



## First report of European mountain ash ringspot-associated virus in serviceberry (*Amelanchier* spp.) in Germany

S. von Bargaen\*, M. Tischendorf and C. Büttner

Humboldt-Universität zu Berlin, Faculty of Life Sciences, Division Phytomedicine, Lentzeallee 55/57, 14195 Berlin, Germany

\*E-mail: susanne.von.bargaen@agrar.hu-berlin.de

Received: 16 Apr 2018. Published: 11 May 2018. Keywords: urban green space, viral plant disease

Chlorotic ringspots and line pattern have been observed in different species of serviceberry (*Amelanchier* spp.), for instance in the Netherlands since 1957 (Cooper, 1979). Being tolerant against various abiotic stresses these shrubs are often used in urban green space and also have decorative flowers, berries and a red leaf colouration in the autumn.

Serviceberry shrubs cultivated in public or private urban areas in the cities of Oldenburg (Lower Saxony) and Berlin, Germany exhibiting chlorotic ringspots, mottle and line patterns on the leaves were sampled (Fig. 1, Table 1). The disease resembled characteristic symptoms caused by *European mountain ash ringspot-associated virus* (EMARaV) in serviceberry (Grimová *et al.*, 2015). To confirm the presence of the virus in these samples, total RNA was extracted and tested by RT-PCR using a genus-specific primer set to detect RNA1 (Elbeaino *et al.*, 2013) and EMARaV-specific primers targeting RNA2, RNA3 (Mielke & Mühlbach, 2008) and RNA4 (Roßbach *et al.*, 2015). All four RNAs of EMARaV were detectable in each of the four serviceberries exhibiting symptoms (Fig. 2) and this was confirmed by sequencing of the PCR products. Complete viral genome segments were amplified by RT-PCR from sample E55282 using primers targeting the conserved terminal regions of emaraviruses (Di Bello *et al.*, 2015). PCR products were cloned and sequenced confirming the complete RNA3 (1.6 kb, GenBank Accession No. LT992915) and RNA4 (1.3 kb, LT992916) of EMARaV. Sequence comparison with respective reference sequences from GenBank and generation of a neighbour-joining phylogenetic tree using complete sequences of the nucleocapsid protein encoded by RNA3 of emaraviruses (Fig. 3) confirmed the virus as EMARaV with a maximum of 99.6 % (RNA3) and 99.1 % (RNA4) identity at the protein level.

This is the first record of EMARaV affecting serviceberry and causing chlorotic ringspots, mottling and line patterns in Germany. The virus has previously been confirmed to be widespread in *Sorbus aucuparia* in Germany (Roßbach *et al.*, 2015). The disease identified in Germany shares characteristic symptoms reported from EMARaV-infected serviceberry in the Czech Republic (Grimová *et al.*, 2015). It has been demonstrated that the causal agent is graft transmissible within species of the *Rosaceae* family including serviceberry and rowan (Cooper, 1979; Grimová *et al.*, 2015).

How the diseased serviceberries acquired the virus in independent locations in Germany is unknown as modes of transmission other than grafting are not well documented for EMARaV.

### Acknowledgements

We thank Dr. T. Brand (LWK Niedersachsen) and Kira Köpke for collection of samples and the German Research Foundation for financial support (DFG-no: BU890/27-1).

### References

Cooper JI, 1979. *Virus Diseases of Trees and Shrubs*. 1st edition. Cambridge, UK: Institute of Terrestrial Ecology.

Di Bello PL, Ho T, Tzanetakis IE, 2015. The evolution of emaraviruses is becoming more complex: seven segments identified in the causal agent of Rose rosette disease. *Virus Research* **210**, 241-244. <http://dx.doi.org/10.1016/j.virusres.2015.08.009>

Elbeaino T, Whitfield A, Sharma M, Digiario M, 2013. Emaravirus-specific degenerate PCR primers allowed the identification of partial RNA-dependent RNA polymerase sequences of Maize red stripe virus and Pigeonpea sterility mosaic virus. *Journal of Virological Methods* **188**, 37-40. <http://dx.doi.org/10.1016/j.jviromet.2012.11.037>

Grimová L, Marek M, Konrady M, Ryšánek P, 2015. Newly identified host range of *European mountain ash ringspot-associated virus* (EMARaV) and its distribution in the Czech Republic. *Forest Pathology* **45**, 177-189. <http://dx.doi.org/10.1111/efp.12151>

Mielke N, Weber M, Kahn S, Mühlbach HP, 2008. Detection of *European mountain ash ringspot-associated virus* (EMARaV) in *Sorbus aucuparia* L. by a specific antiserum and reverse transcription-PCR. *Forest Pathology* **38**, 371-380. <http://dx.doi.org/10.1111/j.1439-0329.2008.00553.x>

Roßbach J, Dieckmann HL, Büttner T, Mühlbach H-P, von Bargaen S, Büttner C, 2015. Genetic variability and phylogeny of *European mountain ash ringspot-associated virus* RNA3 and RNA4. *Forests* **6**, 4072-4087. <http://dx.doi.org/10.3390/f6114072>



Figure 1  
Table 1: Overview of serviceberries (*Amelanchier* spp.) sampled from Germany

Sample ID	Symptoms	Collection date	Location, street	Tree	Area
E55282	chlorotic ringspots	September 2016	Klingenbergstraße	Oldenburg	public space
E55283	no symptoms	September 2016	Klingenbergstraße	Oldenburg	public space
E56730	chlorotic ringspots, line pattern	June 2017	Klingenbergstraße	Oldenburg	public space
E56731	chlorotic ringspots, mottle	June 2017	Brandenburger Straße	Oldenburg	public park
E57062	chlorotic ringspots	June 2017	Döhrenstieg	Berlin	private garden

Figure 4

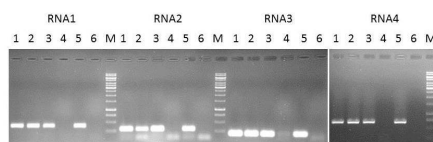


Figure 2

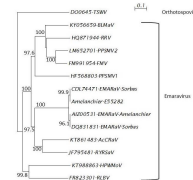


Figure 3

**To cite this report:** von Bargaen S, Tischendorf M, Büttner C, 2018. First report of *European mountain ash ringspot-associated virus* in serviceberry (*Amelanchier* spp.) in Germany. *New Disease Reports* **37**, 19. <http://dx.doi.org/10.5197/j.2044-0588.2018.037.019>  
©2018 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.