Development of Novel Chimeric Vaccine and Delivery System for Classical Swine Fever Virus Pengcheng Shang¹, Adriana Avila², Rui Guo¹, Susan K. Whitaker³, Yanhua Li¹, John Tomich³, Raymond R. R. Rowland¹, Ying Fang¹

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Introduction

Historically, the greatest threats to the US swine production are from foreign animal diseases, such as CSF. CSF is caused by CSFV, which is highly contagious to pigs and wild boars. CSFV belongs to the family Flaviviridae, genus Pestivirus. The genome of pestiviruses is a positive-sense, single stranded RNA. It comprises a single open reading frame (ORF) that encodes a polyprotein of about 4000 amino acids, which is processed by viral and cellular proteases to generate four structural proteins (C, Erns, E1 and E2) and nine nonstructural proteins. Infections caused by CSFV can have a deleterious effect on swine production, causing excessive morbidity and mortality. Modified live virus (MLV) vaccines for CSF, such as the CSFV C-strain, are effective in controlling infection but are not routinely used, primarily due to the lack of diagnostic tools that can differentiate infected from vaccinated animals (DIVA), which hampers disease control measures relying on serology. Furthermore, MLV vaccines may pose concerns on biosecurity issues, since virulent revertant virus can be easily obtained (and modified) from the vaccinated animals in the field. There are restrictions on the international trade in pig products from countries using CSF MLV vaccines. To circumvent these problems, subunit vaccines have been developed. Previous studies demonstrated that the CSFV E2 protein can induce specific neutralizing antibodies. Subunit vaccines based on the E2 glycoprotein have been tested extensively for their efficacy, and the companion DIVA test has been developed based on ELISAs that detects E2 and another viral glycoprotein, the Erns. In comparison to MLV vaccines, the main limitation of these subunit vaccines is their higher cost for production of antigens; on the other hand, immunization with E2 protein alone is less efficacious and the early stimulation of innate immune responses is needed for initial protection of immunized animals. To overcome these limitations, this proposed study will use a novel approach for CSF vaccine development, in which a modified live PRRSV will be used as a vector to express CSFV E2 protein. To eliminate the need for a cold chain, DNA vaccine approach was used, in which nanoparticles composed of branched amphiphilic peptide capsules (BAPCs,) were employed as the DNA delivery agent. The candidate vaccine and the DNA delivery system were evaluated in *in vitro* expression system and a nursery pig model. This study developed a candidate viral vectored chimeric vaccine and established DNA vaccine delivery system, which could be applied to other emerging and transboundary swine pathogens in the future.



nm. (B) Cluster of BAPCs interacting with DNA. Scale bar = 100 nm. (C) Schematic representation of potential BAPC-DNA interactions.



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