Pectin methylesterases contribute the pathogenic differences between races 1 and 4 of *Fusarium oxysporum* f. sp.*cubense*

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Table S1 Primers for quantitative real-time PCR

Gene	GeneBank number	Primer sequences (Forward/ Reverse)	Product size (bp)
MaPME1	FJ264505.1	GCTACCAAGACACCCTCTAC	94
		GTACAAGTTGCAGTTCTGGAGC	04
RPS2	HQ853246.1	TAGGGATTCCGACGATTTGTTT	122
		TAGCGTCATCATTGGCTGGGA	122
CHS2	XM_009405523.2	GCATCACCAGGAGCAACCAC	107
		CATGTACGCGCCGATGTTAG	137
СНЅЗ	XM_009406457.2	CGGCAGGTGCTGAGCGAGTA	101
		GTGGTTGCCTTTCCGTCCTC	
CalS5	XM_018828485.1	GACCTTATAAGTGACAGGAAAGAGATGG	175
		CACAGATGCGTTTCCACAGATCAGAATC	175
PAL	XM_009419417.2	ACATATCCTAAGAAGCCCCACCACC	206
		TCGTCAAAGTGGGCACAAGAGAG	200
JAZ	XM_009411294.2	TGCAAGAGGTTCATCATAGGACAGT	109
		TCTTCAGACATGGAAGTGGCCTC	190
GST	XM_009409198.2	CCTTCAGATGTTCTTCCAGTGCATG	185
		TGTTATCACGGCGATCATTGAGGAG	

Table S2 Antibodies recognizing different HGs and their antigens

Antibody	Antigen	Secondary antibody	
JIM5 ¹	Partially methyl-esterified HG epitope:		
	unesterified and partially esterified residues	Anti-rat-IgG-FITC	
	(up to 40%)		
JIM7 ¹	Partially methyl-esterified HG epitope:		
	methyl-esterified residues (up to 80%)	Anti-rat-igG-FITC	
LM19 ²	Low methyl-esterified HG	Anti-rat-IgG-FITC	
LM20 ²	Highly methyl-esterified HG	Anti-rat-IgG-FITC	
2F4 ³	Un-esterified/Calcium ion cross-linked HG	Anti-mouse-IgG-FITC	



Figure S1 The relative fluorescence intensity of PME in banana (*Musa* spp. AAA) roots after infection with *Fusarium oxysporum* f. sp.*cubense*. *Foc*1: *Fusarium oxysporum* f. sp.*cubense* race 1; *Foc*4: *Fusarium oxysporum* f. sp.*cubense* race 4. For quantification of fluorescence signal, the mean relative fluorescence intensity (n =3 sections of roots) was measured with ZEN 2012 software.







Figure S3 The relative fluorescence intensity of pectins recognized by LM19 in banana (*Musa* spp. AAA) roots after infection with *Fusarium oxysporum* f. sp.*cubense*.













sections of roots) was measured with ZEN 2012 software.

References:

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