Molecular characterization of sweet potato leaf curl virus isolate from China (SPLCV-CN) and its phylogenetic relationship with other members of the Geminiviridae

Yu Shi Luan · Juan Zhang · Dan Mei Liu · Wen Li Li

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Erratum to: Virus Genes
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- 1. On 3rd page, left column, bottom line 8, 2771 should be changed to 2788.
- 2. On 4th page, right column, middle, 3rd paragraph, bottom line 2, AC1(89%) should be changed to AC1(86%), bottom line 7, AC1(87%) should be changed to AC1(84%).
- 3. Table 2, column 3, identity of AC1 and column 5, similarity of AC1 should be read as follows:

Identity		Similarity	
AC1	AV1	AC1	AV1
68%		68%	
70%		73%	
72%		72%	
72%		70%	
70%		70%	
70%		70%	
69%		69%	
69%		69%	
70%		71%	
62%		62%	
61%		61%	
63%		62%	
64%		63%	
66%		66%	
66%		64%	
66%		66%	
62%		63%	

Table 3, column 3, percent identity (nt/aa) of AC1 should be read as follows:

The online version of the original article can be found under doi:10.1007/s11262-007-0084-1.

Y. S. Luan · J. Zhang · D. M. Liu · W. L. Li (✉)
Department of Bioscience and Bioengineering, Dalian University of Technology, Dalian 116024, P.R. China
e-mail: liwenli2002@yahoo.com

D. M. Liu
Eastern Liaoning University, Dandong 118001, P.R. China

AC1	AC2	AC3	AC4
84%/77%			
85%/86%			
86%/84%			

4. Fig. 1, Genome organization of the SPLCV-CN component A, the numbers representing the position of

nucleotide on the genome is incorrect, the corrected figure is shown here.