COMMENTS

## Comments on "Determination of heterozygosity for avirulence/virulence loci through sexual hybridization of *Puccinia striiformis* f. sp. *tritici*"

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Over the past decades Puccinia striiformis f.sp. tritici (Pst) has developed into one of the most, if not the most important fungal pathogen in wheat production worldwide. In China, Pst has caused numerous epidemics with partially devastating yield losses<sup>[1]</sup>. The occurrence of the "warrior" race in Europe in 2011 also caused significant problems<sup>[2]</sup>. *Pst*, like other obligate biotrophs, is characterized by a high degree of genetic variability, especially with respect avirulence/virulence development on specific host varieties. This variability may be caused by mutations, somatic recombination, or recombination during the sexual stage of the fungus. The discovery of the hitherto unknown alternate host for Pst in 2010<sup>[3]</sup> put a new focus on the role of sexual recombination in this variability<sup>[4,5]</sup>. With now 35 barberry species identified as potential alternate host for *Pst* in one of China's bread baskets (Gansu, Sichuan, Shaanxi, Yunnan and Tibet)<sup>[6,7]</sup>, elucidating the role of the sexual cycle of *Pst* has become even more important. The work entitled "Determination of heterozygosity for avirulence/virulence loci through sexual hybridization of Puccinia striiformis f. sp. tritici" by Yuan TIAN, Gangming ZHAN, Xia LU, Jie ZHAO, Lili HUANG, and Zhensheng KANG, in this issue (DOI: 10.15302/J-FASE-2016114), is an important work in the light of giving breeders some indication as to which wheat varieties to use for future crosses. From the 25 wheat varieties tested, 17 turned out to carry resistance genes for which the corresponding Pst avirulence/virulence genes turned out to be heterozygous. Such lines should be excluded from future breeding programs as chances are high that resistances generated in the new cultivars will be rapidly overcome. As such this paper constitutes a major advance in understanding the intricate interaction of Pst and its host wheat, and at the same time provides practical cues for future breeding programs.

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acquisition, metabolism and suppression of plant resistance mechanisms in rust fungi. He habilitated in 2008 at the University of Konstanz with the monograph "The Role of Haustoria in the Biotrophic Interaction of the Rust Fungus *Uromyces fabae* and Its Host Plant *Vicia faba*". He obtained the *venia legendi* for Phytopathology and Microbiology. Since 2010 he is Chair for Phytopathology at the Institute of Phytomedicine, Faculty of Agricultural Sciences of the University of Hohenheim, Germany. From 2010 to 2014 he was CEO of the Institute of Phytomedicine, and since 2014 he is Deputy-CEO of this institute. In 2015 he was elected Dean of the Faculty of Agricultural Sciences of the University of Hohenheim. Ralf Voegele is member of the Vereinigung für Allgemeine und Angewandte Mikrobiologie (VAAM), the American Society for Microbiology (ASM), the International Society for Molecular Plant-Microbe Interactions (IS-MPMI), the American Phytopathological Society (APS), and the Deutsche Phytomedizinische Gesellschaft (DPG). Since 2014 he is director of the working group "Host-Parasite Interactions" of the DPG. Over the past 25 years Ralf Voegele published more than 50 peer reviewed scientific articles and has numerous collaborators in the US, Brazil, Europe, Africa, Australia, New Zealand, and China.