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.