

ABSTRACT

Molecular identification and pathological characterisation of fungal pathogens associated with wilting of eggplants (*Solanum* spp.) is a necessary precondition for efficient disease management and for the development of durable host resistance in eggplant cultivars. A study was carried out to identify and differentiate *Fusarium* isolates from eggplant fields in the Arumeru district, Tanzania, based on morphological characters, molecular sequence data of the ITS and elongation factor 1-alpha gene exons and introns and pathogenicity tests. Twenty-nine *Fusarium* isolates were studied. They belonged to the *Fusarium oxysporum*- (52 %), *F. solani*- (24 %) and *F. incarnatum-equiseti* (24 %) species complexes. Pathogenicity tests indicated that representatives of all three species groups were pathogenic to eggplants. The tested members of *F. incarnatum-equiseti* and *F. oxysporum* were pathogenic to African eggplant (*S. macrocarpon*, MM 1131) and *S. anguivi* (N 19). *Fusarium solani* caused wilting in *S. anguivi*. Two isolates were identified as *Fusarium oxysporum* f. sp. *melongenae* (FOM). This is the first report of *Fusarium oxysporum* f.sp *melongenae* and *F. solani* causing wilting of eggplant in Tanzania and the first report of FIESC causing wilting in African eggplant.

Key words: *Fusarium equiseti*, *F.oxysporum*, *F. solani*, *Fusarium* wilt, *Solanum* species