

1 First Report of *Rhizopus oryzae* Causing Potato Soft Rot in the  
2 Hebei Province of China

3

4 **W. G. Cui<sup>1,2</sup>, H. L. Zheng<sup>2</sup>, F. B. Zhang<sup>1,2</sup>, B. Swingle<sup>3</sup>, H. T. Zhu<sup>1</sup>, M. Gao<sup>1</sup>§**

5 <sup>1</sup>Key Laboratory of Microbial Resources Collection and Preservation, Ministry of  
6 Agriculture / Institute of Agricultural Resources and Regional Planning, Chinese  
7 Academy of Agricultural Sciences, Beijing 100081, PR China

8 <sup>2</sup>College of Agronomy, Inner Mongolia Agricultural University, Hohhot, Inner  
9 Mongolia Autonomous Region 010018, China

10 <sup>3</sup>Emerging Pests and Pathogens Research Unit, Robert W. Holley Center, United  
11 States Department of Agriculture-Agricultural Research Service, Ithaca, New York,  
12 USA, 14853 / Department of Plant Pathology and Plant-Microbe Biology, Cornell  
13 University, Ithaca, New York, USA, 14853

14

15 W. G. Cui and H. L. Zheng are co-first authors.

16 §Corresponding author: M. Gao, Email: [gaomiao@caas.cn](mailto:gaomiao@caas.cn)

17

18 **Funding:** The National Key Research and Development Program of China, Project  
19 No.: 2018YFD0201203.

20

21 *Rhizopus* soft rot occurs on the succulent tissues of vegetables, fruits and ornamental  
22 plants throughout the world. In September 2016, a disease outbreak in potato  
23 suspected as *Rhizopus* soft rot occurred in two fields in Guyuan County, Zhangjiakou  
24 City, Hebei Province of China. There were soft water-soaked lesions on the surface of  
25 diseased tubers, and the interiors were black and soft. Infected plants were sampled to  
26 isolate and determine the identity of the disease-causing organism. Symptomatic  
27 tubers were surface-sterilized with 1% NaClO for 1 min, rinsed three times with  
28 autoclaved distilled water. Small pieces of the inner decayed tissue were removed and  
29 cultured on potato dextrose agar (PDA) at 28°C for 2-3 days. Fifteen white colonies

30 grew rapidly on PDA, they became brownish-grey to blackish-grey mycelium. One  
31 colony was isolated (MLS-22-1-z15) for identification. Under the microscope,  
32 non-septate rhizoids, sporangia and sporangiospores were observed. Sporangiospores  
33 were unequal, numerous irregular, sub-globose or oval, and  $5.7 (3.5 \text{ to } 8.2) \times 4.7 (3.1$   
34  $\text{ to } 6.5) \mu\text{m}$  ( $n = 50$ ) in diameter. The sporangia were globose, black,  $120.2 (64.5 \text{ to}$   
35  $165) \mu\text{m}$  ( $n = 50$ ) in diameter. We analyzed the sequence of the internal transcribed  
36 spacer (ITS) region between genes coding for the large and small rRNA genes to  
37 determine the identity of the MLS-22-1-z15 isolate. The sequence of ITS was  
38 determined by PCR and Sanger sequencing using ITS4/ITS5 primers (Abe et al.  
39 2010). BLAST analysis of the MLS-22-1-z15 isolate ITS sequence (accession no.  
40 MH046841) revealed 100% sequence identity with *Rhizopus oryzae* (accession no.  
41 KU203330.1). Based on the morphological characteristics (Zhang et al. 2013; Suyala  
42 et al. 2008) and rDNA-ITS sequencing, the MLS-22-1-z15 isolate was identified as *R.*  
43 *oryzae*. To test whether we had isolated the disease-causing organism, we inoculated  
44 healthy tubers with isolate MLS-22-1-z15 by using a wound inoculation method  
45 (Choiseul et al. 2007) and observed disease development and progression. The  
46 disease-free potato tubers were rinsed three times with autoclaved distilled water, and  
47 surface sterilized with 75% ethanol. A 3 mm plug was removed from the surface  
48 sterilized tubers using a cork borer, and a 3-mm-diameter agar plug from the edge of a  
49 2-day-old colony was placed in the hole. An uninoculated agar plug was inserted into  
50 the wound as a negative control. These results were replicated in three independent  
51 trials, with each including three tubers, for both treatments. After incubation at 28°C  
52 for 3 days, all of the inoculated tubers had extensive decay and gave off a  
53 foul-smelling odor. The control tubers had no obvious symptoms. *R. oryzae* was  
54 re-isolated from the symptomatic potato tubers and confirmed as such based on  
55 colony and sporangia morphology and ITS sequence. To our knowledge, this is the  
56 first report of *R. oryzae* associated with soft rot on potatoes in China.

57

## 58 **References**

59 Abe, A., et al. 2010. Bioscience, Biotechnology, and Biochemistry. 74:1325.

- 60 Choiseul, J., et al. 2007. *Potato Research*. 49:241.
- 61 Suyala, S., et al. 2008. *Songklanakarinn Journal of Science and Technology*. 30:121.
- 62 Zhang, X., et al. 2014. *Journal of Phytopathology*. 161:745.