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pandemic potential than some other influenza strains due to its ability to infect both pigs and humans. The 2009 H1N1 pandemic served as a real-world example of how swine influenza can rapidly spread worldwide, causing significant illness and mortality. Since the 2009 pandemic, the H1N1 strain of swine influenza has continued to circulate as a seasonal influenza virus alongside other influenza strains. This theory highlights the importance of ongoing surveillance and vaccination efforts to monitor and control the spread of swine influenza. Research in the field of swine influenza includes theories and studies on improving vaccination strategies.

This includes developing more effective vaccines, understanding the long-term immunity provided by previous exposure or vaccination, and assessing the need for regular booster shots. The "One Health" approach is a theory that emphasizes the interconnectedness of human, animal, and environmental health. This theory underscores the importance of studying swine influenza within the broader context of the ecosystem, considering factors such as agricultural practices, human-animal interactions, and environmental changes that may influence the emergence and spread of the virus. Ongoing research and surveillance aim to identify potential new strains or variants of swine influenza with pandemic potential. Scientists are particularly concerned about the possibility of reassortment events involving multiple influenza strains, which could lead to the emergence of highly contagious and virulent viruses. In conclusion, swine influenza research involves various theories and scientific investigations to better understand its behavior, evolution, and impact on public health. This knowledge is crucial for developing effective prevention and control strategies to mitigate the potential threat of swine influenza outbreaks and pandemics in the future. Swine influenza is believed to have originated in pigs and can be transmitted from pigs to humans. However, it can also spread efficiently from human to human through respiratory droplets [5-7].

The symptoms of swine influenza are similar to those of seasonal influenza and include fever, cough, sore throat, body aches, and fatigue. Gastrointestinal symptoms like nausea and diarrhea can also occur. Vaccination is a crucial preventive measure against swine influenza, with the H1N1 strain typically included in seasonal influenza vaccines. Good hygiene practices, such as handwashing and respiratory etiquette, are essential for reducing transmission. Antiviral medications like oseltamivir (Tamiflu) can help reduce the severity and duration of symptoms if administered early in the course of the illness. Swine influenza continues to circulate as a seasonal influenza virus. Surveillance and monitoring efforts remain in place to track the virus and adapt vaccines accordingly. The 2009 swine influenza pandemic highlighted the importance of global pandemic preparedness and cooperation in responding to infectious diseases. Swine influenza research includes theories related to zoonotic origins, genetic reassortment, antigenic shift and drift, and potential pandemic threats. The One Health approach emphasizes the interconnectedness of human, animal, and environmental health in understanding and managing the virus. In summary, while swine influenza is no longer the pandemic threat it once was, it continues to be a public health concern. Vaccination, public health measures, ongoing research, and international collaboration are essential components in managing and preventing the spread of swine influenza. The lessons learned from the 2009 pandemic have contributed to improved preparedness for future infectious disease outbreaks. The global distribution and ongoing evolution of type A swine influenza virus (IAV-S) continue

to pose significant challenges against developing broadly protective vaccines to control swine influenza. This study focuses on the hemagglutinin (HA) consensus-based approach towards developing a more broadly protective swine influenza vaccine against various H3 strains circulating in domestic pig populations. By computationally analyzing >1000 swine H3 full-length HA sequences, we generated a consensus H3 and expressed it in the context of influenza A WSN/33 reverse genetics system [8-10].

## Conclusion

The derived recombinant chimeric swine influenza virus with the consensus H3 was inactivated and further evaluated as a potential universal vaccine in pigs. The consensus H3 vaccine elicited broadly active hemagglutination inhibition (HI) antibodies against divergent swine H3N2 influenza viruses including human H3N2 variant of concern, and strains belong to genetic clusters IV, IV-A, IV-B, IV-C, IV-D and IV-F. Importantly, vaccinated pigs were completely protected against challenge with a clinical swine H3N2 isolate in that neither viral shedding nor replication in lungs of vaccinated pigs were observed. These findings warrant further study of the consensus H3 vaccine platform for broad protection against diverse swine influenza viruses.

## Acknowledgment

None

## Conflict of Interest

None

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