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COVER IMAGE



Cover photograph: Influenza viruses replicate within and transmit between a wide range of avian and mammalian host species. Here, some of these species have gathered to commiserate. A common feature of influenza virus infection is the production of defective interfering particles (DIPs) that harbor large deletions in one or more genome segments. The causes and consequences of DIP formation remain poorly understood. Alnaji et al. developed a next-generation sequencing pipeline to facilitate high-resolution analysis of DIPs present in influenza A and B virus populations. (See related article at e00354-19.) (Courtesy of Allegra Raff; reprinted with permission.)

SPOTLIGHT

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