Localization of EMARaV proteins in planta by Agrobacterium-mediated transformation

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Inferences about the function of the p4 protein of EMARaV are expected by investigating the localization of the protein *in planta*.

European mountain ash ringspot-associated virus (EMARaV) infects European mountain ash (S. aucuparia L.) and causes chlorotic ringspots and mottling of leaves (Figure 1). The virus is suspected to influence the decline of branches and even the entire tree (Benthack et ctio al., 2005). EMARaV is composed of four ss(-)RNA genome segments and is assigned to the genus *Emaravirus* (Mühlbach & Mielke-Ehret, 2011). Each of the four viral RNAs is coding for one protein (p1-p4). Currently it is unknown, which function the RNA4 encoded p4 protein 0 adopts in the infection process. For many plant viruses the existence ntl of a gene silencing suppressor and a movement protein is essential. These functions could not be associated with the proteins encoded by





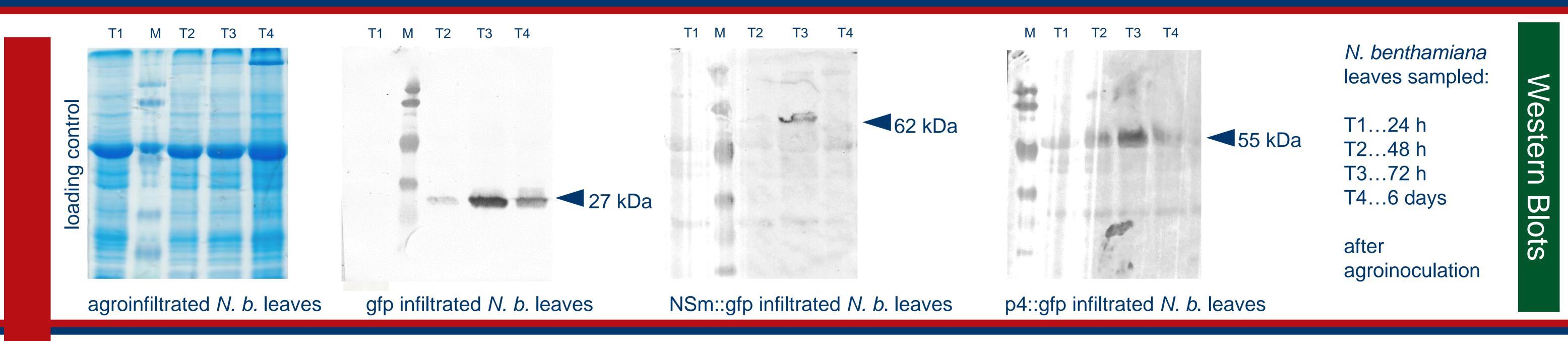
RNA1-RNA3. It can be assumed that EMARaV RNA4 encodes a gene silencing suppressor and/or a movement protein (MP).

For localization purposes the p4 protein of EMARaV were C-terminal (TSWV) was cloned as a reference for putative movement function of õ EMARaV p4. Agrobacteria were transformed with gfp constructs and Nicotiana benthamiana leaves were subsequently agroinfiltrated. \geq Expression of the gfp-fused construct were checked by western blot analysis by use of an anti-gfp antibody (Roche).

Figure 1: EMARaV infected rowan with symptoms of mottling and chlorotic ringspots on leaves.

For Co-localization a construct of the MP of *Cucumber mosaic virus* fused with gfp. Additionally, the MP of the Tomato spotted wilt virus fused with a red fluorescence protein (CMV_MP::RFP) was used. Plasmolysis treatments were conducted by incubation of small leaf sections (diameter of 0.5 cm) in 4% NaCl for 20 min according to Oomen et al., 2011. Confocal laser scanning microscopy was used for localization of viral proteins.

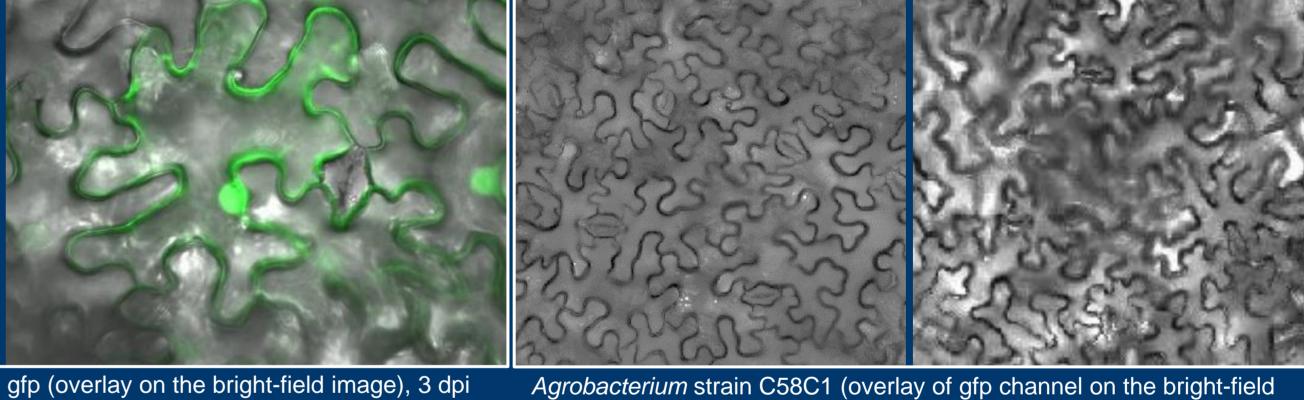




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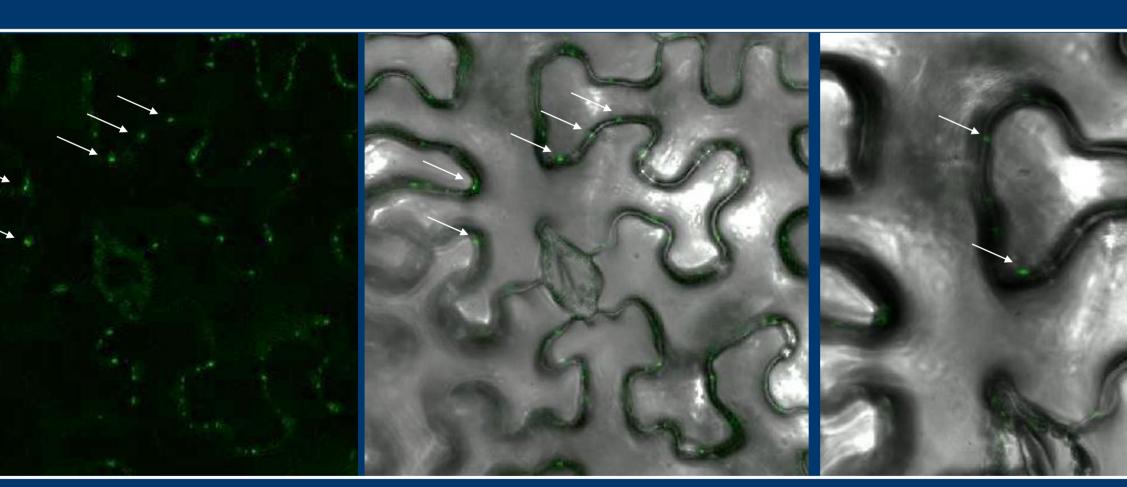
Known

MPs



Agrobacterium strain C58C1 (overlay of gfp channel on the bright-field image), 1 dpi (left) and 3 dpi (right)





TSWV_NSm::gfp signal, 2 dpi

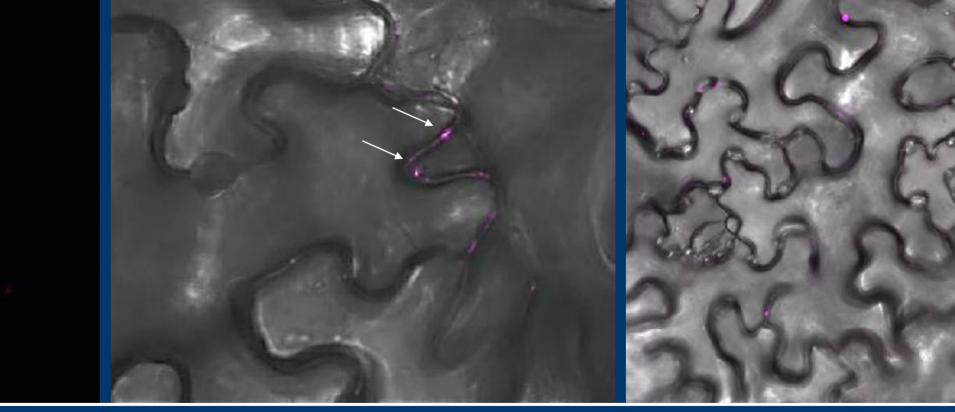
gfp signal, 1 dpi

TSWV_NSm::gfp (overlay on the bright-field image), 2 dpi

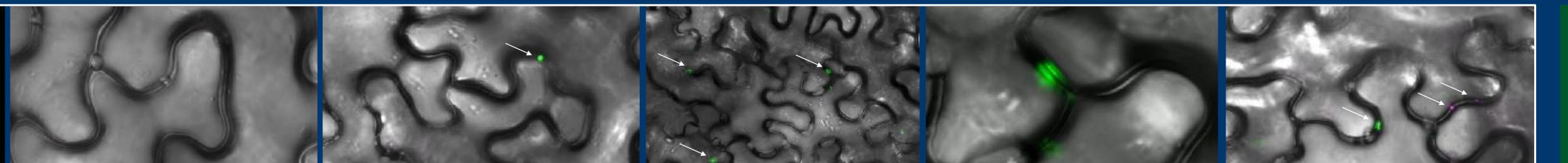
gfp (overlay on the bright-field image), 1 dpi gfp signal, 3 dpi

CMV_MP::rfp signal, 2 dpi





CMV_MP::rfp (overlay on the bright-field image), 2 dpi (left), 3 dpi (right)



EMA

EMARaV_p4::gfp signal, 2 dpi

EMARaV_p4::gfp (overlay on the bright-field image), 2 dpi (left), 3 dpi (right)

EMARaV_p4::gfp (overlay on the bright-field image), 3 dpi, plasmolysis treatment EMARaV_p4::gfp and CMV_MP::rfp coinfiltration (overlay of all channels), 2 dpi

Expression in N. b. of gfp fusion constructs of EMARaV p4 protein and TSWV NSm protein were confirmed with an anti-gfp antibody in western blot analysis > p4 protein was expressed 24 hours after agroinoculation > A significant expression of NSm was observed after three days Both proteins could be localized after two days by CLSM > TSWV NSm and CMV MP showed a regular distribution of punctuated spots \succ along the cell wall, which is typical for virus MPs References

Benthack W, Mielke N, Büttner C, Mühlbach H-P. 2005. Archives of Virology 150: 37-52. Mühlbach H P, Mielke-Ehret N. 2011. Emaravirus. Elsevier Academic Press, San Diego/USA: 767-770. Oomen R, Séveno-Carpentier E, Ricodeau N, Bournaud C, Conéjéro G, Paris N, Berthomieu P, Marquès L. 2011. New Phytologist 192:140-150.

> The p4 protein is located as punctuated spots at the cell wall, too > But it is not regular distributed along the whole cell wall Plasmolysis treatment did not affect localization of p4 at the cell wall

> Co-localization of EMARaV p4 or TSWV NSm with the MP of CMV was not successful

Co-infiltrated constructs were found solely in different cells.

Acknowledgement

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