

Article supplementary materials

Supplementary materials belonging to

Trends In Molecular Diagnosis And Diversity Studies For Phytosanitary Regulated *Xanthomonas*

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Supplementary Table 2: Characteristics of MLVA schemes developed for *Xanthomonas* spp.

Target	Number of TR loci	Range of repeat unit sizes (bp)	Reference
<i>X. albilineans</i>	15	3-36	[1]
<i>X. arboricola</i>	26	7-20	[2]
<i>X. arboricola</i> pv. <i>pruni</i>	6	6-20	[3]
<i>X. arboricola</i> pv. <i>pruni</i>	23	6-116	[4]
<i>X. citri</i> pv. <i>citri</i>	14	6-7	[5]
<i>X. citri</i> pv. <i>citri</i>	31	10-217	[6]
<i>X. citri</i> pv. <i>mangiferaeindicae</i>	12	6-7	[7]
<i>X. citri</i> pv. <i>viticola</i>	8	7	[8]
<i>X. fragariae</i>	36	3-33	[9]
<i>X. euvesicatoria</i>	16	6-8	[10]
<i>X. oryzae</i>	16	6-12	[11]
<i>X. oryzae</i> pv. <i>oryzicola</i>	25	5-9	[12]
<i>X. phaseoli</i> pv. <i>manihotis</i>	22	6-26	[13]
<i>X. phaseoli</i> pv. <i>manihotis</i>	15	6-7	[14]
<i>X. vasicola</i> pv. <i>musacearum</i>	19	6-12	[15]

Genotyping schemes analysing a relatively large number of TR loci are desirable as they allow to partially control their homoplastic evolution [16].

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