Check for updates

OPEN ACCESS

EDITED BY Wentao Li, Huazhong Agricultural University, China

REVIEWED BY Gaurav Kumar Sharma, Indian Veterinary Research Institute (IVRI), India Baochao Fan, Jiangsu Academy of Agricultural Sciences (JAAS), China

*CORRESPONDENCE Jianchao Wei jianchaowei@shvri.ac.cn Zhiyong Ma zhiyongma@shvri.ac.cn

SPECIALTY SECTION

This article was submitted to Veterinary Experimental and Diagnostic Pathology, a section of the journal Frontiers in Veterinary Science

RECEIVED 17 June 2022 ACCEPTED 05 September 2022 PUBLISHED 23 September 2022

CITATION

Yang Y, Xia Q, Sun Q, Zhang Y, Li Y, Ma X, Guan Z, Zhang J, Li Z, Liu K, Li B, Shao D, Qiu Y, Ma Z and Wei J (2022) Detection of African swine fever virus antibodies in serum using a pB602L protein-based indirect ELISA. *Front. Vet. Sci.* 9:971841. doi: 10.3389/fvets.2022.971841

COPYRIGHT

© 2022 Yang, Xia, Sun, Zhang, Li, Ma, Guan, Zhang, Li, Liu, Li, Shao, Qiu, Ma and Wei. This is an open-access article distributed under the terms of the Creative Commons Attribution License

(CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

Detection of African swine fever virus antibodies in serum using a pB602L protein-based indirect ELISA

Yang Yang¹, Qiqi Xia¹, Qin Sun^{1,2}, Yan Zhang^{1,2}, Yuhao Li¹, Xiaochun Ma¹, Zhixin Guan¹, Junjie Zhang¹, Zongjie Li¹, Ke Liu¹, Beibei Li¹, Donghua Shao¹, Yafeng Qiu¹, Zhiyong Ma^{1*} and Jianchao Wei^{1*}

¹Shanghai Veterinary Research Institute, Chinese Academy of Agricultural Sciences, Shanghai, China, ²College of Animal Science, Yangtze University, Jingzhou, China

African Swine Fever (ASF) is an acute, highly contagious and deadly infectious disease that has a huge impact on the swine industry. It is caused by the African swine fever virus (ASFV). The most acute forms of ASF in domestic pigs have mortality rates of up to 100%. The lack of a commercial vaccine and effective therapeutic drugs has brought great challenges to the prevention and control of ASF. Current, the African swine fever virus requires a huge amount of detection, so there is a need for more sensitive and accurate detection technology. The protein pB602L, as a late non-structural protein, has a high corresponding antibody titer and strong antigenicity in infected swine. In this research, the B602L gene was constructed into the pColdI prokaryotic expression vector, and prokaryotic expression of the soluble pB602L protein was induced by IPTG. Western blot analysis demonstrated that the protein had strong immunogenicity. We established an indirect ELISA method for the detection of anti-ASFV using purified recombinant pB602L protein as antigen. The detection method showed excellent specificity without cross-reactions with antibodies against PRRSV, CSFV, JEV, and GETV. The method could detect anti-ASFV in serum samples that were diluted up to 6,400 times, showing high sensitivity. The coefficients of variation of the intra-assay and inter-assay were both <10%. The assays had excellent specificity, sensitivity, and repeatability. In summary, we developed an accurate, rapid, and economical method for the detection of anti-ASFV in pig serum samples with great potential for ASF monitoring and epidemic control.

KEYWORDS

African swine fever virus, indirect ELISA, B602L, prokaryotic expression system, antibodies in serum

Introduction

African swine fever (ASF) is a highly contagious hemorrhagic and transboundary viral disease of swine for which currently no preventive vaccine is available (1). It is a devastating disease with mortality rates close to 100% in young herds, causing severe socioeconomic losses. Since it was first reported in Kenya in 1921 (2), ASF has become endemic in many African countries (3). Recently, ASF emerged in Asian countries, including China (4), where it spread rapidly to many parts of the country (5). At present, ASF continues to place a huge burden on the pig industry in the affected areas (6).

Current approaches to control and prevent ASF include early diagnosis, rapid elimination of infectious sources, and interruption of transmission routes. Therefore, the development of rapid, sensitive, and field-deployable African swine fever virus (ASFV) assays are important for disease surveillance and control (7, 8). Currently, there are several early diagnostic techniques available to detect ASFV, including PCR, loopmediated isothermal amplification, fluorescence quantitative PCR, colloidal gold rapid strip, hemadsorption, enzymelinked immunosorbent assay (ELISA), and other serological tests (9). Serological testing is been widely used in ASF eradication and control programs; since no vaccine is available, the presence of ASF antibodies implies previous infection. Therefore, serology is suitable for detecting potential viruscarrying animals in epidemic situations. ELISA is the most commonly used serological test, so it can be widely used for screening purposes (10).

ASFV is the sole member of the genus Asfarvirus within the family Asfarviridae and is a unique and genetically complex virus (11). ASFV is a large, double-stranded, linear DNA virus with a complex structure. The genome size is about 170-190 kb. ASFV encodes more than 150 proteins, of which more than 50 are structural proteins (12). ASFV can be divided into 23 different genotypes according to the ASFV B646L gene encoding the p72 major capsid protein (13). Morphologically, ASFV has a icosahedral shape and a complex structural architecture composed of several concentric layers (14, 15). ASFV infects mononuclear macrophages mainly through megalocytosis and clathrin-mediated endocytosis (1). pB602L is a nonstructural protein that functions as a molecular chaperone of the major structural protein p72, forming aberrant "zipper-like" structures instead of icosahedral virus particles in the absence of pB602L (16). B602L contains a central variable region (CVR) that allows frequently subgenotyping of ASFV isolates based on this region (17). The central variable region (CVR) within the B602L gene of the African swine fever virus (ASFV) is highly polymorphic within the ASFV genotypes defined by sequencing of the Cterminal end of the p72 locus (17, 18). Previous studies have shown that pB602L is strongly antigenic and can be used to develop diagnostic tools for ASFV. Previously, a pB602L-based ELISA assay was employed to detect serum antibodies against ASFV, and the test results were mostly consistent with those obtained using the gold standard, western blot assays (16). B Gutiérrez-Castañeda et al. showed that pB602L is recognized by hyperimmune antisera from domestic pigs and bushpigs at late time points after infection, suggesting it may be useful diagnostically to distinguish animals persistently infected with virus (17). pB602L may be a suitable target for the development of diagnostic tools to assess the humoral immune responses to these vaccines, as antibodies against pB602L are only produced after this protein is expressed in host cells (1, 17).

In this study, we successfully expressed and purified the ASFV pB602L protein and established an indirect ELISA method for the detection of anti-ASFV using purified pB602L as a diagnostic antigen, laying the foundation for the development of early diagnosis kits for ASFV and functional research of the pB602L protein.

Materials and methods

Serum samples

ASFV (SY-18) infection inactivated sera (n = 30) were obtained from the China Animal Health & Epidemiology Center (Qingdao, China). All positive sera were tested by gold standard. And all positive sera were tested by the Svanova ASFV p30 antibody detection kit and the Ingenasa ASFV p72 ELISA kit.

Positive sera against pathogenic Japanese encephalitis virus (JEV), classical swine fever virus (CSFV), Getah virus (GETV), Porcine circovirus 2 (PCV2), Porcine parvovirus (PPV)-, and Pseudorabies virus (PRV) were prepared in our laboratory.

All 120 negative serum samples were collected before the outbreak of ASFV in China from 2016 to 2017 and were confirmed to be negative by the Svanova ASFV p30 antibody detection kit and the Ingenasa ASFV p72 ELISA kit.

Cloning and expression of pB602L

The transmembrane region of pB602L protein was predicted through TMHMM-2.0 (Denmark, http://www.cbs.dtu.dk/services/TMHMM/). The full-length B602L coding region of ASFV SY-18 (GenBank accession, MH766894.1) was synthesized by Tsingke Biotechnology (Shanghai, China) and amplified using primers. F (5[']-CTCGGTACCCTCGAGGGATCCATGGCAGAATTTAATATT GA-3[']) and R (5[']-GACTGCAGGTCGACAAGCTTCTACAATT CTGCTTTTGTAT-3[']).

The recombinant expression plasmid was constructed by ligating the PCR product of the B602L gene into the vector pCold I. The ligation product was transformed into *E.coli* DH5 α , and the recombinant plasmid was successfully constructed by sequencing. Next, the constructed plasmids were transformed

into *E.coli* BL21, and then protein expression was induced for 14–16 h with 0.5 mM IPTG at 16°C. The recombinant pB602L protein was purified by Ni⁺ affinity chromatography and identified by SDS-PAGE and western blots. ASFV-positive pig serum (dilution of 1:5,000) and anti-His (dilution of 1:5,000) were used as the primary antibodies for the western blot assays.

Establishment of an indirect ELISA antibody detection method for ASFV pB602L protein

Square titration

The working concentration of pB602L protein antigen and serum dilution were optimized using square titration. In short, the antigen pB602L was coated on plates at concentrations of 0.625, 0.32, 0.16, 0.08, and 0.04 μ g/mL. Anti-ASFV positive and negative standard sera were diluted at 1:50, 1:100, 1:200, and 1:400. The condition with the highest OD450 ratio of positive and negative sera (P/N value) and an OD450 value of positive sera close to 1.0 was selected as the best working condition.

Working conditions for pB602L-ELISA

In addition to optimal pB602L protein coating concentration and serum dilution, different blocking solutions (5% skim milk, 1% gelatin, and 2% BSA) and reaction times of various materials were also explored, and the optimal reaction time of serum was also tested. The optimal concentration and reaction time of HRP-conjugated anti-pig antibody were tested using the following dilutions: 1:2,500, 1:5,000 and 1:10,000. Furthermore, the TMB response time was optimized. The OD450 and P/N values of these groups were compared to determine the best working conditions.

Cut-off value for pB602L-ELISA

The optimal cutoff value for iELISA was determined using a total of 64 negative serum samples from ASFV-uninfected swnie. The OD450 values were measured, and statistical analysis was performed to calculate the average value (AV) and standard deviation (SD) of the OD450 values. The cut-off value was determined as AV + 3SD (19, 20).

Specificity and sensitivity of pB602L-ELISA

The constructed pB602L-ELISA method was applied against porcine reproductive and respiratory syndrome virus (PRRSV)-, Japanese encephalitis virus (JEV)-, classical swine fever virus (CSFV)-, Getah virus (GETV)-, Porcine circovirus 2 (PCV2)-, Porcine parvovirus (PPV)-, and Pseudorabies virus (PRV)positive sera,ASFV-positive and -negative sera. We diluted ASFV-positive serum from 1:100 to 1:204,800 to determine the highest dilution of serum. We diluted ASFV-positive serum from 1:100 to 1:204,800 to determine the highest dilution of serum. Based on the critical value, the sensitivity of the constructed ELISA method was appraised. Statistical analysis Coefficient of Variation.

Reproducibility of pB602L-ELISA

Six random serum samples were randomly selected, and the established pB602L-ELISA assay was used to conduct intraassays repeatability tests on three different ELISA plates coated in the same batch. In addition, three different batches of coated ELISA plates were randomly selected for inter-assay repeatability experiments. Statistical analysis was performed to calculate intra- and inter-assays variation [coefficient of variation (CV)] between runs.

Comparison of pB602L-ELISA with commercial kits

All 150 clinical serum samples were tested applying the pB602L-ELISA method. The results were compared with the results of a commercial ELISA kit for anti-ASFV detection to evaluate the performance of pB602L-ELISA by calculating the concordance rate as follows: [(true positive + true negative) / (true positive + true negative + true negative) + true positive + true negative + false positive) \times 100%].

Statistical analysis

All data were analyzed using Prism 5 software (GraphPad Software, La Jolla, CA, USA). All data were analyzed using a two-tailed Student's *t*-test. P < 0.05 was considered statistically significant.

Results

Expression and purification of pB602L protein

The pB602L protein had no transmembrane region (Figure 1A). In this study, we expressed the full length of pB602L protein (Figure 1B). The 65-kDa ASFV pB602L protein was successfully expressed. It was detected in the soluble supernatant fraction by WB analysis applying anti-His and ASFV-positive pig serum (Figure 1C).



Establishment of the pB602L-ELISA method

The working concentration of pB602L protein antigen and serum dilution were optimized using square titration. The experimental results show that the optimal coating concentration of antigen for the pB602L-ELISA method is 0.04 μ g/mL, the optimal dilution of serum is 1:100 (Table 1). In addition, other conditions of the pB602L-ELISA method were improved. In brief, the best blocking solution is 2% BSA in PBST, and optimal blocking incubation is 2 h at 37°C (Figure 2A). The optimal reaction time for serum was 1 h at 37°C (Tigure 2B). The optimal dilution of HRP-conjugated anti-pig was 1:5,000, and the optimal reaction condition was 1 h at 37°C (Table 2). Finally, the optimal reaction time for TMB solution was 15 min at 37°C (Figure 2C).

Optimization of the cut-off value

The average OD450 value of negative sera (n = 64) was 0.192, and the SD was 0.049, so the cut-off value of the pB602L-ELISA method was AV + 3SD = 0.340 (Figure 3A). Therefore, samples with an OD450 value of the tested serum of \geq 0.340 are considered positive. Conversely, samples with an OD450 value of <0.340 are considered negative.

Diagnostic specificity of pB602L-ELISA

The established pB602L-ELISA method was used to test anti-PRRSV-, anti-JEV-, anti-GETV-, anti-CSFV-, anti-PCV2-, anti-PPV-, and anti-PRV-positive pig serum. Except for the ASFVpositive control, all sera were negative (OD450 < 0.340), which

| Serum dilution | | | Antigen coating concentration (μ g/mL) | | | | | |
|----------------|-----|--------|---|--------|--------|--------|--|--|
| | | 0.625 | 0.320 | 0.160 | 0.080 | 0.040 | | |
| | Р | 3.585 | 3.209 | 2.715 | 2.424 | 2.363 | | |
| 1:50 | Ν | 0.086 | 0.105 | 0.086 | 0.08 | 0.088 | | |
| | P/N | 41.686 | 30.562 | 31.570 | 30.300 | 26.852 | | |
| | Р | 3.095 | 2.410 | 1.775 | 1.555 | 1.483 | | |
| 1:100 | Ν | 0.093 | 0.120 | 0.088 | 0.066 | 0.080 | | |
| | P/N | 33.279 | 20.083 | 20.170 | 23.561 | 18.538 | | |
| | Р | 2.359 | 1.764 | 1.211 | 1.116 | 1.056 | | |
| 1:200 | Ν | 0.076 | 0.078 | 0.08 | 0.084 | 0.073 | | |
| | P/N | 31.039 | 22.615 | 15.138 | 13.286 | 14.466 | | |
| | Р | 1.599 | 1.240 | 0.951 | 0.740 | 0.769 | | |
| 1:400 | Ν | 0.077 | 0.080 | 0.072 | 0.066 | 0.055 | | |
| | P/N | 20.766 | 15.500 | 13.208 | 11.212 | 13.982 | | |

TABLE 1 P/N values at different conditions.



FIGURE 2

Optimization of pB602L-ELISA method conditions. (A) Optimization of the type of pB602L-ELISA blocking solution and working conditions. The best blocking solution was 2% BSA in PBST, and optimal blocking conditions are 2 h at 37°C. (B) Determination of serum action conditions. The optimal reaction time for serum was 1 h at 37 °C. (C) Determination of TMB solution action conditions. The optimal reaction time for TMB solution was 15 min at 37°C.

TABLE 2 HRP-conjugated anti-pig working conditions.

HRP anti-pig antibody dilution

| HRP anti-pig antibody dilution | | HRP anti-pig antibody action conditions | | | | | |
|--------------------------------|-----|---|----------|------------|----------|--|--|
| | | 37°C 0.5 h | 37°C 1 h | 37°C 1.5 h | 37°C 2 h | | |
| 1:2,500 | Р | 1.082 | 1.268 | 1.263 | 1.464 | | |
| | Ν | 0.300 | 0.255 | 0.233 | 0.268 | | |
| | P/N | 3.607 | 4.973 | 5.421 | 5.463 | | |
| 1:5,000 | Р | 0.774 | 1.048 | 1.192 | 1.248 | | |
| | Ν | 0.183 | 0.163 | 0.195 | 0.203 | | |
| | P/N | 4.230 | 6.429 | 5.519 | 6.148 | | |
| 1:10,000 | Р | 0.526 | 0.784 | 1.003 | 1.065 | | |
| | Ν | 0.169 | 0.175 | 0.234 | 0.176 | | |
| | P/N | 3.112 | 4.480 | 4.286 | 6.051 | | |
| | | | | | | | |



indicated that the established ELISA method has good specificity (Figure 3B).

Diagnostic sensitivity of pB602L-ELISA

The ASFV-positive serum was diluted according to different dilutions. The results showed that the maximum dilution of serum that still had an OD450 value above the cutoff value of 0.340 was 1:6,400, so the sensitivity of the established pB620L-ELISA method was 1:6,400 (Figure 3C), indicating that the established indirect ELISA method is extremely sensitive.

Repeatability of the pB602L-ELISA method

Intra-assay variation and inter-assay variation were calculated to verify the repeatability of the pB602L-ELISA method. The maximum CV of the intra-assay repeatability test was 4.27% (Table 3), and the maximum CV of the inter-assay repeatability test was 5.82% (Table 3), both <10%. These results indicate that the established ELISA method has good repeatability.

Comparison of the pB602L-ELISA method to commercial kits

The performance of the pB602L-ELISA method was compared with that of a commercial ELISA kit (ID Screen $^{(R)}$ African Swine Fever Competition) for anti-ASFV detection in

terms of specificity and accuracy. The results are shown in Table 4. The concordance rate of the detection results of these two methods is 97.33%.

Discussion

ASF is a devastating and economically significant disease that can affect domestic and wild swine (7, 8). It was introduced in China in 2018, and because of its high fatality rate, disease prevention and control has become extremely important for the Chinese pig industry (4). At present, the control of the disease mainly relies on rapid diagnosis and culling of infected pigs and those in close contact with them. ASFV has caused huge economic losses to the global swine industry (21).

ASFV is complex, and its genome encodes many proteins related to immune evasion. There is currently no safe and effective vaccine available (3132). Consequently, it is very significant to develop a method that can quickly and accurately detect ASFV. At present, serological detection techniques and molecular diagnostic methods are still regarded as the main means of identifying infected animals and preventing and controlling ASF (12, 22). There are many methods for detecting viral DNA, including real-time quantitative PCR and in situ hybridization with nucleic acid probes (23). Although these methods have high sensitivity, they also have high requirements on experimental operation technology and the experimental environment (23, 24). Serological diagnosis based on ELISA is not only stable detection means, but it also has good specificity, high sensitivity, simple operation, and low cost, so it is appropriate for the detection of large batches of samples, and hence it is widely used in practice (22). Therefore, the

| Sample number | The result of intra | | | The result of inter | | |
|---------------|---------------------|--------------------|-----------------------|---------------------|--------------------|--------------------------|
| | Average value | Standard deviation | Coefficient variation | Average value | Standard deviation | Coefficient variation |
| 1 | 2.062 | 0.088 | 4.27% | 2.674 | 0.099 | 3.73% |
| 2 | 1.809 | 0.012 | 0.68% | 2.512 | 0.057 | 2.25% |
| 3 | 1.754 | 0.045 | 2.56% | 1.218 | 0.049 | 4.09% |
| 4 | 2.160 | 0.044 | 2.05% | 2.541 | 0.078 | 3.08% |
| 5 | 3.064 | 0.047 | 1.55% | 0.877 | 0.051 | 5.82% |
| 6 | 0.805 | 0.017 | 2.05% | 1.129 | 0.033 | 2.95% |

TABLE 3 The results of the repeating test.

TABLE 4 Comparison of the pB602L-ELISA method and a commercial ELISA kit.

| Detection method | | Commercial ASFV ELISA antibody kit | | | |
|----------------------|----------|---------------------------------------|----------|-------|--|
| | | Positive | Negative | Total | |
| pB602L-ELISA | Positive | 29 | 3 | 32 | |
| | Negative | 1 | 117 | 118 | |
| | Total | 30 | 120 | 150 | |
| Relative sensitivity | | | 96.66% | | |
| Relative specificity | | | 97.50% | | |
| Compliance rate | | | 97.33% | | |

OIE considered ELISA as the first serological method for the diagnosis of ASF (23). Most commercial ELISA assays have been developed based on recombinant ASFV structural proteins such as p72, p54, and p30, and these assays have been used for clinical diagnosis of porcine ASFV infection (25). At present, studies have shown that in addition to the structural proteins of ASFV, such as p72, p54, and p30, its non-structural proteins, such as pK205R and pB602L, also have good immunogenicity, so an effective ELISA method could be established based on these proteins (17, 23).

The pB602L protein of ASFV is encoded by the B602L gene, which contains a central variable region and is frequently used for subgenotyping of ASFV isolates (26). One of the proven markers to characterize this virus is the central variable region (CVR) within the B602L gene (17, 27). pB602L is an important nonstructural protein and a molecular chaperone of the major structural protein p72, which is required for the formation of the icosahedral capsid structure of ASFV (16). The B602L gene contains a central variable region, so the amino acid sequence alignment analysis of the B602L protein of different genotypes of ASFV strains was carried out. The results are shown in of the Supplementary Figures 1, 2. The B602L protein of the strain has only a few amino acid differences, and the B602L protein has a high homology in different genotype. We also

performed an amino acid sequence alignment analysis of the B602L protein of the genotype II ASFV strain prevalent in China. The results are shown in of the Supplementary Figure 3. The alignment results show that the B602L protein of the known prevalent strains in China is not different. In conclusion, the overall antigenicity of ASFV B602L protein is relatively high, and there is only genotype II ASFV strain in China, so the selection of B602L protein has high sensitivity. Previous studies have shown that pB602L has good antigenicity and immunogenicity; the protein can be recognized by domestic pig and wild boar immune sera, and it may serve as a candidate antigen for the development of diagnostic methods for ASFV (28). Consistently high serological responses against pB602L were observed at time points later in infection than responses against the structural proteins in domestic pigs and bushpigs infected with a range of different viral isolates (29, 30). B Gutiérrez-Castañeda et al. show that ASFV B602L protein can be applied to antibody kinetics (17). Consistently high serological responses against pB602L was observed at time points later in infection than responses against the structural proteins in domestic pigs and bushpigs infected with a range of different viral isolates (17). So Gutiérrez-Castañeda et al. (17) suggests that an ELISA against these proteins may be of use in distinguishing animals persistently infected with virus from animals immunized with vaccines incorporating virus structural proteins (17). This suggests that an ELISA against these proteins may be of use in distinguishing animals persistently infected with virus from animals immunized with vaccines incorporating virus structural proteins. Previously, a pB602L-based ELISA assay was used to detect ASFV serum antibodies, which also distinguished pigs persistently infected with native ASFV strains from pigs immunized with subunit vaccines of structural proteins (16). Compared with eukaryotic expression systems, prokaryotic expression systems have lower production costs, simple operation, stable protein preparation, and easy purification and are suitable for large-scale production (31, 32). Therefore, in this study, a prokaryotic expression system was used to express the ASFV pB602L protein. The results revealed that the obtained B602L protein was soluble

and strong antigenicity, and the B602L protein could be served as an ELISA antigen to create and improve an indirect ELISA detection assay.

The indirect ELISA antibody detection method established in this study has a sensitivity of up to 1:6,400. The established method does not cross-react with antibodies against other related swine viruses such as PRRSV, JEV, CSFV, GETV, PCV2, PPV, and PRV. The CV in not only intra-batch but also inter-batch repeatability tests were <10%, which indicated that the pB602L-ELISA method constructed in this research has favorable repeatability. The concordance rate with a commercial ELISA kit (ID Screen[®] African Swine Fever Competition) for anti-ASFV detection was 97.33%. In conclusion, the recombinant ASFV pB602L protein prepared in this study has good antigenicity, and the established indirect ELISA antibody detection method has good sensitivity, good specificity, and high efficiency, and hence it is suitable for the on-site detection of anti-ASFV. This will provide an experimental basis for the detection of ASFV and the specific diagnosis of the disease. It has the potential to become an effective tool for evaluating the humoral immune response in pigs and the protective effects of live attenuated vaccines. Similarly, the construction of ELISA detection method is of great implication for the prevention and treatment of ASF.

Conclusion

In this study, the pB602L-ELISA method for indirect ASFV detection was successfully established. It has good reproducibility and specificity, and production and operation are simple. This study lays the foundation for the development of a kit for large-scale detection of anti-ASFV.

Data availability statement

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found in the article/Supplementary material.

Author contributions

YY: formal analysis, investigation, and validation. QX: formal analysis. QS: methodology and writing-review and

editing. YZ: formal analysis and investigation. YL, ZG, and JZ: investigation and validation. XM: validation. BL and YQ: validation and writing-review and editing. KL: validation. DS: methodology. ZM: conceptualization, funding acquisition, and project administration. JW: conceptualization, formal analysis, funding acquisition, investigation, methodology, and writing-original draft. All authors contributed to the article and approved the submitted version.

Funding

The work was supported by grants from the National Key Research and Development Program of China (Nos. 2021YFD1801300 and 2021YFD1801401 awarded to JW and YQ, respectively), the National Natural Science foundation of China (No. 31941012 awarded to JW), the Project of Shanghai Science and Technology Commission (No. 20392002400 awarded to JW), and the Agricultural Science and Technology Innovation Program (CAAS-ZDRW202203 awarded to ZM). The funders had no role in the study design, data collection and analysis, the decision to publish, or preparation of the manuscript.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/ fvets.2022.971841/full#supplementary-material

References

1. Sang H, Miller G, Lokhandwala S, Sangewar N, Waghela SD, Bishop RP, et al. Progress toward development of effective and safe african swine fever virus Vaccines. *Front Vet Sci.* (2020) 7:84. doi: 10.3389/fvets.2020.00084

2. Nix RJ, Gallardo C, Hutchings G, Blanco E, Dixon LK. Molecular epidemiology of African swine fever virus studied by analysis of four variable genome regions. *Arch Virol.* (2006) 151:2475–94. doi: 10.1007/s00705-006-0794-z

3. Costard S, Mur L, Lubroth J, Sanchez-Vizcaino JM, Pfeiffer DU. Epidemiology of African swine fever virus. *Virus Res.* (2012) 173:191-7. doi: 10.1016/j.virusres.10030

4. Zhou X, Li N, Luo Y, Liu Y, Miao F, Chen T, et al. Emergence of African swine fever in China, 2018. *Transbound Emerg Dis.* (2018) 65:1482–4. doi: 10.1111/tbed.12989

5. Wang T, Sun Y, Qiu HJ. African swine fever: an unprecedented disaster and challenge to China. *Infect Dis Poverty.* (2018) 7:111. doi: 10.1186/s40249-018-0495-3

6. Teklue T, Sun Y, Abid M, Luo Y, Qiu HJ. Current status and evolving approaches to African swine fever vaccine development. *Transbound Emerg Dis.* (2020) 67:529–42. doi: 10.1111/tbed.13364

7. Wang D, Yu J, Wang Y, Zhang M, Li P, Liu M, et al. Development of a real-time loop-mediated isothermal amplification (LAMP) assay and visual LAMP assay for detection of African swine fever virus (ASFV). *J Virol Methods.* (2020) 276:113775. doi: 10.1016/j.jviromet.2019.113775

8. Wang T, Sun Y, Huang S, Qiu HJ. Multifaceted immune responses to African swine fever virus: implications for vaccine development. *Vet Microbiol.* (2020) 249:108832. doi: 10.1016/j.vetmic.2020.108832

9. Ward MP, Tian K, Nowotny N. African swine fever, the forgotten pandemic. *Transbound Emerg Dis.* (2021) 68:2637–9. doi: 10.1111/tbed.14245

10. Miao F, Zhang J, Li N, Chen T, Wang L, Zhang F, et al. Rapid and sensitive recombinase polymerase amplification combined with lateral flow strip for detecting african swine fever virus. *Front Microbiol.* (2019) 10: 1004. doi: 10.3389/fmicb.2019.01004

11. Alonso C, Borca M, Dixon L, Revilla Y, Rodriguez F, Escribano JM, et al. ICTV virus taxonomy profile: asfarviridae. *J Gen Virol.* (2018) 99:613–4. doi: 10.1099/jgv.0.001049

12. Wang F, Zhang H, Hou L, Yang C, Wen Y. Advance of African swine fever virus in recent years. *Res Vet Sci.* (2021) 136:535–9. doi: 10.1016/j.rvsc.2021.04.004

13. Awosanya EJ, Olugasa BO, Gimba FI, Sabri MY, Ogundipe GA. Detection of African swine fever virus in pigs in Southwest Nigeria. *Vet World.* (2021) 14:1840–5. doi: 10.14202/vetworld.2021.1840-1845

14. Jacobs, S. C., Dixon, L. K., Brookes, S. M., and Smith, GL. Expression of African swine fever virus envelope protein j13L inhibits vaccinia virus morphogenesis. *J Gen Virol.* (1998) 79(Pt 5):1169–78. doi: 10.1099./0022-1317-79-5-1169

15. Salas ML, Andres G. African swine fever virus morphogenesis. Virus Res. (2012) 173:29–41. doi: 10.1016/j.virusres.2012.09.016

16. Wang P, Liu C, Wang S, Wen L, Shi Z, Chi Y, et al. Production and application of mouse monoclonal antibodies targeting linear epitopes in pB602L of African swine fever virus. *Arch Virol.* (2022) 167:415-424. doi: 10.1007/s00705-021-05335-0

17. Gutierrez-Castaneda B, Reis AL, Corteyn A, Parkhouse RM, Kollnberger S. Expression, cellular localization and antibody responses of the African swine fever virus genes B602L and K205R. *Arch Virol.* (2008) 153:2303–6. doi: 10.1007/s00705-008-0246-z

18. Simulundu E, Sinkala Y, Chambaro HM, Chinyemba A, Banda F, Mooya L, et al. Genetic characterisation of African swine fever virus from 2017 outbreaks in Zambia: Identification of p72 genotype II variants in domestic pigs. *Onderstepoort J Vet Res.* (2018) 85:e1–5. doi: 10.4102/ojvr.v85i1.1562

19. Zhang S, Wang D, Jiang Y, Li Z, Zou Y, Li M, et al. Development and application of a baculovirus-expressed capsid protein-based indirect ELISA for detection of porcine circovirus 3 IgG antibodies. *BMC Vet Res.* (2019) 15:79. doi: 10.1186/s12917-019-1810-3

20. Zhong K, Zhu M, Yuan Q, Deng Z, Feng S, Liu D, et al. Development of an indirect ELISA to detect african swine fever virus pp62 protein-specific antibodies. *Front Vet Sci.* (2021) 8:798559. doi: 10.3389/fvets.2021.798559

21. Liu Y, Zhang X, Qi W, Yang Y, Liu Z. An T, et al. Prevention and Control Strategies of African Swine Fever and Progress on Pig Farm Repopulation in China. *Viruses.* (2021) 13:2552. doi: 10.3390./v13122552

22. Arias M, Jurado C, Gallardo C, Fernandez-Pinero J, Sanchez-Vizcaino JM. Gaps in African swine fever: analysis and priorities. *Transbound Emerg Dis.* (2018) 65 Suppl 1:235–47. doi: 10.1111/tbed.12695

23. Gaudreault NN, Madden DW, Wilson WC, Trujillo JD, Richt JA. African swine fever virus: an emerging DNA arbovirus. *Front Vet Sci.* (2020) 7:215. doi: 10.3389/fvets.2020.00215

24. Fernandez-Pinero J, Gallardo C, Elizalde M, Robles A, Gomez C, Bishop R, et al. Molecular diagnosis of african swine fever by a new real-time PCR using universal probe library. *Transbound Emerg Dis.* (2013) 60:48–58. doi: 10.1111/j.1865-201201317.x

25. Petrovan V, Yuan F, Li Y, Shang P, Murgia MV, Misra S, et al. Development and characterization of monoclonal antibodies against p30 protein of African swine fever virus. *Virus Res.* (2019) 269:197632. doi: 10.1016/j.virusres.05010

26. Owolodun OA, Bastos AD, Antiabong JF, Ogedengbe ME, Ekong PS, Yakubu B, et al. Molecular characterisation of African swine fever viruses from Nigeria (2003–2006) recovers multiple virus variants and reaffirms CVR epidemiological utility. *Virus Genes.* (2010) 41:361–8. doi: 10.1007/s11262-009-0444-0

27. Gallardo C, Mwaengo DM, Macharia JM, Arias M, Taracha EA, Soler A, et al. Enhanced discrimination of African swine fever virus isolates through nucleotide sequencing of the p54, 72. and pB602L (CVR) genes. *Virus Genes*. (2009) 38:85–95. doi: 10.1007/s11262-008-0293-2

28. Lokhandwala S, Waghela SD, Bray J, Sangewar N, Charendoff C, Martin C, et al. Adenovirus-vectored novel African swine fever virus antigens elicit robust immune responses in swine. *PLoS ONE.* (2017) 12:e0177007. doi: 10.1371/journal.pone.0177007

29. Minoungou GL, Diop M, Dakouo M, Ouattara AK, Settypalli TBK, Lo M, et al. Molecular characterization of African swine fever viruses in Burkina Faso, Mali, and Senegal 1989-2016: genetic diversity of ASFV in West Africa. *Transbound Emerg Dis.* (2021) 68:2842–52. doi: 10.1111/tbed.14240

30. Onzere CK, Bastos AD, Okoth EA, Lichoti JK, Bochere EN, Owido M, et al. Multi-locus sequence typing of African swine fever viruses from endemic regions of Kenya and Eastern Uganda (2011–2013) reveals rapid B602L central variable region evolution. *Virus Genes.* (2018) 54:111–23. doi: 10.1007/s11262-017-1521-4

31. Laage R, Langosch D. Strategies for prokaryotic expression of eukaryotic membrane proteins. *Traffic.* (2001) 2:99–104. doi: 10.1034/j.1600-0854.2001.020204.x

32. Porowinska D, Wujak M, Roszek K, Komoszynski M. Prokaryotic expression systems. *Postepy Hig Med Dosw.* (2013) 67:119–29. doi: 10.5604/17322693.10 38351