

# USDA National Surveillance Program for Influenza A Virus in Swine

Technical Review, Evaluation, and Recommendations  
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## Executive summary

The National Surveillance Program for influenza A virus in the U.S swine population is risk-based and monitors the genetic evolution of endemic influenza A virus in swine to better understand endemic and emerging influenza virus ecology. The program is precedent setting because it is a USDA-APHIS-VS first for an endemic disease and involves a non-regulatory approach to a potentially regulated disease and has swine industry support and engagement.

In my opinion, the overall program has been successful because of:

- New knowledge created through the phylogenetic studies of co-circulating influenza viruses in swine, as part of cooperative work with the Agricultural Research Services
- Trust and goodwill which has developed between animal and public health partners
- Political value of the program and partnerships created

To inform discussion within USDA-APHIS, I recommend changes to components of the surveillance program, with the longer-term goal of maximizing efficiency and cost-effectiveness with regards to sampling design, data collection, data management and analysis, interpretation of results, and dissemination of information to stakeholders. My 26 recommendations are summarized on pages 21-23 of this report and are categorized as to whether they apply to the existing program or to a revised program that addresses proposed funding cuts.

My highest-ranked priorities for action are to:

- **Identify failure/inefficiency points** throughout the existing program and strategies that might be feasible to implement in order to improve program efficiency.
- **Undertake a cost-effective analysis of the testing algorithm** and this task should include deciding on the most appropriate number of samples for full-genome sequencing to achieve program objectives
- **Develop an impacts statement** for internal and external stakeholders that summarizes and promotes the animal and public health benefits of the current program; this should include data on how influenza isolates have been used by biologics companies
- **Develop a strategy for increasing accessions from the traceable stream**, including use of incentives and perhaps shifting responsibility and costs to the swine industry; this may include **having the swine industry provide representative regional samples to ensure adequate geographical coverage of the U.S. herd**
- Agree as to **what epidemiological analyses are realistic with the currently-available**

**data or after modification** of the program, given the reluctance of the swine industry to allow state-level analyses

- **Provide more explicit description of goals, objectives, and outcomes/deliverables** for a revised program including a reassessment of the importance of geographical coverage and representativeness as part of the programmatic goals
- **Define key attributes and associated metrics** for a revised program
- **Develop case-definitions that include genomic data** at the sample, pig, and herd levels for purposes of epidemiological analysis
- **Develop a comprehensive data management system** to allow more streamlined integration of data from multiple sources, including laboratory and field data

## Recommendations

	Number	For EXISTING Program	For REVISED Program
<b>Priorities and objectives</b>	1		<ul style="list-style-type: none"> <li>• Deliverables for each objective</li> <li>• Listing current priorities in reports</li> <li>• Traceability of changes</li> </ul>
	2		<ul style="list-style-type: none"> <li>• Swine Health Information Center (SHIC) as a possible stakeholder</li> </ul>
<b>Short- and long-term outcomes</b>	3		<ul style="list-style-type: none"> <li>• Update or change outcomes and define practical deliverables</li> </ul>
	4		<ul style="list-style-type: none"> <li>• Identify knowledge gaps in IAV-S ecology &amp; epidemiology that can be realistically addressed</li> </ul>
	5	<ul style="list-style-type: none"> <li>• Spatiotemporal analysis of variant IAV-S at state or regional levels</li> </ul>	
	6	<ul style="list-style-type: none"> <li>• Answer questions related to geographical objectives</li> <li>• Assess impact of data pooling across states into regions</li> <li>• Sample-size questions</li> </ul>	
	7		<ul style="list-style-type: none"> <li>• Discussions with swine industry regarding data management &amp; analysis for IAV-S epidemiology</li> </ul>
<b>Target and source populations</b>	8	<ul style="list-style-type: none"> <li>• Spatial coverage (preferably state-level)</li> <li>• Retrospective evaluation (at ARS) of variant IAV-S</li> </ul>	
	9	<ul style="list-style-type: none"> <li>• Describe criteria for selection of viruses for full genome sequencing</li> </ul>	
<b>Case definition</b>	10		<ul style="list-style-type: none"> <li>• Include genomic data</li> <li>• Review case definitions for clarity, accuracy, and acceptability</li> </ul>

<b>Data management and analysis</b>	11		<ul style="list-style-type: none"> <li>• Develop a comprehensive data management system for integrating field and laboratory data</li> </ul>
	12		<ul style="list-style-type: none"> <li>• Standardize data submissions from farms and electronic transmission of data by all parties</li> </ul>
	13	<ul style="list-style-type: none"> <li>• Evaluate completeness and timeliness of NAHLN's data submissions from 2014 on and investigate worst 20% for impacts</li> </ul>	
	14	<ul style="list-style-type: none"> <li>• Update swine industry representatives about limited epidemiological data unless more traceable herds and possible follow-up can be included in program</li> </ul>	<ul style="list-style-type: none"> <li>• Discuss alternative models; with the swine industry leading epidemiologic studies and USDA providing a supporting role</li> </ul>
<b>Result interpretation</b>	15	<ul style="list-style-type: none"> <li>• Include pattern-interpretation disclaimer for sequence data in annual reports</li> </ul>	
<b>Testing protocol/cost effectiveness</b>	16	<ul style="list-style-type: none"> <li>• Cost-analysis of full sampling and testing algorithm</li> </ul>	
	17	<ul style="list-style-type: none"> <li>• Calculate number of isolates to be sequenced to meet the desired objectives of the surveillance program including phylogenetic analyses</li> </ul>	
<b>Sources of system inefficiency &amp; failures</b>	18	<ul style="list-style-type: none"> <li>• Identify and rank failures and inefficiencies for each step in entire surveillance system</li> </ul>	
	19	<ul style="list-style-type: none"> <li>• Set metrics for times within each step, as appropriate</li> <li>• Define unacceptable failure percentage for each step</li> </ul>	
	20	<ul style="list-style-type: none"> <li>• Consider cost-effective and feasible mitigations for identified failure/inefficiency points</li> </ul>	

	21	<ul style="list-style-type: none"> <li>Impacts statement summarizing benefits to stakeholders and focusing on public health and animal health goals</li> </ul>	<ul style="list-style-type: none"> <li>Future discussions with stakeholders and modifications of the existing system (to ensure more focus on deliverables and outcomes) informed by the impacts statement</li> </ul>
<b>Success and impact metrics</b>	22	<ul style="list-style-type: none"> <li>Survey biologics companies to determine use of IAV-S from NVSL for vaccine development</li> </ul>	
<b>Dissemination to stakeholders</b>	23	<ul style="list-style-type: none"> <li>Describe communication process to the swine industry for a unique strain</li> <li>Describe roles of responsible parties in the communication process</li> </ul>	
<b>Cost-sharing</b>	24	<ul style="list-style-type: none"> <li>Initiate discussions with swine industry representatives about sharing testing costs</li> </ul>	
<b>Surveillance attributes</b>	25	<ul style="list-style-type: none"> <li>Identify and rank the most important attributes to be assessed in the surveillance program</li> </ul>	<ul style="list-style-type: none"> <li>Identify potential metrics for a modified program based on these assessed attributes</li> </ul>
<b>Modification process</b>	26		<ul style="list-style-type: none"> <li>Develop a Gantt chart showing the sequential and simultaneous steps in the modification process and the timeline for each</li> <li>Develop a written plan for regular internal evaluation of surveillance system attributes</li> </ul>