



## First report of *Golovinomyces sordidus* causing powdery mildew on plantain in Vietnam

L.T.T. Tam<sup>1\*</sup> and H.V. Cuong<sup>2</sup>

<sup>1</sup> Division of Plant Pathology, Plant Protection Research Institute (PPRI); Duc Thang, Bac Tu Liem, Hanoi, Vietnam; <sup>2</sup> Research Centre for Tropical Plant Pathology (RCTPP), Vietnam National University of Agriculture (VNUA), Trau Quy, Gia Lam, Hanoi, Vietnam

\*E-mail: [thanhtamle10\\_2012@yahoo.co.uk](mailto:thanhtamle10_2012@yahoo.co.uk)

**Received:** 27 Aug 2015. **Published:** 20 Dec 2015. **Keywords:** *Erysiphe sordida*, medicinal crops, *Plantago major*

*Plantago major* (plantain) is one of the most abundant and widely distributed medicinal crops in the world. Since 2013, typical symptoms of powdery mildew were observed in plantain growing wild in fields as well as cultivated for medicine production, the latter with a large area in Hanoi, Vietnam. Disease incidence was observed at around 80% and the disease severity almost 100% with almost all aerial parts of the plant (i.e. flowers, leaves and stem), covered by white fungal colonies (Fig. 1). As a result, yield at harvest was dramatically decreased. A representative diseased specimen was deposited in the PPRI Herbarium (Accession No. PPRI-PM050).

Conidiophores were composed of one to three cells, and measured (82.4-)112.2-147.1(-172.1) µm long. Footcells of conidiophores were cylindrical, straight, or curved at the base and 50-62.5 x 10-12.5 µm long (Fig. 2). Conidia, produced in chains, were variable in shape, oval to cylindrical, oval or doliform, 30-35 x 17.5-20 µm with a length/width ratio of 1.6 to 2.0. No chasmothecia were found. The morphological characteristics were consistent with descriptions of the genus *Golovinomyces* (Braun & Cook, 2012). To confirm the identity of the causal fungus to the species level, the complete ITS region of rDNA from diseased plant accession PPRI-PM050 was amplified with our universal primer pairs HF1 (5'-GGATCCTCGTAACAAGGTTTCCGTAG-3')/HR4 (5'-CTGCAGCTCCGCTTATTGATATGCTT-3') (Tam *et al.*, 2015) and directly sequenced. The resulting sequence (636 bp) was deposited in GenBank (Accession No. KM260734). A BLAST search using the present data revealed that the ITS sequence shares 99% identity with those of *Erysiphe sordida* and *Golovinomyces sordidus* (AF011309, AB077658 respectively) in which *E. sordida* and *G. sordidus* are synonyms (Saenz & Taylor, 1999; Matsuda & Takamatsu, 2003).

Pathogenicity was confirmed through inoculation by gently pressing

diseased leaves of plantain onto young leaves of ten potted three-week-old seedlings of *P. major*. Ten non-inoculated seedlings were used as controls. Plants were maintained in a greenhouse in PPRI at 26-28°C. Inoculated leaves developed symptoms after nine days, whereas the control plants remained symptomless. The fungus present on the inoculated leaves was morphologically identical to that observed on the original diseased leaves, with the same sequence following PCR as above, fulfilling Koch's postulates. *G. sordidus* is also known to infect several host plants in the Plantaginaceae and is widely distributed in North America, Europe, Central Asia, China, Japan, Korea and New Zealand (Braun & Cook, 2012). To our knowledge, this is the first report of *G. sordidus* infecting *P. major* in Vietnam. The disease seems to be an important threat to the production of pharmaceuticals from plantain in Vietnam.

### References

- Braun U, Cook RTA, 2012. Taxonomic Manual of the *Erysiphales* (Powdery Mildews), CBS Biodiversity Series No. 11. Utrecht, The Netherlands: CBS-KNAW Fungal Biodiversity Centre.
- Matsuda S, Takamatsu S, 2003. Evolution of host-parasite relationships of *Golovinomyces* (Ascomycete: Erysiphaceae) inferred from nuclear rDNA sequences. *Molecular Phylogenetics and Evolution* 27, 314-327. [http://dx.doi.org/10.1016/S1055-7903\(02\)00401-3](http://dx.doi.org/10.1016/S1055-7903(02)00401-3)
- Saenz GS, Taylor JW, 1999. Phylogeny of the Erysiphales (powdery mildews) inferred from internal transcribed spacer ribosomal DNA sequences. *Canadian Journal of Botany* 77, 150-168. <http://dx.doi.org/10.1139/b98-235>
- Tam LTT, Dung PN, Liem NV, 2015. First report of powdery mildew caused by *Erysiphe cruciferarum* on *Brassica juncea* in Vietnam. *Plant Disease* (in press). <http://dx.doi.org/10.1094/PDIS-06-15-0678-PDN>



Figure 1



Figure 2

**To cite this report:** Tam LTT, Cuong HV, 2015. First report of *Golovinomyces sordidus* causing powdery mildew on plantain in Vietnam. *New Disease Reports* 32, 33. <http://dx.doi.org/10.5197/j.2044-0588.2015.032.033>

©2015 The Authors

This report was published on-line at [www.ndrs.org.uk](http://www.ndrs.org.uk) where high quality versions of the figures can be found.