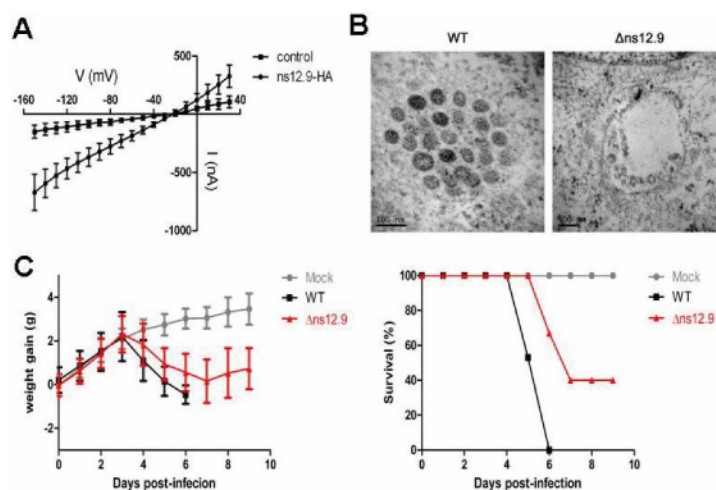


## The viroporin function of ns12.9 in HCoV-OC43 morphogenesis and pathogenesis

With the support by the National Science and Technology Major Project, the research team led by Prof. Sun Bing (孙兵) from the Institut Pasteur of Shanghai, Chinese Academy of Sciences, and Prof. Xiong Si-Dong (熊思东) from Soochow University, found that the ns12.9 accessory protein of human coronavirus OC43 (HCoV-OC43) is a viroporin involved in virion morphogenesis and pathogenesis. This wonderful work was recently published in *Journal of Virology* (2015, 89: 11383–11395), and selected as a Spotlight Feature article.

HCoV-OC43 was first isolated in the 1960s and is a major agent of the common cold. The functions of HCoV-OC43 structural proteins have been well studied, but few studies have focused on its accessory proteins. In the present study, we identified that the ns12.9 accessory protein is a newly recognized viroporin. Then, we engineered a recombinant mutant virus lacking the ns12.9 protein (HCoV-OC43- $\Delta$ ns12.9) to characterize the contributions of ns12.9 during HCoV-OC43 infection, and found that HCoV-OC43- $\Delta$ ns12.9 presents a growth defect *in vitro* and *in vivo*. Systematic dissection of single-cycle replication revealed the important functions of ns12.9 viroporin in virion morphogenesis. Furthermore, the mice infected with HCoV-OC43-ns12.9 exhibited reduced inflammation and virulence accompanied by a lower titer in the brain than that of wild-type-infected mice, suggesting ns12.9 viroporin influences virus pathogenesis. These findings contribute to the knowledge on the functional relevance of viroporins to viral infectivity, and provide a potential target for developing antiviral drugs.



**Figure** The HCoV-OC43 ns12.9 protein is a viroporin involved in virion morphogenesis and pathogenesis. (A) The *I/V* curve of voltage dependencies of steady-state currents in control oocytes (filled squares) and ns12.9-expressing oocytes (filled circles). (B) Transmission electron micrographs of virions in the infected RD cells. (C) The weight variations and survival curves after wild type and mutant virus infection.