

## Bursaphelenchus hofmanni Braasch, 199 growth substrate in hops nurseries

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To date, seven Bursaphelenchus species have been reported in surveys of the Czech Republic (Čermák et al., 2013). However, the occurrence of some species, such as Bursaphelenchus hofmanni Braasch, 1998, is limited to single detections in imported coniferous wood (unpubl. data in Braasch, 2001). During a survey to determine pathogenic agents on hops (Humulus lupulus L.) conducted by State Phytosanitary Administration (CZ) in hops fields and nurseries in 2012, B. hofmanni was found in a mixture of peat and soil in a hops seedling nursery in Zatec (Bohemia, Czech Republic). The occurrence of Bursaphelenchus species in non-woody plants or substrates has been previously reported for species of the fungivorus group, such as B. gonzalezi Loof, 1964, B. hunti Giblin & Kaya, 1983 and B. fungivorus Franklin & Hooper, 1962. Species belonging to the hofmanni group are often associated with wood products such as packaging material (Gu et al., 2006) and, to our knowledge, this is the first time that this species has been found associated with peat substrate and soil.

Nematodes were isolated from 60 g of peat substrate (3

and extracted to Specimens below fixed in hot 4%: according to De two males were Botryotinia fuck (MEA).

Nematode ide phological and ITS, 18S and 28 lows: single spe traction buffer (1% Triton X-10 1.5 ml Eppendo a micropestle an quently at 95°C DNA template for of rDNA were a DNA polymeras (NEB). GoTaq II a terminal 10 m



## Short communication

## 8 associated with peat in the Czech Republic

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ising the Baermann funnel technique, iging to Bursaphelenchus were killed and formalin and transferred to pure glycerin Grisse (1969). Two mature females and used to establish a culture maintained in eliana, growing on 5% malt extract agar

ntification was confirmed by both mormolecular analyses (sequencing of the S rDNA loci). DNA was extracted as folcimens were collected into 20 μl of ex-10 mM Tris-HCl, pH 8.8; 1 mM EDTA; 0 (v/v); 100 μg ml<sup>-1</sup> Proteinase K) in a rf tube. Each sample was ground using indincubated at 55°C for 1 h and subsefor 10 min. The mixture was used as a for PCR. The SSU, LSU and ITS regions amplified using a Phusion High Fidelity in the (having 3' → 5' exonuclease activity) NA polymerase (Promega) was added to in/72°C elongation step to create an A-

osequent 1A cioning. The primers used