CORRECTION

Correction To: Detection and diversity of maize yellow mosaic virus infecting maize in Nigeria

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In the original publication of the article, Figures 1 and 2 were interchanged. The correct Figs. 1 and 2 are given below:

The original article can be found online at <https://doi.org/10.1007/s13337-019-00555-0>.

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Fig. 1 Virus-like symptoms observed on maize plants in Kaduna and Katsina states, Nigeria. The observed symptoms included (a) wide yellow stripes that may sometimes be accompanied with (b, c) stunted

growth. Symptomatic plants tested positive by RT-PCR for maize yellow mosaic virus (MaYMV)

Fig. 2 Unrooted minimum evolution phylogenetic tree (K2 + G + 1) depicting evolutionary relationships among global isolates of maize yellow mosaic virus (MaYMV) based on analysis of ~ 1100 bp nucleotide sequences encompassing the partial P1–P2 fusion protein and coat protein genes of MaYMV. The analysis was performed with the MEGA7 program [15] with 1000 bootstrap replications. *MZ* maize, *SC* sugarcane, *IGRS* itch grass, *SG* Sorghum, *PN* Panicum