solani*, *Thielaviopsis* sp., *Sclerotium rolfsii* and *Pythium aphanidermatum* or Foliar blight caused by *Colletotrichum dematium* and *Botrytis cinerea*^[22]. Others fungi such as *Sclerotinia sclerotiorum* are regarded as a plant pathogen on Periwinkle^[23] and *Macrophomina phaseolina* (Tassi) Goid that was proved to be pathogenic to *C. roseus* by causing leaf blight^[24]. Madagascar periwinkle is also affected by *Phytophthora tropicalis* responsible for leaf blight disease symptoms^[25]. Sharma^[21] have recoded twig blight on *C. roseus* induced by *Colletotrichum gloeosporioides*. Garibaldi^[26] described leaf blight symptoms on *Catharanthus roseus* plant caused by *Rhizoctonia solani*. Baysal-Gurel^[27] have identified the causal agent for *Phytophthora* aerial blight symptom on annual vinca as *Phytophthora nicotiana*. Also, Intaparn^[28] have reported root and stem rot induced by *Phytophthora* sp.

Generally, when inhabiting host plant, endophytes and pathogens can be related via a various interactions which probably determine survival, pathologies appearance and growth performance of plants. In Morocco, no study was carried out to determine the fungal community in association with *Catharanthus roseus*.

Against this background, the aim was to describe and identify the fungal organisms involved in the necrotic damage observed on this plant species.

2. Materiel and Methods

2.1. Samples Collection

Symptomatic *Catharanthus roseus* leaves, tiges, pods and seeds showing necrotic lesions were detached from twenty *Catharanthus* plant samples which were collected from gardens in Kenitra city, Morocco. Sterilized polythene bags were used to collect the samples. They were brought out to the laboratory, and they were processed within twenty-four hours of being collected.

2.2. Isolation and Identification

Fungi were isolated from aerial parts of *C. roseus* as described by^[29]. Different tissues surface was first disinfected by immersing them in 70% ethanol for 60 seconds, then 90 seconds in 70% sodium hypochlorite, and finally in sterile distilled water for 60 seconds before air-drying for 5 minutes on filter paper (150 mm) under sterilization conditions. Disinfected leaf tissues were cut into 1cm × 1cm segments and ten pieces were placed on petri dishes. At room temperature, the dishes were incubated for 14 days then repeatedly observed for fungal growth. To obtain a pure culture, an inoculum of each new colony was continuously transferred onto a fresh PSA medium (Potato Sucrose Agar: 200 g potato, 20 g sucrose, 15 g Agar-agar, 1000 ml distilled water) added with 100 mg/L of chloramphenicol.

Diseased seedlings were rooted out and cut at a height of 5 cm from crown. The roots were then rinsed under running water and cut into pieces 0.5 cm long. Samples were surface disinfected with sodium hydrochlorite (0.5% chlorine) for three minutes, then rinsed twice with distilled water for a further two minutes and dried with sterile filter paper. Root fragments were then placed in 9 cm Petri dishes containing potato sucrose agar (PSA). Plates were kept in incubation at 25 ± 1 °C for 4 days, after which the fungi were purified by cutting out a bit of mycelia taken from the edge of colony and cultured in the center of a petri dish containing the PDA and incubated for 7 days^[30]. Subsequently, the purified fungal collections were identified using fungal recognition keys derived from phenotypic hallmarks, such as fungal mycelial structure and spore morphology, fungal colony color, branching nature of the mycelium, conidiophore shape and size, spores formed and ability to form sclerotia^[31–34]. Cultivation set-ups on slides were used to study the arrangement of conidia under the light microscope. Periodic examination of the

slides was performed using the lactophenol cotton blue stain when sufficient growth had been achieved. Fungal types and species were selected by light microscopy in accordance with the key determinations of^[35–40]. The isolation percentage and/or contamination, which defines the incidence of isolation of the different fungi from 100 lesions, was then calculated using Ponchet's formula^[41]: $PC = (NLI/NTL) \times 100$ PC: Percent of infection and/or contamination; NLI: The number of lesions containing the fungal species. NTL: Total number of segments with lesions used for fungus isolation.

2.3. Statistical Analysis

Analysis was performed using a one-way (ANOVA) and an LSD test at the 5% test level. Per cent were transformed into $\text{Arcsin } \sqrt{P}$ (where P is the isolation percentage).

3. Results and Discussion

Infected plants showed symptoms mainly on the leaves. Lesions are circular (approx. 0.1 to 0.3 cm long), brownish-yellow to blackish. These spots lengthen and become elliptical or rectangular (2 cm long), with a dark brown center. Occasionally, the attacked leaves become twisted, deformed, and desiccated. Leaves then rot and become covered with a velvety, grayish felt (**Figure 1**).



Figure 1. Various symptoms on *Catharanthus roseus* plants. (a) Leaf brown spots and yellowing; (b) desiccation symptoms of leaves detached from infected plants of *Catharanthus roseus* (L.) G. Don. in the gardens of the city of Kenitra.

It has been observed that infection generally starts at the upper end of flower buds and pods, and subsequently, the whole pod may be attacked (**Figure 2**). On the stems, infected parts are covered with a conidiation greyish felting (**Figure 3**).

A diverse fungus complex was found on the leaves, stems, flower buds, pods, and roots. As a result, *C. roseus*

recorded a large and varied fungal community (Table 1). From the host's leaf, stem, and root tissues, a total of 17 species, representing 13 fungal groups, were identified (Table 2). The most dominant fungal organisms were *B. cinerea* and *Alternaria alternata*. *B. cinerea* was found on leaves, floral buds and pods with respective percentage of contamination of 88%, 53% and 49% compared to 67,3%; 36,66% and 47,33% for *A. alternata* undetectable at twigs level of infected *Catharanthus roseus*. The appearance of a fungus on a tissue did not express a close and specific relationship with *Catharanthus* tissues. Lesions in one organ can contain many species that also appear in other organs, despite having different frequencies. A complex of species is transmitted regardless the microenvironment within each organ. In general, 13 species were isolated from leaves, 4 from flower buds, 3 from pods, 1 from stems, 10 from roots, and no species from seeds.



Figure 2. Necrosis on surface of floral buds. (a) pods; (b) of *Catharanthus roseus* plants samples.



Figure 3. Infected plants of *Catharanthus roseus* showing greyish to black mycelia and fruiting bodies covering stems.

Table 1. Fungal isolates recovered from leaves, stems, pods, seeds, and roots of symptomatic *Catharanthus roseus* Cv *rosea* planted in gardens area, Kenitra city.

Fungal Species	Stems	Leaves	Floral Buds	Pods	Roots	Seeds
<i>Botrytis cinerea</i>	BcS1, BcS1', BcS1"	BcS2, BcS2', BcS2"	BcS3, BcS3', BcS3"	BcS4	–	–
<i>Aspergillus niger</i>	–	An fS1, AnfS2	–	–	An rS1, An rS2	–
<i>Alternaria alternata</i>	–	Aa fS1, AafS2, AafS3	Aa bS1, AabS2, Aab S3	Aa gS1, AagS2	–	–
<i>Cladosporium herbarum</i>	–	ChS1, ChS2, ChS3, Ch4	–	–	–	–
<i>Fusarium oxysporum</i>	–	Fo f	–	–	Fo r	–
<i>Curvularia lunata</i>	–	Cl f	Cl b	–	Cl r	–
<i>Trichoderma harzianum</i>	–	ThS1, Th S2, ThS3, ThS4	–	–	–	–
<i>Penicillium sp</i>	–	Pn S1sp, PnS2sp	–	–	–	–
<i>Fusarium chlamydosporum</i>	–	Fch f	–	–	Fch r	–
<i>Fusarium vertillioides</i> ,	–	Fv f	–	–	Fv r	–
<i>Epicoccum nigrum</i>	–	EnfS1, Enf S2, EnfS3	En bS1, EnbS2	EngS1, EngS2	–	–
<i>Penicillium frequetans</i>	–	–	–	–	PfS1, PS2	–
<i>Fusarium solani</i>	–	FsS1, FsS2	–	–	–	–
<i>Fusarium nivale</i>	–	–	–	–	Fn	–
<i>Fusarium subglutinans</i>	–	Fsb f	–	–	Fsb r	–
<i>Aspergillus flavus</i>	–	–	–	–	Af	–
<i>Aspergillus fumigatus</i>	–	–	–	–	Afu	–

(-): non-isolated species; (S): strains. BC: *Botrytis cineria*, Af: *Aspergillus flavus*, An: *Aspergillus niger*, Af: *Aspergillus fumigatus*, Pn sp.: *Penicillium* sp., Cl: *Curvularia lunata*, Th: *Trichoderma harzianum*, Fc: *Fusarium chlamydosporum*, Aa: *Alternaria alternata*, Ch: *Cladosporium herbarum*, Fn: *Fusarium nivale*, Fo: *Fusarium oxysporum*, Fv: *Fusarium verticillioideis*, Fs: *Fusarium solani*, Fsb: *Fusarium subglutinans*, Pf: *Penicillium frequetans*, En: *Epicoccum nigrum*.

Table 2. Percentage of contamination of *Catharanthus roseus* by different fungal species isolated using Blotter method.

Fungal Species	Twigs (%)	Leaves (%)	Floral Buds (%)	Pods (%)	Roots (%)
<i>Botrytis cinerea</i>	95 ^a	88 ^a	53 ^{cd}	49 ^{cd}	—
<i>Aspergillus niger</i> *	—	71,66 ^b	—	—	72,33 ^b
<i>Alternaria alternata</i>	—	67,3 ^b	36,66 ^{ef}	47,33 ^d	—
<i>Cladosporium herbarum</i>	—	61 ^{bc}	—	—	—
<i>Fusarium oxysporum</i> *	—	50,66 ^{cd}	—	—	67,3 ^b
<i>Curvularia lunata</i> *	—	49,66 ^{cd}	18,33 ^{fg}	—	49,66 ^{cd}
<i>Trichoderma harzianum</i> *	—	40 ^e	—	—	70 ^b
<i>Penicillium sp.</i>	—	27 ^f	—	—	—
<i>Fusarium chlamydosporum</i> *	—	20 ^{fg}	—	—	26 ^f
<i>Fusarium verticillioides</i> *	—	15,66 ^{fg}	—	—	35 ^{ef}
<i>Epicoccum nigrum</i>	—	57,66 ^c	27 ^f	20,66 ^{fg}	—
<i>Penicillium frequentans</i>	—	—	—	—	47,33 ^d
<i>Fusarium solani</i>	—	10 ^g	—	—	—
<i>Fusarium nivale</i>	—	—	—	—	40 ^e
<i>Fusarium subglutinans</i>	—	26 ^f	—	—	—
<i>Aspergillus flavus</i>	—	—	—	—	47,66 ^d
<i>Aspergillus fumigatus</i>	—	—	—	—	37,66 ^{ef}

* At the 5% level, there was no significant difference between two results that were read on the same column.

At the root level, out of 10 retrieved fungal species, the genus *Fusarium* was represented by three species: *Fusarium oxysporum* (67.33%), *F. nivale*, and *F. verticillioides* (35%), with *Fusarium chlamydosporium* (26%). The highest frequency was found in *A. niger* (72.33%), followed by *Trichoderma harzianum* (70%) and 49.6% for *C. lunata*, which significantly displayed an equal isolation frequency to those of *P. frequentans* and *A. flavus*. Both macroscopic and microscopic morphological characteristics have been described for fungal species grown on Potato sucrose agar culture medium.

The morphological examination of *Alternaria alternata* grown on PSA showed brownish olivaceous colony. Conidiophores were usually short being either ramified or simple, solitary or grouped mostly curved but some are straight. Conidia produced usually in long branched strings in diverse shapes; oval, obclavate, obpyriform or ellipsoidal with average size $4\text{--}38 \times 7\text{--}14 \mu\text{m}$ with 1 to 8 septa. They possessed 1 to 8 transverse septa with a short apical beak (**Figure 4a,b**).

Epicoccum nigrum colony is fast growing, suede-like to downy, with a strong yellow to orange-brown diffusible pigment. Conidiophores vary in size and occur in dense, pale-colored clusters measuring $12\text{--}14 \times 5\text{--}4 \mu\text{m}$. They are grouped into aggregates (sporodochia). The conidia, $15\text{--}25 \mu\text{m}$ in diameter, are globe-shaped, pyriform, and lightly pigmented with a bright yellow to orange-brown color (**Figure 4c**).

Botrytis cinerea had fluffy colony, with gray-white

aerial mycelia on PSA, then turned gray and formed ovate to elliptical conidia on the mycelia after 7 days of incubation. It develops tufts of upright, greyish (brownish grey or ash grey) conidiophores with branches at the top. The conidia and conidiophores appear grayish in mass when young, turning a light to medium brown on maturation with size ranging from 12–18 to 13.59 mm in length by 5–12 μm in width (**Figure 4d**). The fungus generates black sclerotia. These structures ensure the survival of the fungus during unfavourable periods.

Curvularia lunata forms black colonies. Conidiophores are straight or flexible, geniculate, usually smooth and brown in color. Conidia are solitary, simple, usually curved, elliptical, fusiform, ovoid, measuring $17\text{--}30 \times 4\text{--}13 \mu\text{m}$ with 3 transverse septa, pale to dark brown, smooth or verruculose (**Figure 4e**).

Culture of *Fusarium verticillioides* has white mycelia, cottony, ranging in color from white to salmon. The short conidiophores (phialides), which taper towards the tip with rough collars, are solitary or in branched clusters. Microconidia are oval, 0-septate borne from monophialides measuring $7\text{--}10 \times 2,5 \text{ à } 3,2 \mu\text{m}$. This fungus produces oval or club-shaped macroconidia, $31 \text{ à } 58 \times 2,7 \text{ à } 3,6 \mu\text{m}$, usually 3–7 septate sometimes forming chains. No chlamydospores were observed (**Figure 4f**).

The olivaceous-brown, slow-growing *Cladosporium herbarum* colony turns powdery as a result of the creation of

a lot of conidia. It is olivaceous-black on the back. Conidiophores can be upright, straight or flexuose, unbranched or just branching in the apical region, and in certain species, they

have geniculate sympodial elongation. Conidia are ovoid to cylindrical, 4–11 μm in size, and formed in branched chains with a noticeable black hilum (**Figure 4g**).

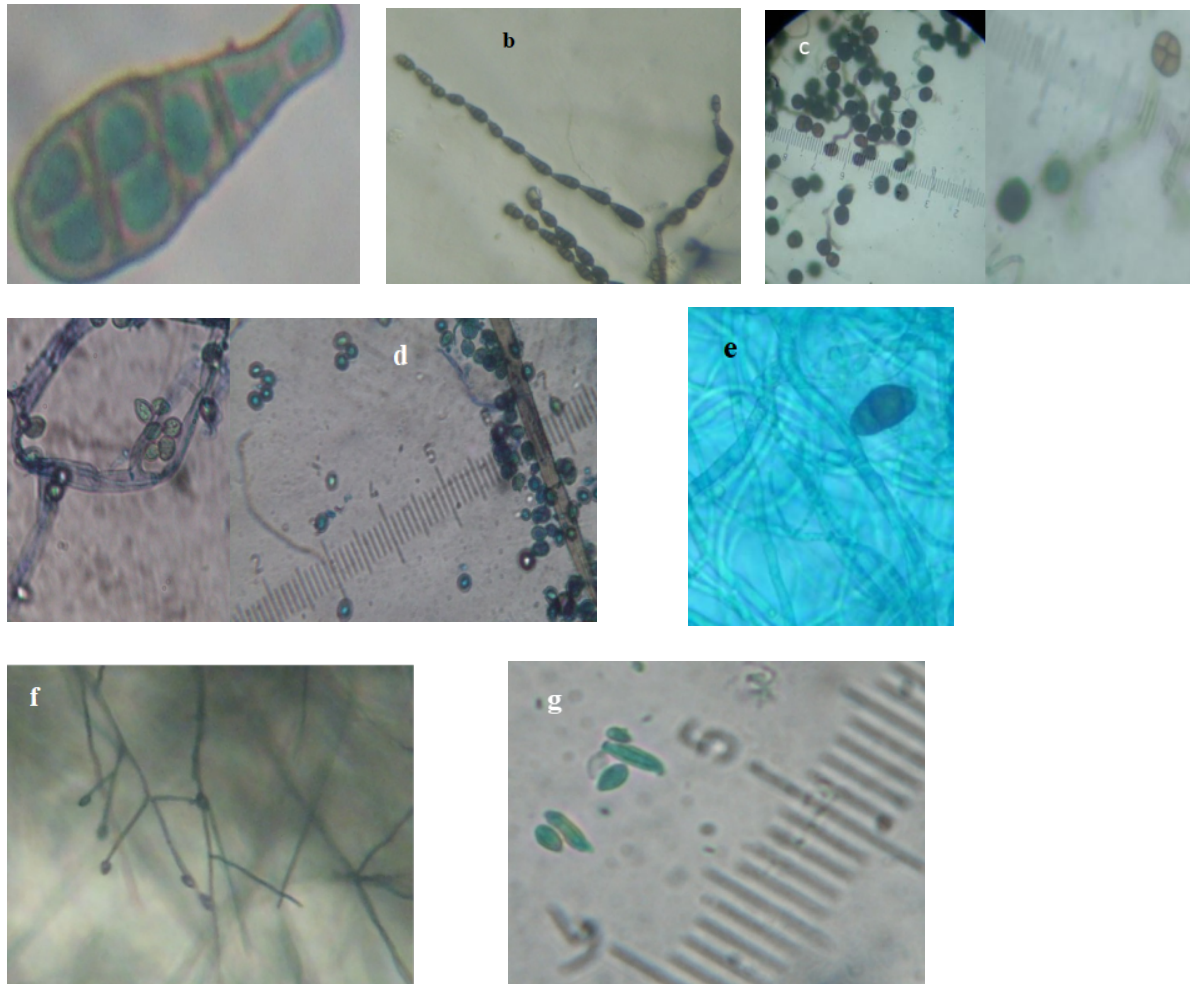


Figure 4. Conidia and conidiophore morphology of fungal species isolated from *Catharanthus roseus*. (a) conidia; (b) chain of *Alternaria alternata* conidia; (c) conidiophore and conidia of *Epicoccum nigrum*; (d) conidiophore and conidia of *Botrytis cinerea*; (e) conidia of *Curvularia lunata*; (f) conidiophores and conidia of *Fusarium verticillioides*; (g) conidia of *Cladosporium herbarum* observed under light microscope at ($\times 400$) magnification, Mounting liquid: Bleu coton.

The *Trichoderma harzianum* colony grows quickly. Initially, the mycelium is white, but it later turns green and powdery. The conidiophore is branched and produces phialides, which bear conidia at the top that group together into bundles. The conidia are unicellular, and they are globular or ovoid in shape (**Figure 5A₁, A₂**).

The aerial mycelium of *Fusarium oxysporum* is white and dense. The mycelium turns violet or blue when there are a lot of chlamydo spores, but it turns yellowish brown when there are a lot of sporodochia. Usually abundant, microconidia (3–15 \times 3–5 μm) are carried on branched conidiophores

or short, simple phialides. They are typically unicellular, ellipsoidal to cylindrical, straight, or curved, and they are never generated in chains. Macroconidia are fusiform, somewhat curved, pointed at the ends, and frequently triseptate (20–35 \times 3–5 μm). On the mycelium, chlamydo spores can be terminal or intercalary, spherical, solitary, or in clusters of two to three, and hyaline with smooth or rough walls (**Figure 5B**).

The colony of *Fusarium solani* grows rapidly with abundant mycelium. Microconidia are sparse, with a single cell. The macroconidia are large, with 3 to 4 septa. The spore wall are parallel and the apical and basal cells are

slightly domed (Figure). Chlamydoconidia are solitary (7.3 μm) or in pairs 8 to 10.6 μm long, abundant, with smooth walls (Figure 5C).

Fusarium nivale (Fr.) Sorauer 1901; has a yellowish-white colony with an orange underside. Aerial mycelium is sparse. Microconidia are absent. Macroconidia are present, small, curved with 1 to 3 septa (Figure 5D₁, D₂).

The *Fusarium subglutinans* culture appears white with a dark purple underside. It contains abundant oval microconidia, most of which are formed by a single cell. The macroconidia are also abundant, almost straight, thin-walled,

and with the two surfaces of the spore almost parallel (Figure 5E₁, E₂).

The *Fusarium chlamydosporium* culture grows quickly on PDA. The mycelium is flaky, fairly dense, off-white in color, and turns violet in the older parts of the colony, with a reverse of red to brown shades. Microconidia are abundant straight to comma-shaped, usually without any septation. Macroconidia were sparse, moderately curved with 2 to 3 septa. Chlamydoconidia were found in large number and were single, in chains and intercalary. Conidiophores are short and branched with mono- and polyphialides (Figure 5F₁, F₂).

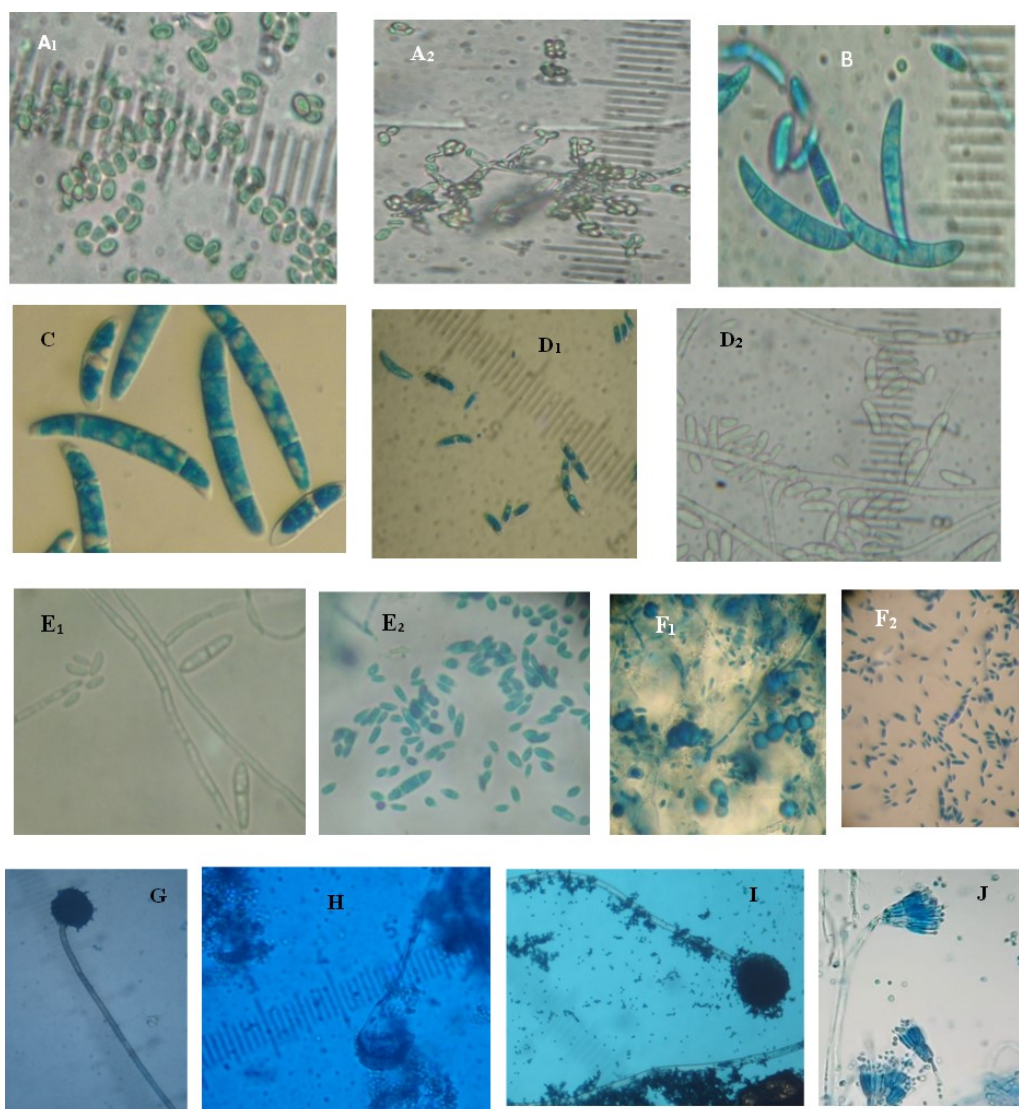


Figure 5. Micromorphology (after methylene blue staining) of fungal species isolated from *Catharanthus roseus* plants. (A₁,A₂) Conidia and conidiophores of *Trichoderma harzianum*; (B) conidia of *Fusarium oxysporum*; (C) conidia of *Fusarium solani*; (D₁,D₂) Conidia and conidiophores of *Fusarium nivale*; (E₁,E₂) conidiophore and conidia of *F. subglutinans*; (F₁,F₂) Chlamydoconidia and macro-microconidia of *Fusarium Chlamydosporium*; (G):conidia and conidiophore of *Aspergillus niger*; (H): conidiophore of *Aspergillus flavus*; (I) conidiophore and conidia of *Aspergillus fumigatus*; (J) conidiophore and conidia of *Penicillium* sp. obtained in PSA medium at ($\times 400$) magnification.

The colony of *Aspergillus niger* starts off as white and translucent, then turns black as it produces spores. *Aspergillus* species are identified by a conidiophore with a swollen end (or vesicle). This end is spherical (40 µm in diameter) and has bottle-shaped phialides arranged all around it. The conidiophores are smooth, clear, brownish at the top, and very long (1 to 3 mm). The conidia are single-celled and single-nucleated, round, 2 to 3 µm in diameter, black or dark brown, and covered with spines and projections (echinulate form). They are often found in chains and spread by the wind (**Figure 5G**).

The colony of *Aspergillus flavus* Link, 1809 grows quickly, displaying sparse, greenish-yellow aerial mycelium with a light yellow reverse side. The conidiophore gradually enlarges upwards to form a vesicle 10–40 µm in diameter. The conidia are almost round, greenish-yellow in color, and vary greatly in size from $2 \times 3.3 \times 4.6$ µm to 5×6 µm in diameter (**Figure 5H**).

The colony of *Aspergillus fumigatus* has an olive-green color with a yellowish underside. The conidiophore is short and septate, gradually enlarging towards the apex to form a vesicle 20–30 µm in diameter. The conidia are blackish-green, globose, and 2 to 3.5 µm in diameter (**Figure 5I**).

The *Penicillium* sp. colony is green. The mycelium hyphae are septate. The straight conidiophore is generally unbranched, with whorls at the top. The conidia are globular in shape, smooth-walled, and arranged in chains (**Figure 5J**).

4. Discussion and Conclusion

No research on fungal infections of *Catharanthus roseus* has been done in Morocco. The acquired data demonstrated how the fungal community's makeup affects the health of these attractive plants. The presence of a varied fungal complex on the plant organs under study can be explained by the fact that a lesion might be started by a single infection and then colonized by others. Ellis^[37] asserts that each species' frequency is determined by the temperature and humidity levels that are conducive to its growth^[42]. There are fewer species detected on roots than leaf lesions, which is consistent with findings by^[43]. As per the hypothesis of^[15], the *Catharanthus* plant's bark has the highest species variety and richness, followed by the leaf and stem tissues.

The obtained results confirmed previous studies that

have already reported some of our isolated fungal species while others were encountered for the first time on *Catharanthus roseus* in Morocco. The lesions observed on the aerial parts were mainly dominated by *B. cinerea*. In fact, this species has a wide range of plant hosts such as vegetables, grapevine and fruit trees^[44–46], strawberry plants^[47, 48], tomatoes^[49, 50], lettuce (Beneden et al., 2009)^[51], ornamentals^[52], including *Rosa hybrida*^[53], and *Hibiscus* twigs and branches^[54]. *B. cinerea* caused necrosis on the stems of *Catharanthus roseus*, which later spread to the leaf stalks and, finally, the entire leaf. On the afflicted tissues, the fungus subsequently produce a thin, fragile, gray mycelium, which was especially dispersed on the stems^[55]. Generally, *B. cinerea* infection symptoms include leaf spot; blight; stem canker or rot, and damping-off of young seedlings. Blight is common symptoms begin as small water-soaked lesions or areas^[56, 57] on flower petals, leaves, or stems. Similar to this, numerous crops and ornamental plants are seriously threatened by *Alternaria alternata*, a pathogen that is well-known for its cosmopolitan nature and broad host range. rice seeds^[58], tomato, potato^[59], saffron^[60, 61]. This fungus can also cause latent infections, which can result in post-harvest spoilage on citrus^[62] or pears^[63]. *A. alternata* targets its hosts' leaves, stems, and fruits. On the leaves, lesions manifest as roughly round patches with distinct black patches^[64]. Also, it was reported that this fungus caused different types of rose periwinkle leaf blight in Iraq^[65]. This author claims that the symptoms began as tiny, irregular, round, light brown dots that eventually grew larger and darker to cover the majority of the leaf surface. The brown or black spots on *C. roseus* leaves does not necessarily indicate that *B. cinerea* or *A. alternata* are responsible for these troubles, because other species such as *Curvularia lunata* can also induce the same symptoms. *Curvularia lunata* has been retrieved from leaves of *C. roseus*^[66, 67]. This species has been recorded among the recoverable endophytic fungi colonizing the roots, leaves and stems of *C. roseus*^[68]. Pandey^[69] have reported *Curvularia* sp. among 7 fungal endophytes isolated from the leaves of this plant species. In Morocco, it was detected on aerial parts of *Hibiscus rosa-sinensis*^[54]. Lamrani^[58] noted *C. lunata* as a rice-seed contaminant. The infected tissues manifested elongated or circular brown-to-black lesions on the leaves and stems of certain varieties of rice, which later expand and become merged^[70].

The genus *Fusarium* was well represented in this study, with six identified species. Of them, *Fusarium oxysporum* was the most prevalent fungus. *F. oxysporum* is common in soil and is widely distributed in nature, where it act as a saprophyte or parasite, causing root rot in attractive plants such as *Tulipa gesneriana* L., *Cyclamen persicum*, *Gladiolus callianthus*, *Acacia dealbata*^[71], but it was also recovered as endophytic fungus in *C. roseus*^[72]. Symptoms infection induced by this fungus often manifest as browning of the vessels and yellowing of the leaves, which can result in complete desiccation of the plants^[40]. It results in reddish-brown rot with a wet appearance on the stems, where clusters of spores, often pink in color, emerge in various spots^[73].

Fusarium oxysporum has been reported to affect strawberry in a specialized form known as *F. oxysporum* f. sp. *fragariae*^[74]. It has also been found on banana plants (*Musa accuminata* L.)^[75] and on the stems and branches of *Hibiscus rosa-sinensis*, where it appears as orange-colored domes^[54]. For *Fusarium solani*, it was isolated from *Catharanthus*'s bark in India^[15] and earlier in Malaysia from roots^[16]. Yasir and Almaliky^[76] also stated the presence of *F. solani* alongside *F. equiseti* and *Rhizoctonia solani* when diagnosing the cause of root rot and wilt disease observed on *C. roseus*. This species also attacks crops such as cotton^[77] and has been found in the mycoflora associated with banana plants (*Musa accuminata* L.)^[76]. In Morocco, Rieuf^[78] reported this species as responsible of *Fusarium* attacks in *Geranium rosea*.

However, *Catharanthus roseus* has never been mentioned as a host plant for *F. subglutinans* and *F. nivale* in addition to *Fusarium verticillioides* and *F. chlamyosporium* showing moderate isolation frequencies. This latter fungus could damage seed and affect vigour of rice seedling^[79, 80]. *F. verticillioides* can infect all plant parts of Maize^[81], roots^[82], stems^[83], and ears^[84] during every growth stage.

We can suggest that different pathogens can cause similar symptoms, and the same pathogens can lead to different symptoms depending on the conditions when the disease starts. Many observations pointed out that lesions can be inhabited by both pathogens and saprophytes, such as *Trichoderma harzianum* et *Fusarium* species.

T. harzianum is known as a biological control agent^[85]. In some cases, the antagonistic effect of *T. harzianum* is simply associated with parasitic behavior manifested by

the hyphae wrapping around the filaments of the parasitic fungus^[86]. Numerous studies have demonstrated that *Trichoderma* has interesting potential for controlling various pathogens^[87].

Aspergillus niger was strongly represented with high frequencies on roots and on leaves but was exclusively found on the leaves and roots of the plant under investigation. This particular species is known to cause black crown rot on groundnuts, leading to plant death^[88]. For *Aspergillus flavus*, it was reported in Brazil by^[89] on *Musa* sp., rice (*Oryza sativa*), and coconut (*Cocos nucifera*). This fungus can affect several banana species such as *M. textilis* in the Philippines^[90] and *M. accuminata* in Thailand (Lumyong et al., 2003)^[91]. The third *Aspergillus* species viz, *A. fumigatus* retrieved from vegetative parts, is commonly encountered in air and soil^[92]. The genus *Penicillium* was found on the leaves and roots but recently^[68] have isolated *P. citrinum* associated with endophytic fungi of leaves, stems and roots of *C. roseus*.

Epicoccum nigrum is known to cause discoloration in rice seeds and can also affect their germination capacity^[93].

Although a variety of fungal communities may inhabit in medicinal plants^[94] who have reported a total of 153 fungal isolates which were isolated from thirty leaves of the *C. roseus* plant, including 11 endophytic and 142 exophytic isolates. Other studies revealed the occurrence of endophytic group including *Lasiodiplodia theobromae*, *Cophinforma mamane*, *Phoma putamium* with *F. solani*, *Colletotrichum* sp., *Alternaria longipes*^[15]. However, the expression, the activity and the impact of these organisms vis a vis *C. roseus* are still a concern for debate.

The study indicated that the lesions found on *Catharanthus roseus* are not monospecific, but are instead colonized by a diverse fungal complex. This complex includes several fungi known to be pathogenic while others were categorized as non-pathogenic endophytes. The different organs yielded variable results vis-à-vis the genera of fungal isolates. This study represents the first report of fungal diversity from diseased *Catharanthus roseus* plants in Morocco. However, our findings are insufficient to determine the specific causal agent responsible for the symptomatic areas on the organ surface. Therefore, it is necessary to conduct a precise and adequate characterization of the biology, physiology, and pathogenicity of isolates of these fungi. Examining the evo-

lution of the fungal flora and analyzing the plants before and after plant transplanting could help determine the origin of infections in ornamental plants and facilitate the adoption of more appropriate and timely control methods.

Furthermore, there are many chances to find novel metabolites with potential bioactivity by investigating endophytic fungi that live inside plant species that have therapeutic qualities.

Author Contributions

Conceptualization, A.O.T. and A.D.; methodology, R.B.; software, J.M.; validation, N.A. and M.E.; writing—original draft preparation, N.M.; writing—review and editing, A.D. and A.O.T. All authors have read and agreed to the published version of the manuscript.

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