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COVER IMAGE



Cover photograph: Bats are a reservoir of zoonotic viruses, including filoviruses and henipaviruses; thus, viral coinfections are likely. Hendra and Nipah viruses are deadly henipaviruses that share a high degree of sequence similarities in their attachment (G) and fusion (F) glycoproteins. Heterologous combinations of Nipah and Hendra viral G and F proteins were analyzed by Bradel-Trethewey et al. and found to yield distinct hyper- or hypofusogenic phenotypes. Further, the authors mapped the regions of the glycoproteins responsible for such phenotypes, taking into consideration the cell surface expression levels of these proteins via a homologous tagging approach. (See related article at e00577-19.) (Acrylic on canvas. Courtesy of Armando Pacheco; reprinted with permission.)

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AUTHOR CORRECTION

Correction for Sun et al., “Risk Assessment of Fifth-Wave H7N9 Influenza A Viruses in Mammalian Models” **e00476-19**

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