

Author's Correction

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Molecular Characterization of a dsRNA Mycovirus, *Fusarium graminearum* Virus-DK21, which Is Phylogenetically Related to Hypoviruses but Has a Genome Organization and Gene Expression Strategy Resembling Those of Plant Potex-like Viruses

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In the article by Kwon et al. that appears in the journal (*Mol. Cells* 23, 304–315, June 30, 2007), an overall text should be revised as written below because the released full length sequences of FgV-DK21 in this paper has a changed information.

Page 304, The ninth line in the Abstract section should be changed “The FgV-DK21 dsRNA consists of 6621 nucleotides” instead of “The FgV-DK21 dsRNA consists of 6624 nucleotides”.

Page 304, The twelfth line in the Abstract section should be changed “respectively, and four putative open reading frames.” instead of “respectively, and five putative open reading frames”.

Page 304, The twentieth line in the Abstract section should be changed “deduced amino acid sequences of ORFs 2 through 4 revealed” instead of “deduced amino acid sequences of ORFs 2 through 5 revealed”.

Page 306, the forty-seventh line of left column in the Materials and Methods section should be changed “which are specific for putative ORFs 1, 2-3 and 4” instead of “which are specific for putative ORFs 1, 2-3 and 5”.

Page 306, the sixth line of right column in the Materials and Methods section should be changed “ORF2 (nt 4833 to 4849), ORF4 (nt 5366 to 5383).” instead of “ORF2 (nt 4833 to 4849) and ORF4 (nt 5366 to 5383), and ORF5 (nt 5525 to 5542)”.

Page 308, the third line of right column in the Result section should be changed “through 4 were screened” instead of “through 5 were screened”.

Page 308, the fiftieth-first line of right column in the Result section should be changed “by ORFs 2, and ORF4 are synthesized through” instead of “by ORFs 2, 4, and 5 are synthesized through”.

Page 309, the second line of left column in the Result section should be changed “ORF2 and ORF4 containing sequences complementary to nts 4833 to 4849, 5366 to 5383, respectively.” instead of “ORF2, ORF4, and ORF5 containing sequences complementary to nts 4833 to 4849, 5525 to 5542, and 5525 to 5542, respectively”.

Page 309, a sentence in the tenth line of right column in the Result section should be deleted. (In primer extension with a primer specific for ORF5, no prevalent extension products were observed in any FgV-DK21-infected isolate, even though a similar consensus sequence is present (Fig. 5C)).

Page 311, the fourteenth line of right column in the Discussion section should be changed “ORFs 2-4 encode putative proteins of 16.7, 6.2, and 48.4 kDa.” instead of “ORFs 2-5 encode putative proteins of 16.7, 6.2, 4.8, and 41.3 kDa”.

Page 312, the fourteenth line of left column in the Discussion section should be changed “ORFs 2 through 4 identified the conserved 17-nt sequence” instead of “ORFs 2 through 5 identified the conserved 17-nt sequence”.

Page 312, a sentence in the eighteenth line of left column in the Discussion section should be deleted. (A relatively weak promoter sequence was also observed upstream of ORF5, but none was observed upstream of ORF3).

Page 313, the twelfth line of left column in the Discussion section should be changed “ORF2 and ORF4, are not present upstream of ORF3 (Fig. 4B).” instead of “ORF2, ORF4, and ORF5 are not present upstream of ORF3 (Fig. 4B)”.

Page 313, the twentieth line of left panel in the Discussion section should be changed “Whether ORFs 2 through 4 actually” instead of “Whether ORFs 2 through

5 actually”.

Page 311, the legend of Fig. 3 should be changed “and are specific for putative ORF1 (p1) ORF2-3 (p2) and ORF4 (p3)” instead of “and are specific for putative ORF1 (p1) ORF2-3 (p2) and ORF5 (p3)”.

Page 312, the legend of Fig. 5 should be changed “to ORF2 (nt 4833 to 4849; panel A) and ORF4 (nt 5366 to 5383; panel B) were annealed” instead of “to ORF2 (nt 4833 to 4849; panel A), ORF4 (nt 5366 to 5383; panel B) and ORF5 (nt 5525 to 5542; panel C) were annealed”.

Figures should be changed as below.

Fig. 1

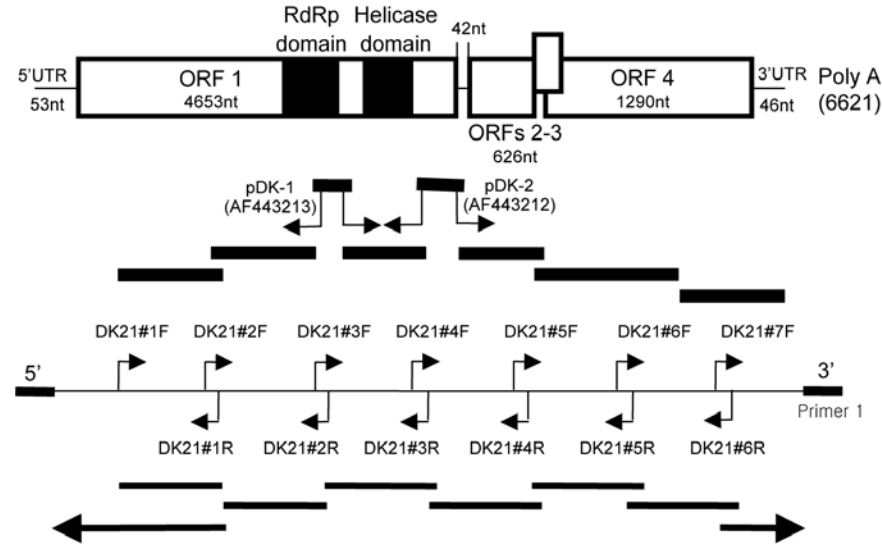


Fig. 4

A				B	
	-6	-3	+4	Putative sgRNA promoter sequences	
Kozak	GCC <u>CCA</u> <u>AUGG</u>			consensus	GAAAGCCCU CGGGCUAG
ORF 1	CCUGAG <u>AUGC</u>			ORF2	GAAAGCCCU CGGGCUAG -12 nt- AUG
ORF 2	GCCAAA <u>AUGG</u>			ORF3	na
ORF 3	CCCCUA <u>AUGA</u>			ORF4	GAAAGCCCU CGGGCUAG -11 nt- AUG
ORF 4	CACACG <u>AUGA</u>				

Fig. 5

