



A multiplex-PCR assay for identification of the quarantine plant pathogen *Xanthomonas axonopodis* pv. *phaseoli*

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Auteur	Boureau, Tristan [1], Kerkoud, Mohammed [2], Chhel, Fabien [3], Hunault, Gilles [4], Darrasse, Armelle [5], Brin, Chrystelle [6], Durand, K. [7], Hajri, A. [8], Poussier, S. [9], Manceau, C. [10], Lardeux, Frédéric [11], Saubion, Frédéric [12], Jacques, Marie-Agnès [13]
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Résumé en anglais	<p>In this study we developed an algorithm to screen for all exact molecular signatures of the quarantine pathogen <i>Xanthomonas axonopodis</i> pv. <i>phaseoli</i> (Xap), based on available data of the presence or absence of virulence-associated genes. The simultaneous presence of genes <i>avrBsT</i> and <i>xopL</i> is specific to Xap. Therefore we developed a multiplex PCR assay targeting <i>avrBsT</i> and <i>xopL</i> for the molecular identification of Xap. The specificity of this multiplex was validated by comparison to that of other molecular identification assays aimed at Xap, on a wide collection of reference strains. This multiplex was further validated on a blind collection of <i>Xanthomonas</i> isolates for which pathogenicity was assayed by stem wounding and by dipping leaves into calibrated inocula. This multiplex was combined to the previously described X4c/X4e molecular identification assay for Xap. Such a combination enables the molecular identification of all strains of <i>Xanthomonas</i> pathogenic on bean. Results also show that assay by stem wounding does not give reliable results in the case of Xap, and that pathogenicity assays by dipping should be preferred.</p>
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Liens

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