

Whole-Genome Characterization of Alfalfa Mosaic Virus Obtained from Metagenomic Analysis of *Vinca minor* and *Wisteria sinensis* in Iran: with Implications for the Genetic Structure of the Virus

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In the first line of the “Materials and Methods” section, the date of collecting samples should only be “In July 2020”, and “In July to June 2020” is incorrect.

Corrected version

In July 2020, *V. minor* and *W. sinensis* leaf samples exhibiting severe virus-like disease symptoms of yellow halo, bright yellow mottle, or mosaic (Fig. 1) were collected from urban areas Mashhad, Razavi Khorasan province, in northeast Iran.

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