## Collaborative One Health approaches can mitigate increasing azole-resistant Aspergillus fumigatus in Africa







Aspergillus fumigatus is a species of fungus that is commonly found in the environment, household dust, soil, and decaying plant matter. The fungus produces spores which are inhaled by humans daily, although the host defence mechanisms of most people limit inflammatory responses and promote fungal clearance. Inhalation of the spores of A fumigatus triggers a wide spectrum of diseases in immunocompromised individuals, depending on the patient group affected and the pathways of pathogenesis. Forms of aspergillosis that are becoming a major public health issue include invasive aspergillosis, allergic bronchopulmonary aspergillosis, and chronic pulmonary aspergillosis.

Estimates suggest that approximately 2.2 million people suffer from invasive aspergillosis annually worldwide,1 and it mostly occurs in immunocompromised individuals with chronic obstructive pulmonary disease, tuberculosis, or leukaemia, and more recently, as a complication of COVID-19. Left untreated, invasive aspergillosis is fatal.

Triazole antifungals are used for the prophylaxis and treatment of invasive fungal infections, acting by inhibiting the ergosterol biosynthesis pathway and disrupting the fungal cell membrane. Surveillance studies from various geographical locations have reported an increase in triazole-resistant A fumigatus (TRAF), raising concerns about the management of diseases caused by the fungus.2

The underlying resistance mechanism commonly involves point mutations in the cyp51A gene, with occasional non-cyp51 mutations. TRAF precludes the use of oral antifungals for treating aspergillosis, leaving only intravenous options, such as amphotericin B and echinocandins, which are not available in many African countries. The growing reports of TRAF have been linked to the use of fungicides in agriculture, with the development of cross-resistance to medical triazoles.

There have been several reports of TRAF from the eastern and western African subregions following the first report from Tanzania in 2014 (appendix).3-10 In Tanzania and Kenya, TRAF was identified in both environmental and clinical samples, with isolation rates ranging from 14% to 27%.3-7 In one study, all five clinical

isolates of A fumigatus were triazole resistant, giving a prevalence of 100%,4 with the TR34/L98H mutation being the predominant resistance phenotype in soils in Tanzania and Kenya. The TR46/Y121F/T289A mutation has also been detected in Tanzania, with all six clinical isolates seen so far-predominantly in cases of otitis media—bearing the TR34/L98H mutation (appendix).

Chowdhary and colleagues<sup>3</sup> found a genetic relatedness of Tanzanian TR46/Y121F/T289A strains to Dutch isolates, and the TR34/L98H isolates were identical to the Indian TR34/L98H genotype.37 These similarities in molecular epidemiology suggest possible migration of isolates harbouring resistance traits from distant locales. In Kenya, although the prevalence of TRAF was higher in fungicide-experienced soil, TRAF was also present in fungicide-naive soil samples.5 This finding implies that TRAF might spread locally from areas of fungicide use to places where fungicides have not been used.

More recently, TRAF has been isolated from environmental samples in Burkina Faso and Nigeria. 9,10 Unlike in east Africa, TRAF isolation rates have been low, ranging from 2.0% to 2.2%, and tandem repeat mutations have not been identified. The cyp51A gene mutations detected in TRAF from Burkina Faso and Nigeria were F46Y/M172V/E427K and M172V respectively (appendix).9,10

Despite the absence of surveillance data on TRAF from other African countries, there is a possibility that TRAF is more widespread, given the extensive use of azole fungicides for agricultural purposes in the continent.9 The inherent ability of A fumigatus to sporulate and survive in almost any environment facilitates its dispersal across long distances, with potential for transmission across national boundaries.

Given the interconnectedness of humans, plants, and animals in shared environments, tackling the problem of TRAF in Africa will require innovative, collaborative, multisectoral, and transdisciplinary One Health approaches. Greater collaboration between See Online for appendix the agricultural and health sectors within Africa is urgently required. One Health policy adoptions and greater funding by African governments will catalyse

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coordinated data sharing, interdisciplinary surveillance, and laboratory collaboration on TRAF. Existing public-private partnerships such as the Africa One Health University Network could be leveraged to promote resource sharing and economies of scale.

Community engagement and advocacy are crucial to promote the adoption of safer and sustainable agricultural practices. Alternative strategies community practices such as the use of azole fungicides should be promoted, along with a more judicious use of triazoles in clinical settings. Chemosensitisation, which involves the use of natural compounds that interact synergistically with antifungals thereby lowering effective dosages and negative effects, holds great promise in this regard. Coordinated capacity-building and research are crucial for identifying and responding to clinical and environmental TRAF in Africa. Urgent action is required to address TRAF and its attendant challenges in Africa, and adopting One Health approaches might be the turning point.

We declare no competing interests.

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- 1 Hammond EE, McDonald CS, Vestbo J, Denning DW. The global impact of Aspergillus infection on COPD. BMC Pulm Med 2020; 20: 241.
- 2 Lestrade PP, Bentvelsen RG, Schauwvlieghe AFAD, et al. Voriconazole resistance and mortality in invasive aspergillosis: a multicenter retrospective cohort study. Clin Infect Dis 2019; 68: 1463–71.
- 3 Chowdhary A, Sharma C, van den Boom M, et al. Multi-azole-resistant Aspergillus fumigatus in the environment in Tanzania. J Antimicrob Chemother 2014; 69: 2979–83.
- 4 Mushi MF, Buname G, Bader O, Groß U, Mshana SE. Aspergillus fumigatus carrying TR34/L98H resistance allele causing complicated suppurative otitis media in Tanzania: call for improved diagnosis of fungi in sub-Saharan Africa. BMC Infect Dis 2016; 16: 464.
- Kemoi EK, Nyerere A, Gross U, Bader O, Gonoi T, Bii CC. Diversity of azole resistant Aspergillus species isolated from experienced and naive soils in Nairobi county and Naivasha sub-county Kenya. Eur Sci J 2017; 13: 201-11
- 6 Kemoi EK, Nyerere A, Bii CC. Triazole-resistant Aspergillus fumigatus from fungicide-experienced soils in Naivasha subcounty and Nairobi county, Kenya. Int J Microbiol 2018; 2018: 7147938.
- 7 Sharma C, Hagen F, Moroti R, Meis JF, Chowdhary A. Triazole-resistant Aspergillus fumigatus harbouring G54 mutation: is it de novo or environmentally acquired? J Glob Antimicrob Resist 2015; 3: 69–74.
- Ashu EE, Korfanty GA, Xu J. Evidence of unique genetic diversity in Aspergillus fumigatus isolates from Cameroon. Mycoses 2017; 60: 739-48.
- 9 Yerbanga IW, Resendiz-Sharpe A, Bamba S, et al. First investigative study of azole-resistant Aspergillus fumigatus in the environment in Burkina Faso. Int J Environ Res Public Health 2021; 18: 2250.
- 10 Resendiz-Sharpe A, Dewaele K, Merckx R, et al. Triazole-resistance in environmental Aspergillus fumigatus in Latin American and African countries. J Fungi (Basel) 2021; 7: 292.