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**UNIVERSIDADE ESTADUAL PAULISTA “JÚLIO DE MESQUITA FILHO”
CAMPUS DE ILHA SOLTEIRA**

MARIA CLARA SHIROMA BURI

**Triazole-Resistance in *Aspergillus fumigatus* in Brazilian Dry Food: Focus on
Yerba Mate, Green Coffee, and Black Pepper**

Ilha Solteira - SP
2025



PROGRAMA DE PÓS-GRADUAÇÃO EM AGRONOMIA

MARIA CLARA SHIROMA BURI

**Triazole-Resistance in *Aspergillus fumigatus* in Brazilian Dry Food: Focus on
Yerba Mate, Green Coffee, and Black Pepper**

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Orientador: Prof. Dr. Paulo Cezar Ceresini

Coorientadora: Prof^a. Dr^a. Katherin Castro-
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AUTORA: MARIA CLARA SHIROMA BURI

ORIENTADOR: PAULO CEZAR CERESINI

COORDINADORA: TATIANE CARLA SILVA

COORDINADORA: KATHERIN CASTRO RIOS

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PAULO CEZAR CERESINI

Data: 09/12/2025 14:54:12-0300

Verifique em <https://validar.itl.gov.br>

Prof. Dr. PAULO CEZAR CERESINI (Participação Presencial)

Departamento de Fitossanidade, Engenharia Rural e Solos / UNESP / Câmpus de Ilha Solteira - FEIS

Documento assinado digitalmente



EDSON LUIZ FURTADO

Data: 09/12/2025 16:03:18-0300

Verifique em <https://validar.itl.gov.br>

Prof. Dr. EDSON LUIZ FURTADO (Participação Virtual)

Departamento de Proteção Vegetal / UNESP / Câmpus de Botucatu - FCA

Documento assinado digitalmente



KARINE ASSIS COSTA

Data: 09/12/2025 15:54:31-0300

Verifique em <https://validar.itl.gov.br>

Profa. Dra. KARINE ASSIS COSTA (Participação Presencial)

Departamento de Biologia e Zootecnia / UNESP / Câmpus de Ilha Solteira - FEIS

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POTENTIAL IMPACT OF THIS RESEARCH

The expected impact on society should be written succinctly, considering the following aspects: the scientific, technical, social, innovative, economic, educational and cultural potential; internationalization; local, regional and national insertion; sustainable development, knowledge and research themes.

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RESUMO

Aspergillus fumigatus é um fungo saprófito e o principal agente causal da aspergilose, uma doença que geralmente se desenvolve após a inalação de esporos. Embora o ambiente seja reconhecido como a principal fonte de exposição, produtos alimentícios secos podem atuar como reservatórios secundários, pois os esporos podem se aerosolizar durante o manuseio. Este estudo teve como objetivo determinar a prevalência e a sensibilidade aos triazóis em *A. fumigatus* de alimentos secos comuns no Brasil, a fim de avaliar seu potencial papel na exposição humana. Analisamos 25 amostras comerciais de erva-mate, pimenta do reino e café verde, e isolamos *A. fumigatus* em 68% dos produtos, obtendo 42 isolados. Todos os isolados apresentaram valores de EC₅₀ abaixo dos limites clínicos de resistência para tebuconazol (0,04–0,70 µg/mL), itraconazol (0,06–0,59 µg/mL) e voriconazol (0,08–0,15 µg/mL). O sequenciamento do gene CYP51A revelou apenas o polimorfismo M172V, que não está associado à resistência. A genotipagem por microssatélites revelou alta diversidade genética, com 33 genótipos multilocus identificados entre os 42 isolados. A rede de mínima expansão mostrou alta conectividade e estrutura populacional limitada, sugerindo fontes ambientais compartilhadas e troca genética contínua. A elevada prevalência e a substancial diversidade genética de *A. fumigatus* nesses alimentos secos confirmam que eles atuam como carreadores de esporos de origem ambiental. Esses achados ressaltam a importância de uma vigilância integrada em *One Health* para detectar mudanças nas populações de *A. fumigatus* e antecipar riscos emergentes na interface entre agricultura, segurança alimentar e saúde pública.

Palavras-chave: *Aspergillus*; *one health*; alimentos secos; resistência a azóis; diversidade genética.

ABSTRACT

Aspergillus fumigatus is a ubiquitous saprotrophic fungus and the primary etiological agent of aspergillosis, a disease that typically arises following inhalation of airborne spores. Although the environment is recognized as the main source of exposure, dry food products may function as secondary reservoirs, as spores can become aerosolized during handling. This study aimed to determine the prevalence and triazole susceptibility of *A. fumigatus* in common Brazilian dry foods to evaluate their potential contribution to human exposure. We analyzed 25 commercial samples of yerba mate, black pepper, and green coffee, and successfully isolated *A. fumigatus* from 68% of them, yielding 42 isolates. All isolates exhibited EC₅₀ values below clinical resistance thresholds for tebuconazole (0.04–0.70 µg/mL), itraconazole (0.06–0.59 µg/mL), and voriconazole (0.08–0.15 µg/mL). CYP51A sequencing revealed only the M172V polymorphism, which is not associated with azole resistance. Microsatellite genotyping uncovered high genetic diversity, identifying 33 multilocus genotypes among the 42 isolates. A minimum-spanning network showed high connectivity and limited population structure, suggesting shared environmental sources and ongoing genetic exchange. The high prevalence and substantial genetic diversity of *A. fumigatus* in these dry foods confirm their role as carriers of environmentally derived spores. These findings underscore the importance of integrated *One Health* surveillance to detect shifts in *A. fumigatus* populations and anticipate emerging risks at the intersection of agriculture, food safety, and public health.

Keywords: *Aspergillus*; *one health*; dry food; azole resistance; genetic diversity.

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LIST OF ABBREVIATIONS AND ACRONYMS

IA	Invasive Aspergillosis
PDA	Potato Dextrose Agar
DNA	Deoxyribonucleic Acid
PCR	Polymerase Chain Reaction
EC ₅₀	Effective Concentration Required to Inhibit 50% of Fungal Growth
STRs	Short Tandem Repeats
M13F	Primer M13 Forward
FAM	Fluorescein Amidite
TET	Tetrachloro-fluorescein
Cy3	Cyanine 3
Cy5	Cyanine 5
MLGs	Unique Multilocus Genotypes
MSN	Minimum Spanning Network
EUCAST	European Committee on Antimicrobial Susceptibility Testing
MIC	Minimum Inhibitory Concentration
ATU	Area of Technical Uncertainty
CPA	Chronic Pulmonary Aspergillosis
IPA	Invasive Pulmonary Aspergillosis
NWT	Non-Wild-Type
ARAf	Azole-Resistant <i>Aspergillus fumigatus</i>

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1. INTRODUCTION

Deeply rooted in history, culture, and culinary traditions, the spices, teas, and coffee hold enduring significance (Musalib et al., 2022). Among them, green coffee (*Coffea*) black pepper (*Piper nigrum* L.) and yerba mate (*Ilex paraguariensis*) have become increasingly prominent in consumer's daily lives (García-Cordero et al., 2023; Song et al., 2025). These products are valued not only for their distinctive flavors and sensory properties but also for their various health benefits, such as antioxidant activity, antiinflammatory properties, and metabolic support (Dellalibera; Lemaire; Lafay, 2006; Santiano et al., 2023). Amid a growing interest in healthier lifestyles, the demand for natural and functional foods has driven the consumption of these products. Furthermore, green coffee, black pepper, and yerba mate hold significant socioeconomic relevance in Brazil, serving as important sources of income for thousands of families and playing a key role in the national agribusiness sector (Costa; Pauletto, 2021; Maguire-Rajpaul et al., 2020).

The black pepper (*Piper nigrum* L.) is a species of a climbing vine native to Asia whose fruit offers differentiated aroma and flavor, which underlies its use as a spice (Hammouti et al., 2019). In Brazil, the introduction of the crop occurred only in 1930 in the Amazon region; since then, due to its economic importance and high market value, the condiment has become an opportunity, especially for family farmers (Moreira et al., 2021). This spice contains approximately 5–9% of piperamide alkaloids, primarily accumulating in the fruit peel and seeds that piperine stands out as the most biologically relevant compound, traditionally utilized for medicinal purposes, it demonstrates multiple pharmacological and ethnomedicinal properties (Batubara; Rafi; Yolanda, 2020; Dewi et al., 2025) including the relief of pain, digestive disorders, and fever (Ansari et al., 2023), the treatment of asthma (Usmani; Jain; Yadav, 2023), and its role in supporting anticancer therapies (Wiraswati et al., 2025).

Brazil stands as the world's foremost coffee producer, a position underpinned by its exceptionally favorable agroclimatic conditions. Currently, an estimated 330.000 rural establishments are dedicated to coffee cultivation, approximately 78% of which are classified as family farms, wherein coffee constitutes the primary source of livelihood (Santana et al., 2025). Furthermore, Coffee is a notable source of bioactive compounds, including chlorogenic acids, caffeine, cafestol, and kahweol. While it is predominantly consumed in its roasted form, green coffee retains significantly higher

levels of chlorogenic acids, which have been linked to antioxidant and antiinflammatory effects. Consequently, green coffee has garnered increasing interest in scientific research and the nutraceutical sector due to its superior preservation of functional properties compared to roasted coffee beans (Humaryanto; Ave, 2019; Santana et al., 2025).

Just like green coffee, yerba mate (*Ilex paraguariensis*) is traditionally used in “Erva mate”, (is a cold aqueous infusion widely consumed in southern Brazil, Paraguay, and Argentina) notably rich in chlorogenic acids and has been garnering increasing scholarly and commercial interest due to its diverse array of bioactive properties (Kautzmann et al., 2024). Prepared from dried leaves of yerba mate and typically served in a “cuia”— a cup typically lined with leather or made from bull horn — with cold water, it offers a refreshing counterpart to chimarrão, the hot infusion made from the same plant. Owing to its complex phytochemical composition. Composing flavonoids, caffeine, theophylline, chlorogenic acids, saponins, vitamins, and trace elements, yerba mate demonstrates a broad spectrum of health-promoting properties. These include potent antioxidant capacity, metabolic stimulation, and antiinflammatory activity, positioning it as a beverage of growing relevance within the context of functional foods and nutraceuticals (Proch et al., 2023; Silvana et al., 2020).

Even though these products are sources of many desirable substances, they may also contain toxic metals (Chore et al., 2024; Górka; Baran; Słowik-Borowiec, 2025), pesticide residues above regulatory limits (Da Silva et al., 2022; Kandaswamy et al., 2021; Nolasco et al., 2022), mycotoxins such as aflatoxin B1 and ochratoxin A (Al Khoury et al., 2022; Attiya et al., 2021; Costa da Silva et al., 2021; Demirhan; Demirhan, 2023) and microorganism harmful to health (Ildris et al., 2020; KoohyKamaly-Dehkordy et al., 2013). Dry food products, such as spices, teas, and coffee beans, are often contaminated with high levels of microorganisms (Maman et al., 2021). This contamination can occur at various stages of the production chain and may involve bacteria, viruses, and fungi, which pose significant health risks, particularly when these products are used in the preparation of ready-to-eat foods (Gryczka et al., 2020). Among pathogenic fungi, the genus *Aspergillus* stands out for including xerophilic and thermotolerant species capable of thriving in low-moisture and hightemperature environments. Some of these species not only produce mycotoxins but can also act as opportunistic pathogens in humans, as is the case with *Aspergillus*

fumigatus (Abdel-Azeem et al., 2016; Brandl; Andersen, 2017). The clinical importance of *Aspergillus fumigatus* lies in its ability to cause a wide range of respiratory conditions, from localized airway inflammation to severe infections, notably invasive aspergillosis (IA). Although this cosmopolitan fungus is not typically a primary human pathogen, it has emerged as a significant opportunistic agent, particularly due to the increasing prevalence of conditions that require immunosuppressive therapies, including but not limited to cancer and organ transplantation (Resendiz-Sharpe et al., 2021). Aspergillosis is an environmentally acquired respiratory disease that encompasses a spectrum of clinical and radiological manifestations, largely determined by the host's immune status (Koren Fernández et al., 2014). It is estimated that approximately 4.8 million people worldwide are affected by some form of aspergillosis, with an estimated 300,000 cases of IA occurring annually, and associated mortality rates ranging between 30% and 95% (Brown et al., 2012; Garcia et al., 2018; Wang et al., 2022).

The first-line treatment for aspergillosis relies primarily on triazole-based antifungal agents, with itraconazole, voriconazole, and posaconazole being the most commonly used (Russo et al., 2020). However, increasing reports of reduced efficacy have been linked to the emergence of triazole-resistant *A. fumigatus* strains. These resistant isolates often harbor mutations in the CYP51 gene, which encodes lanosterol 14 α -demethylase, an essential enzyme in the ergosterol biosynthesis pathway, responsible for producing the primary sterol component of the fungal cell membrane (Simon et al., 2021; Sueth-Santiago et al., 2015). Furthermore, resistance mechanisms to triazoles in *Aspergillus spp.* can be broadly categorized into two types: those associated with CYP51A alterations and those that are independent of this gene (Garcia-Rubio; Cuenca-Estrella; Mellado, 2017).

CYP51A-related resistance mechanisms are the most widespread globally (Macedo et al., 2021). Since their first identification in 1998 at a hospital in the Netherlands, an increasing number of mutations in the CYP51A gene have been reported. These alterations have been found in *A. fumigatus* isolates from patients undergoing long-term triazole therapy, as well as from triazole-naïve patients and environmental samples (Gyurtane Szabo et al., 2023; Snelders et al., 2008). Mutations within the coding sequence of the CYP51A gene such as TR34/L98H, TR34/L98H/S297T/F495I, TR46/Y121F/T289A, and TR53 are commonly associated with pan-azole resistance, conferring reduced susceptibility to all triazole antifungals

(Lucio et al., 2022). Additionally, certain amino acid substitutions, including M220, G54, and G138, have been shown to impair drug–target interaction by diminishing the binding affinity between the antifungal agent and the fungal enzyme (Howard et al., 2009; Verweij et al., 2016). In contrast, other substitutions such as F46Y, M127V, and E427K are typically linked to lower levels of azole resistance (Verweij et al., 2020).

The detection of TR34/L98H and TR46/Y121F alleles in both environmental and clinical isolates of *A. fumigatus* suggests that CYP51A mutations may arise not only in response to prolonged antifungal therapy, but also as a consequence of triazole fungicide application in agriculture, given the structural similarity between medical and agricultural triazoles (Burks et al., 2021; Kang et al., 2021). Although the use of such fungicides is restricted in certain crops—such as black pepper and yerba mate, which are the focus of our investigation—adjacent areas under intensive agricultural practices may act as resistance *hotspots*. Strains selected under fungicide pressure in these regions can subsequently contaminate nearby crops at various points in the production chain. Moreover, the study by Cui et al. (2019) demonstrated that preexposure to the agricultural fungicide tebuconazole in *A. fumigatus* isolates can induce cross-resistance to itraconazole and voriconazole. Therefore, the emergence of azoleresistant *A. fumigatus* strains in the environment should be regarded as an unintended consequence of triazole use in agriculture, especially considering that this fungal species is not a plant pathogen. Although *Aspergillus fumigatus* strains resistant to triazoles have been isolated from soil, plant debris, air, and compost samples (Kang et al., 2021; Kano et al., 2015), only the study by Viegas et al. (2023) reported isolates from the Fumigati section resistant to posaconazole in tea samples. Nonetheless, this study did not investigate potential mutations associated with the observed resistance profile. Yet, the occurrence of *Aspergillus fumigatus* in food products is frequent, posing a potential risk (Castro-Ríos et al., 2025).

Accordingly, the aim of this study was to investigate the resistance status of *Aspergillus fumigatus* isolates sampled from commercial products of yerba mate, black pepper, and green coffee to the medical azoles itraconazole and voriconazole, and to the agricultural azole tebuconazole. We compared the phenotypic and genotypic profiles of fungicide susceptibility. Additionally, we performed sequencing of genes encoding fungicide target proteins, and microsatellite typing based on short tandem

environmental populations, underscoring the potential role of foods as reservoirs for genetically diverse strains that may also circulate in human environments.

5. CONCLUSION

Despite the absence of phenotypically resistant isolates, our study demonstrates that *Aspergillus fumigatus* is consistently present in minimally processed plant products commonly consumed in Brazil, including black pepper, yerba mate, and green coffee. The variability observed in EC_{50} values for tebuconazole and itraconazole, particularly among isolates from plant-based substrates, suggests early shifts in susceptibility that may reflect environmental selective pressures rather than established resistance. Although voriconazole susceptibility remained uniform across isolates, the presence of subtle differences in sensitivity highlights the importance of continuous monitoring, especially considering the fungus's remarkable adaptive capacity.

Genetic characterization revealed high multilocus genotypic diversity among isolates, alongside evidence of clonal expansion within specific substrates. The coexistence of unique and recurrent genotypes, coupled with the absence of major CYP51A mutations, underscores the evolutionary plasticity of *A. fumigatus* populations circulating in food products. Environmental exposure to agricultural triazoles, whether through fungicide drift, soil contamination, or indirect contact in production environments, may be driving early-stage adaptive responses. These findings reinforce that even food items not directly treated with azoles can act as reservoirs for diverse fungal populations that mirror patterns previously reported in clinical and environmental settings worldwide.

From a One Health perspective, the detection of *A. fumigatus* across multiple food matrices, combined with its genetic diversity and evidence of early susceptibility shifts, highlights the need for integrated surveillance strategies spanning agricultural, environmental, and clinical sectors. Ensuring safe production chains, rational fungicide use, and harmonized monitoring policies is essential to mitigate the risk of selecting for azole-resistant fungal populations. As global consumption of plant-derived foods continues to grow, proactive surveillance of *A. fumigatus* in food commodities represents a critical step toward safeguarding public health, supporting sustainable

agriculture, and reducing the potential emergence and dissemination of triazoleresistant strains.

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