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Author Correction: The *Phytophthora cactorum* genome provides insights into the adaptation to host defense compounds and fungicides

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The information in this Article is incomplete. Since the publication of this Article the Whole Genome Shotgun project has been deposited at DDBJ/ENA/GenBank under the accession number WXXXK000000000; the assembled genome and annotations has also been deposited in the National GeneBank DataBase (Project ID: CNP0000756). Therefore, the Data Availability text,

“All raw genome sequence data have been deposited in the Short Read Archive (SRA) at NCBI under accession number SRR3386345 (PRJNA318145). Raw RNA-seq data have been deposited in the SRA under accession number SRP111895.”

should read:

“All raw genome sequence data have been deposited in the Short Read Archive (SRA) at NCBI under accession number SRR3386345 (PRJNA318145). Raw RNA-seq data have been deposited in the SRA under accession number SRP111895. This Whole Genome Shotgun project has been deposited at DDBJ/ENA/GenBank under the accession WXXXK000000000. The version described in this paper is version WXXXK01000000, the gene IDs have been updated according to Supplementary Data (file: Phyca.WXXXK01.p2g.xlsx). The assembled genome and its annotation also were deposited in the China National GeneBank DataBase (Project ID: CNP0000756).”

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