First Report of Leaf Spot Caused by *Colletotrichum kahawae* on Cultivated Rocket (*Eruca sativa*) in Italy.

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Eruca sativa [syn. E. vesicaria subsp. sativa (Miller) Thell.], common name cultivated rocket or 8 9 arugula, is an annual species belonging to the Brassicaceae that is grown widely for fresh consumption, mostly in mixed, ready-to-eat salads. During September and October 2014, in a 10 commercial plastic-house (0.3 ha) in Piedmont, northern Italy, where rocket had been grown for 11 12 7 years in the same soil under intensive cultivation, a leaf spot was observed on rocket plants (cv. Coltivata) from 7 to 35 days after sowing, corresponding to the cotyledon stage to harvest. The 13 incidence of plants affected ranged from 5 to 15% in the more humid area of the tunnel (75 to 14 90% relative humidity) at 20 to 28°C, with 5 to 75% of the affected leaf area showing white 15 spots. Symptoms first appeared as pale white or cream-colored, circular spots, each 5 to 50 mm 16 in diameter, on the tip or margins of the leaf. Each spot had a thin, dark brown or black border. 17 Leaf spots enlarged, with a round or irregular shape, to form extensive dead areas. Symptoms 18 differed from those produced by Alternaria japonica and Fusarium equiseti on this crop (Gilardi 19 et al. 2013). Isolations were carried out from symptomatic leaf sections (each 1 mm<sup>2</sup>) dipped in 20 1% sodium hypochlorite for 1 min, then rinsed in sterilized water, and placed onto PDA 21 amended with 25 mg/liter of streptomycin sulfate. A *Colletotrichum* sp. (Bailey and Jeger 1992) 22

was recovered consistently at a frequency of 70% from twenty leaf sections incubated at 20 to 1 23°C and a 12 h photoperiod/day. Hyaline, cylindrical, aseptate, and thin-walled conidia (9.3 to 2 16.5 x 3.4 to 6.3 um, average 14.4 x 4.7  $\mu$ m; n = 41 conidia) were produced abundantly in 3 acervuli (latter 79.0 to 91.5 µm; n =15) in gray mycelium. Genomic DNA was extracted from 10 4 mg of mycelium collected from PDA plates of one representative, single-conidium isolate. The 5 ITS1-5.8S-ITS2 region of ribosomal DNA (rDNA), and a fragment of the beta-tubulin 2 gene 6 (TUB2) between exons 2 and 6, were amplified using 200 ng genomic DNA as template (Faedda 7 et al. 2011). Purified amplicons (ExoSAP-IT) were sequenced in both directions. Sequence 8 analysis using BLASTN (Morgulis et al. 2008). The 489 and 600 bp fragments of ITS rDNA 9 (GenBank Accession No. KT259854) and TUB2 (KT259853), respectively, showed 100% 10 similarity with ITS and TUB2 sequences of Colletotrichum kahawae (JN715847.1 and 11 KC425710.1, respectively). Pathogenicity tests were performed on healthy, 25-day-old E. sativa 12 (cv. Coltivata) plants by spraying the leaves with a conidial and mycelial suspension (1 x  $10^5$ 13 conidia/ml) prepared from PDA plates of the same sequenced isolate of the pathogen. Control 14 plants were sprayed with sterilized water. Fifteen plants/treatment were used. The plants were 15 16 then covered with plastic bags for 5 days to maintain relative humidity close to saturation, and 17 kept in a growth chamber at 25°C under white fluorescent lamps (12 h photoperiod/day). Small, necrotic spots enlarged, and each formed a distinct black margin similar to those on the original 18 plants, 7 to 10 days after inoculation, while control plants remained asymptomatic. A 19 Colletotrichum sp. with morphological characteristics similar to C. kahawae was re-isolated 20 consistently from inoculated plants only, following the re-isolation protocol described above. No 21 fungi colonies were re-isolated from asymptomatic leaves of control plants. The pathogenicity 22 23 test was repeated with the same results. C. kahawae has been reported in Italy on olive (Olea

*europaea* L.) fruit (Schena et al. 2014). This is, to our knowledge, the first report of *C. kahawae* on *E. sativa* in Italy as well as worldwide. Due to the wide host range of *C. kahawae* and the
 economic value of cultivated rocket, this disease could be a threat in other rocket production
 areas of Italy and rocket, in turn, could be an inoculum reservoir for other susceptible crops.

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## 6 *References:*

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