Porcine Parainfluenza Virus Type 1 (PPIV-1) in U.S. Swine: Summary of Veterinary Diagnostic Laboratory Data

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Porcine Parainfluenza Virus Type 1 (PPIV-1) in U.S. Swine: Summary of Veterinary Diagnostic Laboratory Data

A.S. Leaflet R3196

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Summary and Implications
Porcine Parainfluenza-1 (PPIV-1) is a Paramyxovirus in the genus Respirovirus. To evaluate the frequency of detection of PPIV-1 on farms in the United States, the Iowa State University Veterinary Diagnostic Laboratory (ISU VDL) conducted an analysis of clinical samples submitted from Spring to Fall 2016. Thirty-four percent of diagnostic samples tested both prospectively (70/204) and retrospectively (472/1385) were positive by polymerase chain reaction (PCR), which suggests PPIV-1 is widespread.

Introduction
PPIV-1 was first detected in swine at the University of Hong Kong in 2013, prompting diagnostic laboratories in the United States (US) to develop assays capable of detecting PPIV-1 in clinical samples. However, it is unclear whether PPIV-1 contributes significantly to the porcine respiratory disease complex (PRDC) since PPIV-1 has been detected from both symptomatic and asymptomatic herds. Other viruses comprising the PRDC include porcine reproductive and respiratory syndrome virus (PRRSV), porcine circovirus type 2 (PCV2) and influenza A virus (IAV). The pathogenesis, transmission, and epidemiology of PPIV-1 have yet to be determined.

Materials and Methods
Diagnostic samples submitted to the ISU VDL were tested prospectively and retrospectively by a PPIV-1-specific real time reverse transcriptase PCR (RT-rPCR). In total, 1589 samples collected from 27 US states were tested; these included but were not limited to oral fluid, tissue homogenates, environmental samples, serum, nasal swabs, oropharyngeal swabs, and bronchoalveolar lavage fluid.

Results and Discussion
A total of 34% of the clinical samples tested were RT-rPCR positive for PPIV-1 (Figure 1). These included oral fluid (75.4%), lung (7.2%), nasal swab (13.1%), and environmental samples (2.3%). These data suggest oral fluids are most likely to contain PPIV-1, followed distantly by nasal swabs.

Iowa, North Carolina, and Illinois had the highest percent of PPIV-1 RT-rPCR positive samples by state, regardless of clinical history (Figure 2). This is correlated to the swine population distribution in the US and the relative proportion of samples submitted to the ISU VDL.
appropriate for PPIV-1 detection when associated with clinical disease (Figure 3).

PPIV-1 has also been detected in coinfections with IAV and PRRSV. Although it is still unknown, these viral coinfections could possibly increase severity of respiratory disease. In summary, these data suggest a high prevalence of PPIV-1 in US swine herds that may not be associated with clinical disease. Herds can be asymptomatic but still carry PPIV-1. More work is needed in developing a comprehensive diagnostic array of assays to assist in the detection of PPIV-1 as well as determining its contribution to the respiratory disease complex.

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