

Inbred to well-fed: Maize breeding efforts to improve food security and safety in Africa

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A, Maize leaves B, plated on technical agar amended with 6-benzyl aminopurine in a crisper box before inoculation. This detached-leaf assay allows evaluation of plant–pathogen interactions for up to 30 days. Credit: *Plant Disease* (2022). DOI: 10.1094/PDIS-12-21-2788-RE

Over 300 million people in Sub-Saharan Africa (SSA) depend on maize for nearly 30% of their caloric intake, and the popularity of maize is expected to continue its upward trend (IITA). As the demand for maize increases, so does the need for sustainable and safe production, yet numerous biotic and abiotic stressors threaten this staple crop in SSA.

Researchers from the International Institute of Tropical Agriculture and



Obafemi Awolowo University in Nigeria conducted a recent study to combat these threats. The study, newly published in *Plant Disease*, offers a promising strategy to protect food security and safety of <u>maize</u> in SSA.

Faith Bankole and colleagues set out to find maize cultivars that could resist multiple diseases, which would benefit <u>smallholder farmers</u> and increase the production of quality maize for food and feed. Using rapid, efficient, and cost-effective laboratory methods, they screened over 100 maize genotypes for resistance to foliar diseases and <u>aflatoxin</u> <u>contamination</u>—finding seven early-maturing (EM) and six extra-early (EEM) maize inbred lines with superior resistance to three major pathogens, plus ten inbreds with resistance that also had significantly reduced aflatoxin accumulation.

The EM and EEM inbreds with multiple <u>disease resistance</u> (MDR) discovered in this study are being tested extensively in hybrid combinations. Hybrids with good agronomic attributes, high yield, and MDR will be released for commercialization. Smallholder farmers that implement the use of these maize hybrids can reliably produce greater, safer yields.







Representative reactions of maize inbreds to three foliar pathogens (Exserohilum turcicum, Bipolaris maydis, and Curvularia lunata) using the detached leaf assay. Inbreds were classified as follows: HR = highly resistant, R = resistant, MR = moderately resistant, S = susceptible, and HS = highly susceptible. Photographs were taken at 10 days after inoculation. Inbreds in the E. turcicum row from left to right are TZEI 56, TZEI 32, TZEI 1, TZEI 120, and TZEI 134; those in the B. maydis row are TZEEI 63, TZEEI 75, TZEEI 79, TZEEI 38, and TZEEI 67; and those in the C. lunata row are TZEI 56, TZEI 56, TZEI 51, TZEI 1, TZEI 17, and TZEI 134. Credit: *Plant Disease* (2022). DOI: 10.1094/PDIS-12-21-2788-RE

"Both EM and EEM maize germplasm have not been exploited as potential sources of resistance to multiple diseases," corresponding author Alejandro Ortega-Beltran comments. "Our results demonstrate that these genotypes can significantly contribute to breeding programs to tackle multiple diseases, benefitting tropical and subtropical regions."

According to the authors, large-scale use of these maize genotypes can potentially increase safe maize production and productivity plus reduce losses caused by aflatoxin contamination—ultimately contributing to human health and livelihood in SSA.

More information: Faith A. Bankole et al, Identification of Early and Extra-Early Maturing Tropical Maize Inbred Lines with Multiple Disease Resistance for Enhanced Maize Production and Productivity in Sub-Saharan Africa, *Plant Disease* (2022). DOI: 10.1094/PDIS-12-21-2788-RE

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