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2 **Title**

3 Selection, Design and Immunogenicity Studies of ASFV Antigens for Subunit mRNA
4 Cocktail Vaccines with Specific Immune Response Profiles

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21 **Abstract**

22 Development of safe and effective subunit vaccines for controlling African Swine Fever
23 Virus (ASFV) infection has been hampered by a lack of protective viral antigens, complex
24 virion structures, and multiple mechanisms of infection. Here, we selected ASFV antigens
25 based on their localization on the virion, known functions, and homologies to the subunits
26 of the protective vaccinia virus vaccine. We also engineered viral capsid proteins for
27 inducing optimal antibody responses and designed T cell-directed antigen for inducing
28 broad and robust cellular immunity. The selected antigens in lipid nanoparticle-mRNA
29 formulations were evaluated for immunogenicity in both mice and pigs with concordant
30 results. Different antigens induced divergent immune response profiles, including the
31 levels of IgG and T cell responses and effector functions of anti-sera. We further
32 developed a computational approach to combine antigens into cocktails for inducing
33 specific immune response profiles and validated candidate cocktail vaccines in mice. Our
34 results provide a basis for further evaluating candidate subunit mRNA vaccines in
35 challenge studies.

36
37 **Teaser**

38 Novel strategies to develop subunit vaccines for ASFV and other complex large DNA
39 viruses.

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42 MAIN TEXT

43 44 Introduction

45 African swine fever (ASF) is a highly contagious swine viral disease that causes almost
46 100% mortality in domestic pigs. Currently the United States is free of ASF and incursion
47 to the US swine population could result in approximately \$50 billion losses (1). The
48 causative agent, African swine fever virus (ASFV), is a large DNA virus with a double-
49 stranded DNA genome of 170 to 190 kb, encoding at least 150 proteins (2). Although a
50 live attenuated vaccine was licensed in Vietnam in 2023, it has major safety concerns (3).
51 A safe and effective subunit ASF vaccine is urgently needed.

52 ASF vaccine development faces several major hurdles (4): i) ASFV is classified as a
53 biosafety level 3 (BSL-3) agent. The requirement for BSL-3 containment for any virus
54 work limits the research on the virus and vaccine development. ii) Viral antigens to elicit
55 protective immunity against ASFV infection have not been identified (4), impeding the
56 development of effective subunit vaccines. iii) ASFV exhibits complex multi-layer
57 structures. Both the extracellular enveloped virion (EEV), i.e., with the outer membrane,
58 and intracellular mature virion (IMV), i.e., without the outer membrane, are infectious (5).
59 iv) ASFV can infect host macrophages through multiple entry mechanisms, including
60 receptor-mediated endocytosis, clathrin-mediated endocytosis, macropinocytosis, and
61 phagocytosis (6). As a result, there is no reliable viral neutralization assay for measuring
62 anti-viral antibody responses induced by any single antigen (7). v) Immunological
63 correlates of protective immunity against ASFV are not fully defined. Studies have shown
64 that antibody responses elicited by inactivated virus were not sufficient to protect pigs
65 from infection (8), whereas immune responses by live attenuated virus confer protection
66 (9). Therefore, T cell immunity likely play a pivotal role in the protection. Lymphocyte
67 depletion studies showed that cytotoxic CD8⁺ lymphocytes are important for ASFV
68 clearance (10). IFN- γ responses correlate with the degree of cross-protection against
69 heterologous ASFV challenge (11). Therefore, an effective ASFV vaccine must induce
70 both humoral and cellular immunities.

71 ASFV is the only member in *Asfarviridae*, which belongs to the phylum of
72 nucleocytoplasmic large DNA viruses (NCLDV), characterized by the complex virion
73 structures, large genomes and protein-coding capacity (2). *Poxviridae*, a well-
74 characterized family in NCLDV, is the most closely related to *Asfarviridae*
75 phylogenetically (**Figure S1**). As a key member of *Poxviridae*, the use of vaccinia virus
76 (VACV) as a vaccine has contributed tremendously to the eradication of smallpox disease
77 caused by Variola virus (12). Like ASFV, VACV also produces two infectious virions:
78 EEV and IMV. Studies have shown that subunit vaccines, composing two EEV antigens
79 (B5R and A33R) and two IMV antigens (L1R and A27L), provide complete protection
80 against lethal VACV challenge in mice (13-16), suggesting the feasibility of developing
81 safe and effective subunit vaccines for NCLDV with appropriate viral immunogens.

82 The success of lipid nanoparticle (LNP)-delivered mRNA encoding the spike protein as
83 COVID-19 vaccines motivates broad applications of mRNA-based vaccines. In mRNA
84 vaccines, proteins are synthesized by the host cells and likely maintain native structural
85 conformation, and therefore could induce both humoral and cellular immunities.
86 Antibodies produced by B cells could block viral infection of host cells through
87 neutralization and help to inactivate or eliminate viruses through other effector
88 mechanisms, such as antibody-dependent complement deposition (ADCP), antibody-

89 dependent cellular phagocytosis (ADCP), and antibody-dependent cellular cytotoxicity
90 (ADCC). T cells could recognize virus-infected cells and help to clear the virus through
91 cytokines or directly killing of the infected cells. In addition, mRNA vaccines are
92 especially suited for developing subunit vaccines as multiple mRNAs can be formulated
93 into the same LNP, greatly simplifying the manufacturing process.

94 In this study, we rationally selected and comprehensively evaluated immunogenicity of
95 ASFV antigens, including i) ASFV homologs corresponding to the subunits of the
96 protective VACV vaccine; ii) promising immunogens reported in previous studies; iii)
97 viral capsid proteins in membrane-bound form for more efficient induction of antibody
98 responses; and iv) novel multiple T cell epitopes (MTE) for inducing broad and robust
99 cellular immunity. In mRNA-LNP formulation, the selected ASFV antigens stimulated
100 concordant antigen-specific antibody and T cell responses in both mice and pigs, with the
101 MTE inducing the most potent cellular immunity. By combining the effector function
102 profiles of antigen-specific antibodies and the levels of B and T cell responses induced by
103 different ASFV antigens, we developed optimal antigen combinations for cocktail
104 vaccines with specific immune response profiles, which were further validated in mice.
105 Our study represents a comprehensive investigation of ASF mRNA subunit vaccines
106 incorporating rational antigen selection, protein engineering, T cell-directed antigen
107 design, and profiling of immune responses. Our results provide a basis for further
108 evaluating the subunit mRNA vaccine candidates in challenge studies. The innovative
109 strategies reported here should be applicable to design vaccines for other large complex
110 DNA viruses such as monkeypox.

111 Results

112 Selection and design of ASFV candidate antigens

113 To identify viral antigens for a safe and effective subunit mRNA vaccine for ASF, we
114 selected ASFV antigens using the following approaches. First, we selected ASFV
115 homologs corresponding to the subunits of the protective VACV vaccine because ASFV is
116 most closely related to VACV (**Figure S1**) and both viruses produce two types of
117 infectious virions (EEV and IMV). Four VACV antigens were used in the subunit vaccine,
118 including A33R and B5R from EEV and L1R and A27L from IMV (**Table 1**). Thus, we
119 selected ASFV EP153R, encoding C-type lectin, which is localized on the outer
120 membrane and shares 18% amino acid identity with VACV A33R and ASFV CD2-like
121 protein (CD2v or EP402R), a major outer membrane protein with a size around 41 kDa,
122 which shares 12% amino acid identity with VACV B5R (42 kDa) (**Figure 1A**). L1R,
123 localized on IMV of VACV, is responsible for membrane fusion. In ASFV, both E248R
124 and E199L have been reported to play a similar role (6, 17) and share 14% and 8% amino
125 acid identities with L1R, respectively. VACV A27L exists as a trimer in the capsid on
126 IMV and contributes to attachment to cell heparan sulfate receptor (18, 19). In ASFV, the
127 viral capsid is formed by two proteins, the trimeric P72 and pentameric Penton (20).
128 Because both P72 and Penton are intracellular proteins and exist in multimeric forms in
129 the capsid, we have engineered them as membrane-bound (MB) form to preserve the
130 multimeric structures and to more effectively induce antibody responses (21). In addition,
131 a single mutation (N180Q) was introduced into the MB-Penton to abolish the
132 glycosylation. Therefore, MB-P72 and MB-P_{N180Q} are included here as comparison. For
133 simplicity, MB-P72 and MB-P_{N180Q} are referred to as P72 and Penton in this study.

134 Second, we selected ASFV antigens that have shown to be promising. ASFV P72 and
135 CD2v were reported to induce neutralizing antibodies, which provide partial protection

136 against lethal viral challenge (22-28). ASFV EP153R was shown to contain several T cell
137 epitopes and synergized with CD2v in reducing viremia and disease symptoms (29, 30).
138 ASFV P54 was shown to induce neutralizing antibody response or partial protection
139 against viral challenge (8, 31). ASFV P22 is a highly immunogenic protein with
140 application for serological diagnostics and potential role in maintaining virion structure
141 (32, 33). Hence, besides P72, CD2v and EP153R, P54 and P22, both localized on the viral
142 membrane, were also selected as candidate antigens for evaluation (**Figure 1A**).

143 Third, we developed a novel multi-T cell epitope (MTE) antigen to stimulate strong
144 cellular immunity against ASFV. T cell epitopes were selected based on i) searching
145 IEDB database and literature for experimentally identified epitopes by IFN- γ ELISpot
146 and/or MHC/mass spectrometry using recovered pig lymphocytes, and ii) NetMHCpan
147 prediction of epitopes that bind to the most frequently swine leukocyte antigen (SLA)
148 alleles (SLA-1:0101, SLA-1:0401 and SLA-1:0801) and those with highest binding
149 affinity from abundantly expressed viral proteins (34, 35) were selected. A total of 27
150 epitopes were selected (**Table 2**), including 22 experimentally identified epitopes (6 from
151 EP153R, 4 from PP62, 5 from MGF100-1L, 1 each from A238L, K145R, MGF505-7R,
152 P34, and P37, 2 from P150) (10, 30, 36-40), and 5 predicted epitopes (3 from M448R and
153 2 from MGF505-7R). Notably, the 6 known T cell epitopes from CD2v and 1 from P72
154 were not included in the MTE design as these two antigens were tested separately. All
155 epitopes were fused together with additional 5 endogenous amino acid residues at each
156 end for proteasomal degradation and in an order that avoid disordered polypeptide
157 structure based on Alphafold 2 prediction. In addition, to facilitate the expression of the
158 MTE antigen, the highly expressed ASFV P30 is linked to MTE by a direct GGGs linker
159 (P30-MTE), or a P2A self-cleavage site (P30-P2A-MTE), or an internal ribosome entry
160 site (IRES) signal (P30-IRES-MTE) for direct translation of MTE (**Figure 1B**). Notably,
161 P30 is highly immunogenic and has also been reported to induce neutralizing antibodies
162 (8, 31, 41).

163 In total, we constructed and tested 11 vectors expressing CD2v, EP153R, P72, Penton,
164 P22, E199L, E248R, P54, P30-MTE, P30-P2A-MTE, and P30-IRES-MTE (**Figure 1A-C**
165 and **Figure S2A**). Because specific antibodies are not available for all selected ASFV
166 proteins, except P72, P54 and P30, we added a HA tag at the C-terminus of the most
167 proteins for easy monitoring of their expression. The HA tag was removed for
168 immunogenicity studies in pigs.

169 **Validation of expression of candidate antigens *in vitro***

170 We validated expression of the selected ASFV candidate antigens in cell lines. HEK 293T
171 and/or Vero cells were transfected with individual expression vectors (**Figure S2A**)
172 followed by confocal immunofluorescence imaging with antibodies specific for P72, P54,
173 and P30, and anti-HA antibody for the rest. Abundant fluorescence signals were detected
174 for all vectors in the transfected but not untransfected cells (**Figure 1E**). Expressions of
175 CD2v, P22, P54, EP153R, E199L, and E248R were also detected by Western blotting
176 (**Figure S3**). Notably, CD2v and EP153R were highly glycosylated, consistent with
177 previous reports (42, 43). Expression of P72 and Penton were validated previously (21).
178 To validate co-expression of P30 and MTE, transfected HEK 293T cells were co-stained
179 with anti-P30 and anti-HA antibodies, followed by confocal microscopy. Both P30 and
180 MTE from all three different designs were readily detected in the cytosol of transfected
181 cells (**Figure 1F**). Thus, the selected ASFV antigens can be expressed in cell lines.

182 Next, we prepared mRNA for each candidate antigen by *in vitro* transcription and
183 formulated the mRNA in LNP. Briefly, DNA fragments encoding P72, Penton, CD2v,
184 EP153R, E199L, E248R, P22, P54, P30-MTE, P30-P2A-MTE, and P30-IRES-MTE were
185 inserted into a pUC plasmid containing the T7 promoter, 5'UTR, 3'UTR, and polyA
186 (**Figure 1D** and **Figure S2B**). Plasmids were linearized and used as templates for *in vitro*
187 transcription using T7 RNA polymerase and ATP, GTP, CTP and pseudo-UTP. mRNAs
188 were capped, purified, and formulated into LNPs using ionizable lipid, DSPC, cholesterol
189 and DMG-PEG-2000 as described (44). The LNP-mRNA formulations were further
190 subject to physicochemical analysis and results showed an average particle size of 80 –
191 120 nm in diameter and a low poly dispersity index (PDI) of 0.1 to 0.2 (**Table S1**),
192 suggesting that the LNP-mRNA particles have a uniform particle size distribution, optimal
193 for particle internalization and biodistribution. LNP-mRNAs were transfected into HEK
194 293T cells and expression of all candidate antigens was readily detected by flow
195 cytometry (**Figure 1G**). These results suggest that formulated LNP-mRNAs are efficiently
196 translated.

197 **Induction of antibody and T cell responses in mice by LNP-mRNA vaccination**

198 We first evaluated the immunogenicity of the candidate antigens in mice. BALB/c mice at
199 6-8 weeks of age were divided into 12 groups with 5 mice per group. Eleven groups were
200 immunized with 11 LNP-mRNAs expressing different antigens twice at day 0 and day 21
201 (**Figure 2A**). The other group was injected with sterile PBS and served as control. Blood
202 was collected before immunization and at day 14 and 35 after immunization for assessing
203 antigen-specific IgG responses by ELISA. Spleen was collected at day 35 for assessing
204 antigen-specific T cell responses by ELISPOT. Compared to before immunization,
205 antigen-specific IgG responses, as indicated by the endpoint titers, were significantly
206 induced in all immunized mice 14 days after the first immunization and further boosted at
207 day 35 (**Figure 2B**). However, the levels of IgG titers varied significantly among the
208 different antigens. For example, 14 days after the first immunization antigen-specific IgG
209 titer was only 1.2-fold over the background for E248R whereas the titer was 125.2-fold for
210 P22. Fourteen days following the boost, antigen-specific IgG titer was increased to 126
211 (2.2-fold) for E248R and to 521,675 (83-fold) for P22. Among the 11 different antigens,
212 P22, P30-IRES-MTE and P30-P2A-MTE induced the highest titers of antigen-specific
213 IgG responses after boost, followed by E199L, P72, Penton, P54 and CD2v, with the
214 lowest for EP153R, E248R and P30-MTE (**Figure 2B**). Similarly, LNP-mRNA
215 immunization induced significant antigen-specific T cell responses as indicated by
216 secretion of cytokines IFN- γ and TNF- α following re-stimulation of splenocytes with
217 either specific recombinant proteins or P72 peptides (**Figure 2C-D**). T cell responses also
218 varied significantly among different antigens. For example, CD2v stimulated highest level
219 of cytokine-secreting spots (IFN- γ : 327, TNF- α : 992), followed by EP153R (IFN- γ : 116,
220 TNF- α : 414), P54 (IFN- γ : 59, TNF- α : 346), Penton (IFN- γ : 54, TNF- α : 593), P72 (IFN- γ :
221 30, TNF- α : 216), and P22 (IFN- γ : 29, TNF- α : 223). These results show that mRNA
222 expressing the selected ASFV antigens induce both antibody and T cell responses, but the
223 magnitudes of the immune responses vary considerably among the different antigens.

224 The three MTE designs also induced significantly different immune responses. P30-IRES-
225 MTE induced the highest titer of anti-P30 IgG (**Figure 2B**) and highest numbers of MTE-
226 specific IFN- γ and TNF- α spots (**Figure 2E-F**), followed by P30-P2A-MTE. P30-MTE
227 induced the lowest antibody and T cell responses. Notably, the MTE-specific T cell
228 responses, as indicated by the numbers of IFN- γ and TNF- α ELISpots, were significantly
229 higher than P30-specific responses, indicating that the T cell-directed antigen (MTE) is

230 potent. Compared to P30-MTE, where P30 is fused with MTE via a GGS linker, in P30-
231 P2A-MTE, P30-MTE is synthesized as a single polypeptide that is cleaved at P2A site,
232 and in P30-IRES-MTE, P30 and MTE are synthesized separately through the internal
233 ribosomal entry site (IRES). Our results show that a complete separation of P30 and MTE
234 translation through IRES is most effective for inducing both P30-specific antibody
235 responses and MTE-specific T cell responses.

236 ASFV induced hemadsorption is characterized by the adherence of red blood cells to the
237 surface of infected cells (45). This phenomenon is primarily mediated by viral protein
238 CD2v expressed on the surface of infected cells. To test if CD2v-specific antibodies
239 inhibit hemadsorption, sera from CD2v LNP-mRNA immunized mice were first incubated
240 with CD2v-expressing HEK 293T cells and then pig red blood cells (RBC) were added for
241 surface adherence. Hemadsorption was observed by confocal microscopy after anti-RBC
242 antibody staining. As shown in **Figure 2G**, sera from unimmunized mice did not show
243 significant inhibition of hemadsorption or “rosette” formation surrounding CD2v-
244 expressing cells, while the sera from CD2v LNP-mRNA immunized mice inhibited
245 “rosette” formation. Quantification of percentages of inhibition showed that CD2v
246 immunized sera yielded 64% inhibition of hemadsorption as compared to 13% with
247 control sera (**Figure 2H-I**). These results show that although CD2v-specific IgG titer is
248 relatively low compared to those induced by other ASFV antigens, the antibodies can
249 specifically block CD2v-mediated hemadsorption.

250 **Induction of antibody and T cell responses in pigs by LNP-mRNA vaccination**

251 We next evaluated immunogenicity of the following ASFV antigens in pigs based on their
252 induction of relatively higher levels of antibody and T cell responses in mice, including
253 P72, Penton, P22, E199L, P54, CD2v, EP153R, P30-IRES-MTE, P30-P2A-MTE.
254 Commercial piglets at 6 weeks of age were immunized with 30 μ g mRNA in LNP
255 formulation (4 pigs per group) twice at day 0 and day 21 (**Figure 3A**). Two pigs were
256 injected with sterile PBS and served as control. Blood was collected before immunization
257 and every 7 days after immunization for assessing antigen-specific IgG responses by
258 ELISA. Spleen were collected at day 35 for assessing antigen-specific T cell responses by
259 ELISPOT. No clinical signs were observed throughout the study and immunized pigs
260 gained as much weight as control pigs, suggesting that LNP-mRNA is safe for use in pigs.

261 Antigen-specific IgG responses, as indicated by the endpoint titers, became detectable in
262 all immunized pigs 14 days after the first immunization, increased steadily afterwards, and
263 reached the highest level at day 35, i.e., 14 days after boost (**Figure 3B**). As observed in
264 mice, the levels of antigen-specific IgG titers varied significantly among the different
265 antigens. Among the 9 antigens, P72, P22, E199L and P30-IRES-MTE induced the
266 highest titers of antigen-specific IgG responses at day 35, followed by Penton and P30-
267 P2A-MTE, with the lowest for CD2v, EP153R, and P54. Similarly, LNP-mRNA
268 immunization induced significant antigen-specific T cell responses as indicated by
269 secretion of cytokines IFN- γ following re-stimulation of splenocytes with either
270 recombinant proteins or P72 peptides or MTE peptides (**Figure 3D**). T cell responses also
271 varied significantly among different antigens. For example, P30-IRES-MTE stimulated
272 highest numbers of antigen-specific IFN- γ -secreting spots (103), followed by CD2v (74)
273 and P30-P2A-MTE (73). These results show that mRNA expressing the selected ASFV
274 antigens induces both antibody and T cell responses in pigs, but the magnitudes of the
275 immune responses vary considerably among the different antigens.

276 We analyzed the correlations of antibody and T cell responses for different antigens
277 between pigs and mice. IgG responses induced by the same antigens in pigs and mice
278 were highly correlated with Pearson correlation coefficient of 0.93 ($p < 0.005$) (**Figure 3D**).
279 Similar, T cell-mediated IFN- γ responses induced by the same antigens in pigs and mice
280 were also highly correlated with Pearson correlation coefficient of 0.87 ($p < 0.005$) (**Figure**
281 **3E**). The similar results between mice and pigs cross-validate the two studies and also
282 suggest that mice can be used to replace pigs for assessing immunogenicity of ASF
283 subunit vaccines in most cases.

284 **Distinct effector functions by antibodies specific for different ASFV antigens**

285 Due to the large size and structural complexity of virion, ASFV can infect host
286 macrophages through multiple mechanisms, including receptor-mediated endocytosis,
287 clathrin-mediated endocytosis, macropinocytosis, and phagocytosis. As a result, viral
288 neutralization assay with sera from immunization with a single ASFV antigen is not
289 reliable (7) and antibody-mediated neutralization is not sufficient to confer protection (8).
290 To identify potential host-dependent antiviral functions of antibodies induced by different
291 ASFV antigens, we determined their effector functions, including ADCD, ADCC and
292 ADCP.

293 To measure ADCD, CHO cells stably expressing individual ASFV antigens were
294 incubated with heat-inactivated pig serum, followed by addition of non-heat-inactivated
295 serum from placebo pigs as a source of complement, and cell lysis was quantified by flow
296 cytometry (**Figure 4A** and **S4A**). ADCD activities of sera varied depending on the
297 immunizing antigens, but overall, the sera from EP153R, P72, Penton, and P30-IRES-
298 MTE-immunized pigs had significantly higher ADCD activities than sera from CD2v,
299 P22, P54, E199L, and P30-P2A-MTE-immunized pigs (25-31% vs. 9-17%, **Figure 4B**).

300 To measure ADCC, CHO cells stably expressing individual ASFV antigens were
301 incubated with heat-inactivated pig serum, followed by addition of pig peripheral blood
302 mononuclear cells (BPMCs) as a source of natural killer (NK) cells, and lysis of CHO
303 cells was quantified by a luminescent assay (**Figure 4C**). Sera from P54-immunized pigs
304 induced highest CHO cell lysis (76%), followed by sera from EP153R (64%), Penton
305 (61.2%), P72 (56%), CD2v (51.2%), and P22 (48.6%) immunized pigs, and sera from
306 E199L, P30-IRES-MTE and P30-P2A-MTE had lowest cell lysis (19-30%) (**Figure 4D**).

307 To measure ADCP, fluorescent beads were coated with specific ASFV proteins and
308 incubated with immune sera and a pig macrophage cell line 3D4/31, and phagocytosis of
309 labeled beads was quantified by flow cytometry (**Figure 4E** and **S4B**). Sera from P72 and
310 Penton-immunized pigs induced highest levels of ADCP (**Figure 4F**), which were 2-3-
311 fold higher than the ADCP induced by sera from pigs immunized with CD2v, EP153R,
312 P22, P54 or E199L. Notably, ADCP activity was lowest for sera from pigs immunized
313 with P30-P2A-MTE, despite robust anti-P30 IgG responses.

314 These results show that antibodies induced by different ASFV antigens exhibit different
315 effector functions, which could be exploit for developing cocktail vaccines with desired
316 immune profiles.

317 **Identification of antigen combinations for cocktail vaccines by computational** 318 **analysis**

319 To identify optimal antigen combinations, we performed computational analysis of five
320 immune parameters: T cell response (IFN- γ), antigen-specific IgG level, ADCD, ADCC,
321 and ADCP. Average values from four pigs per antigen were calculated and normalized by
322 ranking each immune parameter from 1 (lowest) to 8 (highest). The ranking revealed
323 distinct patterns: P22 and P30 induced the highest IgG responses; CD2v and EP153R
324 induced the highest T cell response (IFN- γ); Penton and EP153R induced antibodies
325 showing the highest ADCD activities; P54 and EP153R induced antibodies with the
326 highest ADCC activities; and Penton and P72 induced antibodies with the highest ADCP
327 activities (**Figure 5A**). When the magnitude of each immune parameters from individual
328 antigen was taken into consideration, similar pattern was also found in the chord diagram
329 showing correlations between antigens and immunological categories (**Figure 5B**).
330 Correlation analysis of the five immune parameters across all antigens indicated a positive
331 correlation between Fc-mediated ADCD and ADCP, and between ADCC and T cell
332 response (IFN- γ) (**Figure 5C**). However, IgG response was negatively correlated with
333 IFN- γ and ADCC.

334 Optimal antigen combinations were scored by summing the ranks of selected immune
335 parameters. Among the eight antigens tested (CD2v, EP153R, P72, Penton, P22, P54,
336 E199L, P30), we developed all possible 3-way, 4-way, and 5-way combinations based on
337 rank sums of the five immune parameters ("IgG", "IFN- γ ", "ADCD", "ADCC", "ADCP")
338 (**Figure 6**). For the highest response across five, four or three parameters, the optimal 3-
339 way combination was EP153R_P72_Penton (**Figure 6A**) and the optimal 4-way
340 combination was EP153R_P54_P72_Penton (**Figure 6B**). Although the top 5-way
341 combinations across five, four or three parameters were different, but they all included
342 EP153R, P72 and Penton (**Figure 6C**). When ADCP was excluded because of the
343 potential to enhance ASFV infection of macrophages due to antibody-dependent
344 enhancement (46, 47), the best 3-way cocktail remained EP153R_P72_Penton, with
345 addition of P54 for the 4-way and addition of P54 and P30 for the 5-way combinations.
346 Additionally, when only IgG and T cell responses were considered, the top 3-way
347 combinations were CD2v_P30_P72 or CD2v_E199L_P72, the top 4-way combination
348 was CD2v_E199L_P30_P72, and the top 5-way combination was
349 CD2v_E199L_P30_P72_Penton (**Figure 6**). Although MTE-induced T cell responses and
350 CD2v antibody-mediated hemadsorption inhibition activity were not considered here,
351 optimal combinations for targeted immune profiles provide novel insights for developing
352 cocktail mRNA vaccines.

353 **Induction of robust antibody and T cell responses by cocktail mRNA vaccine** 354 **candidates**

355 We evaluated cocktail vaccines with antigen combinations based on computational
356 analysis and the following considerations. All the cocktails contained P30-IRES-MTE and
357 P72 with the former stimulating robust T cell immunity and the latter as a major capsid
358 protein for inducing antibody responses with partial protection (**Figure 7A**). P72 was also
359 identified as one of the three antigens in the optimal 3-way combination based on the
360 highest response across five, four or three parameters above (**Figure 6**). CD2v was
361 included in cocktails 1-3 to provide maximal induction of T cell immunity due to
362 abundant T cell epitopes present in CD2v and induction of antibodies that inhibit
363 hemadsorption, which is critical for controlling viral spread (24, 48). Compared to cocktail
364 1, penton was included in cocktail 2 as penton is a key capsid component, was identified
365 as one of the three antigens in the optimal 3-way combination, and no study has examined

366 whether antibodies against penton is protective. Cocktail 3 contains another highly
367 immunogenic protein, E199L and Cocktail 4 contained P22.

368 We evaluated the immunogenicity of candidate cocktail vaccines in mice with 5 µg of
369 each mRNA in LNP formulations, the same amount as used in individual immunizations.
370 BALB/c mice were immunized with the cocktail vaccines twice at day 0 and day 21 or
371 given PBS as control. Blood was collected before immunization and at day 14 and 35 after
372 immunization for assessing antigen-specific IgG responses by ELISA. Spleen was
373 collected at day 35 for assessing antigen-specific T cell responses by ELISPOT. Antigen-
374 specific IgG was detected 14 days after the first immunization and the titers were
375 significantly increased (6-528-fold) 14 days following boost (**Figure 7B**). At day 35, IgG
376 titers for P30 and P22 were the highest, the same as observed in mice immunized with
377 P30-IRES-MTE or P22 individually (**Figure 2B**). The IgG titers for the rest of antigens
378 were similar, ranging from 2×10^3 to 1.6×10^4 . Similarly, MTE induced the highest level of
379 T cell response as indicated by the numbers of IFN- γ ELISPOTs in all cocktails (**Figure**
380 **7C**), followed by CD2v, with the lowest for P72.

381 To further determine whether the immune responses elicited by specific antigens in the
382 cocktail vaccines parallel to those elicited by individual antigens, Pearson correlation
383 analysis was conducted. The analysis revealed correlation coefficients of 0.95 for IgG
384 antibody responses and 0.80 for T cell responses (**Figure 6D**). Similar levels of antigen-
385 specific IgG and T cell responses induced by specific antigens from immunization with
386 individual LNP-mRNA versus the cocktail LNP-mRNA suggests that the inclusion of 3 or
387 4 mRNA in the same LNP formulation does not interfere with induction of immune
388 responses to each antigen in the cocktail. These results also show that the cocktail ASFV
389 vaccines induce potent antibody and T cell responses.

390 Discussion

391 As the first step toward developing a safe and effective subunit vaccine for ASF, in this
392 study we rationally selected and designed viral antigens, evaluated their immunogenicity
393 in mice and pigs, developed cocktail vaccines with specific immune response profiles, and
394 validated the induction of robust humoral and cellular immunities of the selected
395 combinations. We chose mRNA platform for a potential ASFV subunit vaccine for the
396 following reasons: First, although immunological correlates for a protective ASFV
397 vaccine is not fully defined, studies have shown that antibody responses elicited by
398 inactivated virus were not sufficient to protect pigs from infection (8), whereas immune
399 responses induced by live attenuated virus confer protection (9). Therefore, T cell
400 immunity likely plays a pivotal role in the protection, and an effective ASFV vaccine must
401 induce both humoral and cellular immunities. Second, in mRNA vaccine, immunogens are
402 expressed in native conformation by the host cells and have a higher chance to induce
403 antibodies that recognize the native viral antigens (conformational epitopes), as well as
404 induce CD8⁺ T cell responses that are effective in clearing virus from infected cells. Third,
405 multiple mRNAs can be easily formulated into the same LNP, greatly simplifying the
406 development of a cocktail vaccine.

407 A key aspect of subunit vaccine development is the identification of immunogens to
408 include in the vaccine. This is especially challenging for ASFV. ASFV has a large
409 genome, encoding more than 150 open reading frames, and many viral proteins have not
410 been characterized. Although some viral antigens, such as P72, P30 and P54, have been
411 shown to confer partial protection by inducing neutralizing antibodies, viral antigens that

412 induce protective immune responses have not been defined. In addition, ASFV has
413 complex structures and both EEV and IMV are infectious. Likely, multiple viral antigens
414 must be combined in order to induce protective immunity.

415 In our selection and design of ASFV antigens, we used the following four novel
416 approaches: First, we selected ASFV antigens that are homologs of the subunits of the
417 protective VACV vaccine (13, 14). *Asfarviridae* is phylogenetically most closely related
418 to *Poxviridae*. A VACV subunit vaccine with two antigens from EEV and two antigens
419 from IMV confers complete protection against lethal challenge in mice (14, 15, 49), which
420 was correlated with induction of serum-neutralizing antibodies and vaccinia virus-specific
421 CD8⁺ T cells (50, 51). Thus, we selected the corresponding ASFV antigens based on the
422 same localization on virion, similar functions, and amino acid sequence homology (Table
423 1). Second, we selected ASFV antigens based on the recent identification of P72 as the
424 major capsid protein and Penton as a minor capsid protein (20). In particular, Penton has
425 never been investigated as a vaccine antigen. Third, even for promising ASFV antigens
426 identified previously, we designed antigen through protein engineering in order to induce
427 strong immune responses. For example, we expressed P72 and Penton, both are
428 intracellular proteins, in membrane-bound form, which form multimeric structures without
429 viral chaperone and induced stronger antibody responses (21). Fourth, we designed T cell-
430 directed (MTE) antigen to induce broad and robust cellular immunity. MTE contains 27
431 experimentally identified and computationally predicted T cell epitopes with majority
432 being CD8 T cell epitopes. To facilitate MTE expression, we linked MTE to the
433 abundantly expressed P30 through three strategies, GGS linker, P2A, and IRES. *In vivo*
434 testing of the three MTE designs showed that LNP-mRNA expressing P30-IRES-MTE
435 stimulated strongest T cell responses (Figures 2 and 3), probably due to the direct
436 initiation of MTE expression via IRES element. Based on these four criteria, we selected
437 eleven ASFV antigens: two each from the outer membrane and capsid, four from the inner
438 membrane, and one from the viral shell (Figure 1A).

439 We evaluated the immunogenicity of the selected ASFV antigens in LNP-mRNA
440 formulation in both mice and pigs with concordant results. However, the magnitude of
441 antibody and T cell responses induced by different ASFV antigens was quite different. For
442 example, P30, P22, P72 and E199L induced robust antibody responses, but T cell
443 responses to P30 and P22 were low (Figures 2 and 3). In contrast, CD2v and EP153R
444 induced only modest antibody responses, probably due to their high level of glycosylation
445 (Figure S3), but induced robust T cell responses. Our findings are in line with previous
446 reports of abundant T cell epitopes harbored by CD2v and EP153R (30). Furthermore,
447 consistent with previous observations, anti-sera induced by CD2v mRNA vaccine are
448 effective in inhibiting hemadsorption, suggesting the functionality of the induced
449 antibodies. Immunogenicity of E199L and Penton has not been investigated, our results
450 show that E199L is quite immunogenic and the engineered Penton (in membrane-bound
451 form with N180Q mutation to prevent glycosylation) also induce both antibody and T cell
452 responses. Our results also show that MTE antigen induced the highest level of T cell
453 responses, suggesting the validity of our design. Induction of divergent antibody and T
454 cell responses by different ASFV antigens suggest the need to strategically combine
455 antigens for inducing a balanced immune response for optimal protection (see below).

456 Another major challenge of ASFV vaccine development is a lack of reliable viral
457 neutralizing assay due to infection of host macrophages by both EEV and IMV through
458 receptor-mediated endocytosis, clathrin-mediated endocytosis, macropinocytosis, and

phagocytosis. This is further compounded by the requirement of BSL3 containment for any live ASFV work. To get around these obstacles, we determined effector functions, including ADCD, ADCC and ADPC, of anti-sera from pigs immunized with different ASFV antigens. The same as antibody and T cell responses, anti-sera induced by different antigens exhibited divergent effector functions (Figures 4 and 5). Nevertheless, Fc-mediated effector functions, ADCD and ADPC, were correlated, so as ADCC and T cell response (IFN- γ) (Figure 5C). Interestingly, IgG response was negatively correlated with IFN- γ and ADCC. To develop cocktail vaccines with desired immune response profiles, we used computational analysis to rank order antigen combinations based on the five semi-quantitative immune parameters: the levels of IgG responses, the levels of T cell responses (IFN- γ), ADCD, ADCC and ADPC. Notably, the top 3-way combination based on 5-, 4-, and 3-parameters was EP153R, P72 and Penton (Figure 6A). In the 4-parameter analysis, we removed ADPC for possible concern due to antibody-dependent enhancement (ADE), which has been observed for flavivirus infection of macrophages (52, 53). In the 3-parameter analysis, we further removed ADCD due to its unidentified role in disease protection. In addition, we further considered MTE-induced T cell responses and inhibition of hemadsorption by CD2v-specific antibodies. We tested four cocktail vaccines and found that antibody and T cell responses induced by specific antigen in the cocktails are highly correlated with immune responses induced by single antigen immunization (Figure 7). These results suggest that strategic combination of different antigens in LNP-mRNA vaccines are likely achieve specific immune responses profiles for optimal protection.

In summary, we have used a comprehensive approach to dissect the immune response profiles of rationally selected ASFV antigens, and developed new strategies for antigen combination for developing a safe and effective mRNA-based subunit vaccine for ASFV. The methodologies developed in this study could also be applied to develop subunit vaccines for other pathogens with large genomes, including the monkeypox virus.

Limitation of the study

Our study has two primary limitations: First, we did not assay neutralizing activity of antibodies induced by different ASFV antigens. This is because developing neutralization assays for antibodies elicited by a single ASFV antigen has been challenging due to multiple entry pathways used by different infectious viral particles. The presence of neutralizing antibodies has been associated with live vaccine-induced protection in one report (54), whereas many studies suggest that neutralizing antibodies to P30, P54, and P72 are insufficient for protection (8), highlighting the critical role of T cell responses in vaccine-induced protection (55). Given the multilayered structures of ASFV and its various entry mechanisms, both neutralizing antibodies and cellular immunity are likely required for blocking viral infection and clearing infected cells. Additionally, antibodies can control infection through mechanisms beyond neutralization. Due to the limited studies on ASFV, the roles of Fc-mediated complement recruitment and engagement of innate immune cells, such as NK cells and macrophages, should not be underestimated, as previous reports have shown that antibody effector functions are associated with protection against respiratory syncytial virus (56), influenza virus (57), and malaria parasites (58). Our study delineates the antibody effector functions and other immune parameters by selected ASFV antigens, offering critical insights for the development of cocktail vaccines. Second, we did not assess protective efficacy of cocktail vaccines under challenge conditions due to the select agent status, restricted biocontainment requirements, and high cost for working with ASFV infection in pigs. Clearly, this should be a priority

507 for moving forward. In addition, we selected ASFV antigens based on the current
508 knowledge of ASFV structure and protein functions. It cannot be excluded that other
509 proteins exhibiting similar localization or functions may be discovered in the future and
510 are worthwhile for testing.

512 **Materials and Methods**

513 **Cells and plasmids**

514 Human embryonic kidney cells (HEK-293T) and Vero E6 cells were maintained in
515 minimum essential medium supplemented with 10% fetal bovine serum, antibiotics (100
516 units/ml of penicillin and 100 mg/ml of streptomycin) and fungizone (0.25 mg/ml) at 37
517 °C with 5% CO₂. Porcine macrophage 3D4/31 cell line was maintained in RPMI with
518 essential supplements including FBS, antibiotics, and fungizone. FreeStyle™ CHO-S cells
519 were obtained from ThermoFisher and cultured in Freestyle™ CHO Expression Medium
520 supplemented with 8 mM L-glutamine at 37°C, 8% CO₂ on an orbital shaker platform
521 rotating at 135 rpm. ASFV P72 (B646L), Penton (H240R), CD2v (EP402R), P22
522 (KP177R), EP153R (C-type lectin), E248R, E199L, P54 (E183L), P30 (CP204L) genes
523 from Georgia 2007/1 strain (NCBI Reference Sequence: NC_044959.2) were synthesized
524 by GenScript and inserted into the pHCMV mammalian cell expression vector (MoBiTec).
525 P72 and Penton were engineered for membrane anchoring by addition of secretion signal
526 peptide from human CD8α (GenBank ID: NP_001139345.1) to the N terminus and CD8α
527 stalk region or hinge, transmembrane region, and a short cytoplasmic tail to the C
528 terminus (21).

529 **Design of T cell-directed vaccine**

530 ASFV-derived CD4⁺ and CD8⁺ T cell epitopes were searched on the Immune Epitope
531 Database (IEDB) (<https://www.iedb.org/>) and limited to those with positive experimental
532 results for IFN-γ ELISpot and/or cellular MHC/mass spectrometry ligand presentation. A
533 total of 22 T cell epitopes from PP62, MGF100-1L, A238L, K145R, MGF505-7R, P34,
534 P150, P37, and EP153R were selected for their strong induction of cellular immunity
535 according to previous publications (10, 30, 37, 40, 59, 60). P72 and CD2v contains
536 multiple T cell epitopes which were not included in the T cell-directed vaccine design.
537 Additionally, the online server, NetMHCpan 4.1
538 (<http://www.cbs.dtu.dk/services/NetMHCpan/>) was used to predict binding affinities of all
539 8-11mer peptides in the most abundantly expressed pp220 polyprotein. The most frequent
540 Swine Leukocyte Antigens (SLA-1:0401, SLA-1:0101, SLA-1:0801) were selected for
541 peptide binding. Strong binders with an % Rank-Eluted ligand (EL) <0.1 and affinity <200
542 mM was selected and included 3 epitopes from M448R and 2 epitopes from MGF505-7R.
543 The T cell-directed vaccine construct, multiple-T cell epitope (MTE), contained a Kozak
544 sequence followed by P30, Internal Ribosome Entry Site (IRES) or porcine teschovirus-1
545 2A (P2A) or GGS linker, and individual T cell epitopes with additional 5 amino acid
546 residues (AA) on each side were fused together, plus a HA tag for MTE detection. MTE
547 was synthesized by GenScript and cloned into expression vector with P30, namely P30-
548 MTE, P30-IRES-MTE, and P30-P2A-MTE.

549 **Lipid Nanoparticle (LNP) formulation of mRNA and validation**

550 Genes encoding the eleven candidate antigens were inserted into pUC57 vector, which
551 contains essential elements for mRNA *in vitro* transcription, such as T7 promoter, UTRs,
552 and polyA tail. The linearized plasmids were subject to *in vitro* transcription using

553 pseudouridine-5'-triphosphate or pseudo-UTP (44). After mRNA purification and capping
554 reaction, an aqueous phase of mRNA was prepared by diluting mRNA stock in 10 mM
555 citrate buffer. The organic phase of lipid nanoparticles was prepared by adding 200 proof
556 ethanol with lipid stock solutions which contained ionizable lipid L002 (Advanced RNA
557 Vaccine, ARV), DSPC helper lipid (Avanti Polar Lipids), cholesterol (Avanti Polar
558 Lipids), and DMG-PEG-2000 (Avanti Polar Lipids) . LNP-mRNA formulations were
559 prepared by mixing organic and aqueous phases at a ratio of 1:3. RiboGreen assay
560 (Thermo Fisher) was performed by following manufacture's instruction to quantify
561 mRNA after formulation. Encapsulation efficiency was measured by Picogreen assay. To
562 validate LNP delivery of mRNAs, HEK-293T cells were seeded onto 24-well tissue
563 culture plate and 500 ng of LNP-mRNA diluted in Opti-MEM was added in individual
564 well. Cells were collected in 48 h and protein expression was analyzed by flow cytometry
565 and confocal microscopy as described below.

566 **Flow cytometry**

567 HEK-293T cells seeded in 6-well tissue culture plate and transfected with 3 µg
568 recombinant pHCMV plasmids expressing individual proteins using the linear 25 kDa
569 polyethylenimine (PEI; Santa Cruz Biotechnology) at a 3:1 mass ratio of PEI to DNA. At
570 48 h post transfection, cells were trypsinized and washed with 1 mL FACS buffer
571 followed by direct surface staining of P72 using mouse anti-ASFV P72 mAb
572 (MyBioSource, San Diego, CA), mouse anti-P30 monoclonal antibody (Aviva Systems
573 Biology, OAEF00154), P54 mAb (GeneTex, GTX635690), or mouse mAb specific for
574 HA tag (GenScript). Alexa Fluor™ 488-conjugated Goat anti-Mouse IgG (H+L) was used
575 as the secondary antibody and cells nuclei were counterstained with DAPI (4',6-
576 Diamidino-2-Phenylindole, Dilactate) before data acquisition on a BD LSR Fortessa HTS-
577 2 cytometer. Results were analyzed using FlowJo v10 software.

578 **Confocal microscopy**

579 Vero E6 cells were grown on glass-bottom 35-mm cell culture dishes (MatTek). In 48
580 hours after transfection of MTE expression vectors, cells were fixed by 4%
581 paraformaldehyde at room temperature (RT) for 15 min followed by permeabilization with
582 0.5% Triton X-100 for 10 min and then blocked with 2% bovine serum albumin for 30
583 min. Cells were incubated with mouse anti-P30 mAb (GenScript) and Rabbit anti-HA tag
584 polyclonal antibody (Thermo Fisher) at 37C for 1 hour. Alexa Fluor™ 488-conjugated
585 goat anti-mouse IgG and Alexa Fluor® 594 AffiniPure™ goat anti-rabbit IgG (H+L)
586 (Jackson ImmunoResearch) were used as secondary antibody. Cells were counterstained
587 with DAPI before proceeding with imaging under a confocal microscope (Nikon A1R
588 HD25, Nikon). Images were processed using the program ImageJ (<https://imagej.net/Fiji>).

589 **Mouse study**

590 To assess immunogenicity of individual antigens in mice, six to eight-week-old female
591 BALB/c mice were purchased from Charles River and housed in animal facility at the
592 Massachusetts Institute of Technology (MIT). All animal ethical and welfare standards
593 were met in this study and experiments were approved by the Institutional Animal Care
594 and Use Committee at MIT under protocol number 0322-021-25. Briefly, 5 mice were
595 assigned to each group and immunized intramuscularly with 50 µL 5 µg LNP-mRNA
596 diluted in PBS. For assessing immunogenicity of candidate vaccine cocktails, a total of 50
597 µL composed of 5 µg of each selected antigens were injected to the hind limb of a mouse.
598 All mice were boosted three weeks later. Serum samples were collected from

599 submandibular vein prior to immunization and two weeks after each injection. Mice were
500 euthanized at day 35 and spleen tissue was collected for analysis of cellular immunity.

501 **Pig study**

502 LNP-mRNA vaccination of pigs was performed according to the protocols approved by
503 Committee on Animal care (protocol number 2308000566) and Midwest Veterinary
504 Service (MVS), Inc. (protocol number 24005). A total of 38 four-week-old piglets were
505 randomly assigned into ten groups with four pigs for each of the 9 LNP-mRNAs groups
506 (MB-P72, MB-Penton, CD2v, P22, E199L, EP153R, P54, P30-IRES-MTE, and P30-P2A-
507 MTE) and the rest two pigs assigned to control group injected with sterile PBS. 30 ug of
508 LNP-mRNA expressing each individual antigen was diluted to 1 mL in sterile PBS and
509 injected to the back of ear intramuscularly and boosted three weeks later. Serum samples
510 were collected before vaccination and weekly after each injection. Body weight of each
511 pig was measured before the study and at the termination. All pigs were euthanized two
512 weeks after boost and spleen, draining dorsal superficial cervical (DSCLN) and
513 mandibular (MLN) lymph nodes were collected for testing T cell responses.

514 **Indirect ELISA**

515 Recombinant ASFV proteins with a HIS tag were inserted into PET28a vector and
516 expressed in a BL21 *Escherichia coli* system under induction of isopropyl- β -D-
517 thiogalactopyranoside (IPTG), followed by protein purification via Ni-nitrilotriacetic acid
518 (NTA) agarose. For detection of antibody response in serum from immunized animals,
519 300 ng purified proteins were coated onto 2HB ELISA plate (Thermo Fisher) in antigen
520 coating buffer (35 mM sodium bicarbonate and 15 mM sodium carbonate, pH 8.8) and
521 incubated at 37C overnight (41). Plates were washed with PBST (0.05% Tween 20 in 1 \times
522 phosphate-buffered saline) three times and blocked with 5% non-fat milk in PBST for 2
523 hours at room temperature. 100 μ L two-fold serial dilutions of serum starting from 1:50
524 were added to the coated plate and incubated for 2 hours at room temperature. HRP-
525 conjugated goat anti-mouse or pig IgG (H+L) (Thermo Fisher) was used as the secondary
526 antibody. Colorimetric reaction was developed by ABTS peroxidase substrate system and
527 stopped by ABTS stop solution in 30 min after color development. optical density OD₄₀₅
528 value was obtained to quantify the coloring intensity using the Tecan Infinite® 200 PRO
529 microplate reader. Cutoff values for each protein were determined by OD value of control
530 group animal samples plus three standard deviations. The endpoint titer for each animal
531 that has the highest dilution giving a reading above cutoff was calculated by interpolating
532 from a sigmoidal standard curve using GraphPad Prism.

533 **Enzyme-linked immunospot (ELISPOT) assay**

534 To evaluate T cell immunity, 10⁵ splenocytes from each mouse or pig were cultured in
535 CTL medium and seeded on pre-coated mouse IFN- γ /TNF- α Double-Color ELISPOT
536 plate (Cellular Technology Limited) or IFN- γ single-color ELISpot plate. Cells from each
537 animal were stimulated with 5 μ g/mL P72 peptide cocktail (**Table S2**), MTE peptide
538 cocktail, or purified proteins (CD2v, P22, EP153R, Penton, P54, P30). Cocktail of phorbol
539 12-myristate 13-acetate (PMA) and ionomycin (Thermo Scientific) was used as positive
540 control and cell culture medium as negative control. Plate was incubated at 37 °C with 5%
541 CO₂ for 36 hours followed by washing with distilled H₂O for three times before addition
542 of primary and secondary antibodies based on manufacture's protocol (Cellular
543 Technology Limited). Colored spots were counted by an automated immunospot analyzer
544 (Cellular Technology Limited). The antigen-specific IFN- γ and TNF- α spot-forming cells
545 (SFCs) were calculated and results were analyzed using GraphPad Prism.

546 **Hemadsorption inhibition assay**

547 CD2v has been reported to contribute to hemadsorption which is a term describing ASFV-
548 infected cells attracting red blood cells to attach on cell surface and form “rosette” pattern
549 (45). To establish an assay for measuring anti-CD2v antibody-mediated hemadsorption
550 inhibition, HEK 293T cells transfected with CD2v-expression vector were harvested and
551 mixed with 1:100 diluted serum pretreated with Receptor Destroying Enzyme (RDE) at 37
552 °C for 18h and then 56°C for 1h. After 1 hour incubation of serum with cell, 2% red blood
553 cells (RBC) were added and incubated at 37°C for 24 hours. Monolayer cells were gently
554 washed with PBS for five times and fixed, permeabilized, and blocked for antibody
555 staining. Rabbit anti-RBC polyclonal antibody (LSBio) and mouse anti-HA mAb (Thermo
556 Fisher) were added and incubated for 1 hour at 37°C. Alexa Fluor™ 488-conjugated goat
557 anti-mouse IgG and Alexa Fluor® 594 AffiniPure™ goat anti-rabbit IgG (H+L) (Jackson
558 ImmunoResearch) were used as secondary antibody. Cells were counterstained with DAPI
559 before proceeding with imaging by confocal microscopy (Nikon A1R HD25, Nikon).
560 Images were processed using the program ImageJ (<https://imagej.net/Fiji>) and geometric
561 mean fluorescent intensity of AF594 red fluorescence for RBC were calculated and
562 compared between groups.

563 **Antibody-dependent complement deposition (ADCD) assay**

564 Before testing, pig serum samples were inactivated at 56°C for 30 min. CHO-S cells stably
565 expressing each of the ASFV proteins (CD2v, EP153R, P22, P54, E199L and membrane-
566 bound P72, P30, Penton) were used as target cells and seeded in non-tissue culture treated
567 96-well plates at a density of 1.0×10^5 cells per well. 10-fold diluted inactivated serum
568 were added and mixed briefly with the cells. After incubation for 30 min at 37°C in a
569 humidified 5% CO₂ incubator, 50 µL control pig serum was added into each well as the
570 complement source and plates were incubated at 37C for 18 hours, followed by DAPI
571 staining. Cells were subject to flow cytometry using a BD LSR Fortessa HTS-2 cytometer.
572 Results were analyzed using FlowJo v10 software and the percentage of live CHO cells
573 positive for mCherry (%live) were calculated. The complement-mediated cytotoxicity was
574 calculated according to the formula: cytotoxicity (%) = $((1 - (\% \text{live of sample} / \% \text{live of}$
575 $\text{no serum ctrl})) \times 100$. The antigen-specific cytotoxicity effect was calculated by
576 normalizing to the control group serum.

577 **Antibody-dependent cellular cytotoxicity (ADCC) assay**

578 PBMCs collected from control pigs were used as a source of NK cells and pre-activated
579 with IL-2 (final concentration 20ng/mL) and IL-12 (final concentration 25 ng/mL) at 37°C
580 for 24 hours. CHO cells expressing ASFV proteins were seeded on 96-well plates at a
581 density of 2×10^4 cells per well. Inactivated serum was 1:10 diluted and incubated with
582 CHO cells for 30 min at 37°C. 50 µL of 2% Triton was added as positive control for
583 complete cell lysis. 4×10^5 Pre-activated PBMCs were added to achieve an effector to
584 target cell ratio of 20:1. Following 24 hours incubation at 37°C in humidified incubator, a
585 CytoTox-Glo™ luciferase based assay was used to measure the luminescence of viable
586 and non-viable cells according to manufacturer’s instructions. Percentage of cell lysis (%
587 lysis) was calculated by dead cell luminescence divided by total cell luminescence. The
588 percentage of antigen-specific cytotoxicity was calculated using the formular: (% lysis of
589 testing sample - % lysis of no serum ctrl) / (% lysis of triton - % lysis of no serum ctrl).

Antibody-dependent cellular phagocytosis (ADCP) assay

ASFV proteins were biotinylated using EZ-Link™ Micro Sulfo-NHS-LC-Biotinylation Kit and then conjugated to 1 μm red FluoSpheres™ NeutrAvidin™-Labeled Microspheres by overnight incubation at 4°C. Fluorescent microspheres were centrifuged at 13,000 g for 2 min at 4°C and washed twice with 1% BSA in PBS to remove the excess unbound proteins. The antigen-coated microspheres were resuspended in 1% BSA. Effector cells were porcine macrophage 3D4/31 cell line labeled with CellTrace™ CFSE Cell Proliferation Kit. To perform the phagocytosis assay, CFSE-labeled 3D4/31 cells were seeded on 96-well plate at a density of 1×10^4 cells per well and incubated for 24 hours in 5% CO₂ incubator. Cell culture media were removed and treated with fucoidan (final concentration 100 μg/mL) for 1 hour at 37°C to block scavenger receptors. Inactivated serum was 10-fold diluted and mixed with 10 μL suspension (equivalent to 4×10^6 antigen-coated microspheres). After 30 min incubation at 37 °C, the mixture was transferred to fucoidan-treated cells and incubated for 6 hours. Cells were washed extensively with sterile PBS, trypsinized, and resuspended in 200 μL medium containing DAPI. Cell suspension was subject to flow cytometry. Phagocytosed microspheres were gated as double positives for CFSE-labeled cells and fluorescent beads. Phagocytic score was determined by multiplying the percentage of fluorescent-bead-positive cells by the mean fluorescent intensity (MFI) of this population. The antigen-specific phagocytosis was computed by normalizing the phagocytic score to control group pigs.

Statistical analysis

Data was processed using R version 4.4.1 (61) with tidyverse version 2.0.0 (62). The R environment was executed using singularity version 3.5.0 (63) and this docker container `docker://bumproo/bulk_r441`. Four biological replicates for each antigen, except for P30 which had three replicates, were summarized by averaging. Correlation analysis was done using the `cor` function from the `stats` package in R with the `pearson` method and heatmaps were plotted using the `Heatmap` function from the R package `ComplexHeatmap` version 2.20.0 (64). These averaged values were then normalized by ranking antigen values for each immunological category from 1 (lowest) to 8 (highest). The `ggplot2` library in R is used to generate the Polar bar plot of rank scores of antigens based on the average rank scores. Based on the average Z-scores of antigens in five immunological categories, the `pyCirclize` module in `python3` is used to generate the chord diagram of correlations between antigens and five immunological categories. All possible 5-way, 4-way, and 3-way antigen combinations were then scored by summing the ranks. The sum scores calculated for the following sets of immunological data: ("IgG","IFN-γ","ADCD","ADCC","ADCP"), ("IgG","IFN-γ","ADCC","ADCD"), ("IgG","IFN-γ","ADCC"), and ("IgG","IFN-γ").

All the histogram charts were expressed as means +/- standard error of mean (SEM). Differences between groups were evaluated by one-way analysis of variance (ANOVA) using Prism software version 6 (GraphPad Software). A *P* value of <0.05 was regarded as statistically significant difference.

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359

360 **Author contributions:**

361 Conceptualization: FY, J Cui, J Chen
362 Methodology: FY, J Cui, TW, JQ, JHJ, HD, CAW
363 Investigation: FY, J Cui, J Chen
364 Supervision: J Chen, RX, HC
365 Writing—original draft: FY
366 Writing—review & editing: FY, J Chen

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368 **Competing interests:** F.Y., J. Cui, and J. Chen (inventors) declare that a provisional
369 patent application related to this work has been filed with the U.S. Patent and Trademark
370 Office in 2024. The other authors declare no competing interest.

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972 **Figures and Tables**

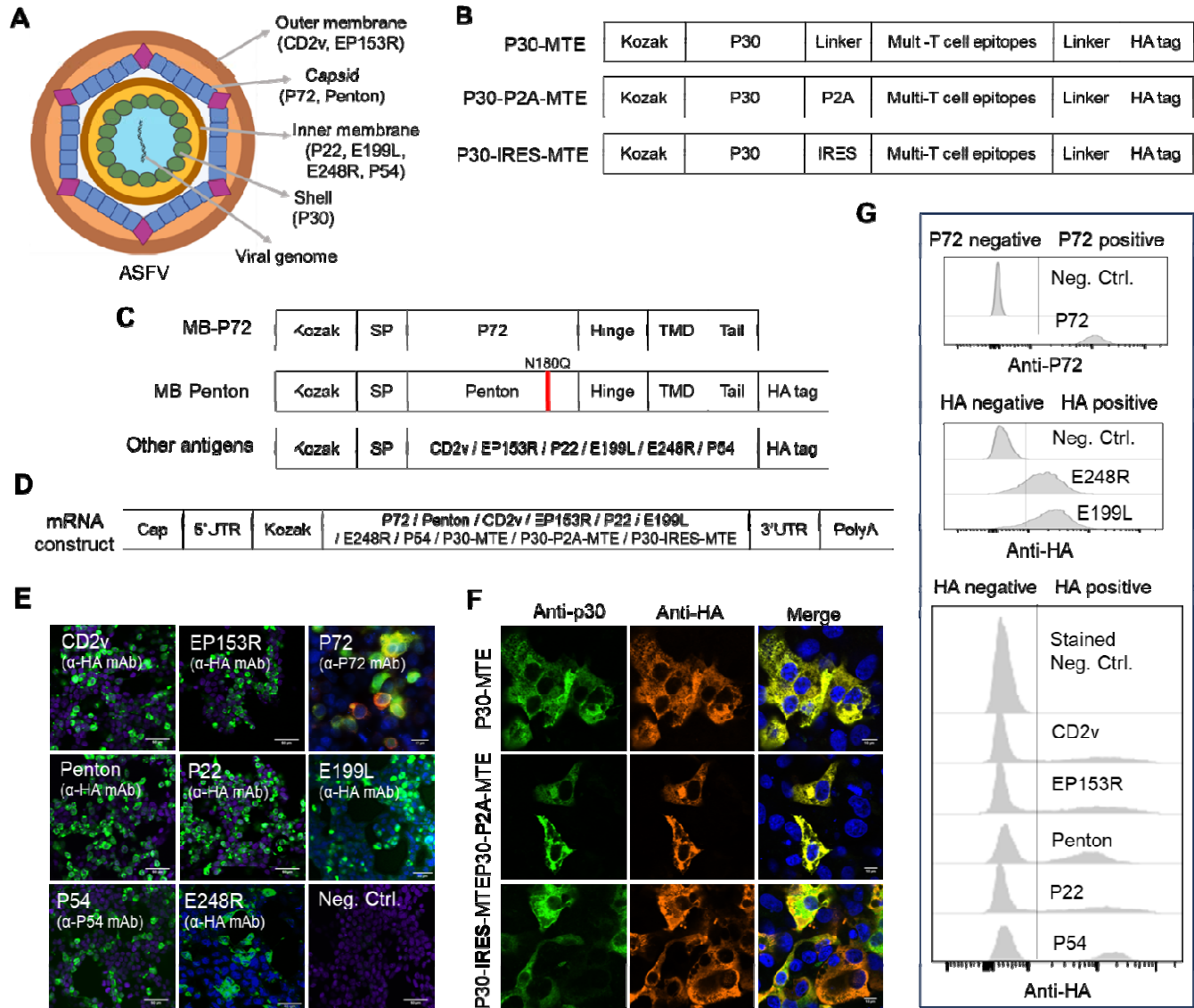


Fig. 1. Selection, design, and in vitro validation of ASFV vaccine candidate antigens.

(A) schematic diagram of the extracellular enveloped form of ASFV. Each layer of the viral particle is shown as distinctive colors, including the outer membrane, capsid layer, inner membrane, core shell, and viral DNA genome. Antigens selected in this study are depicted under each layer. (B) Schematic diagrams showing the three different designs for expressing T cell-directed antigens. In P30-MTE, P30 is fused with multi-T cell epitope (MTE) by a GGGS linker. In P30-P2A-MTE, P30 is linked with MTE via P2A self-cleavage site. In P30-IRES-MTE, MTE is translated separately through IRES. All three constructs contain an HA tag in the C terminus for detecting MTE expression. Kozak sequence was added in the N terminus of all constructs in aid of translation. (C) Schematic diagrams of DNA expression vectors for P72, Penton, CD2v, EP153R, P22, E199L, E248R, and P54. The membrane-anchoring of P72 and Penton was accomplished by addition of signal peptide (SP) in the N terminus, a hinge, transmembrane domain (TMD) and short cytoplasmic tail in the C terminus. The red vertical line represents N180Q mutation to remove the glycan on Penton. A HA tag was added in the C terminus

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of all other ASFV proteins except P72 for easy monitoring expression at the protein level. For P72 detection, a commercial monoclonal antibody was used. **(D)** Schematic diagrams of mRNA construct design. All designed ASFV genes were inserted into a pUC vector between the 5' untranslated region (5' UTR) and the 3'UTR, followed by polyA. A cap was added to the in vitro transcribed mRNA using Vaccinia Capping Enzyme. For testing immunogenicity in mice, all mRNA constructs, except P72, included a HA tag in the C terminus. The HA tag was removed for immunogenicity testing in pigs. **(E-F)** Confocal imaging of candidate antigens in HEK 293T cells transfected with DNA plasmids expressing CD2v, EP153R, P72, Pen.ton, P22, E199L, E248R, and P54 **(E)**, or P30-MTE, P30-P2A-MTE, and P30-IRES-MTE **(F)**. Scale bar is 50 μ m. **(G)** Flow cytometry analysis for expression of ASFV antigens following lipid nanoparticle (LNP)-mRNA transfection of HEK 293T cells. For confocal microscopy and flow cytometry, monolayer HEK 293T cells were transfected with DNA plasmids by polyethylenimine or LNP-mRNAs. Cells were harvested and fixed at 48 hours post transfection and stained with mouse mAbs specific for P72, P54, P30, and mouse anti-HA tag mAb for CD2v, EP153R, P22, E199L, E248R. A rabbit anti-HA tag polyclonal antibody was used to stain MTE antigens. AF488-conjugated goat anti-mouse IgG was used as secondary antibody and an additional AF594 conjugated goat-anti-rabbit IgG was used for **F**. Nuclei were stained with DAPI before confocal imaging.

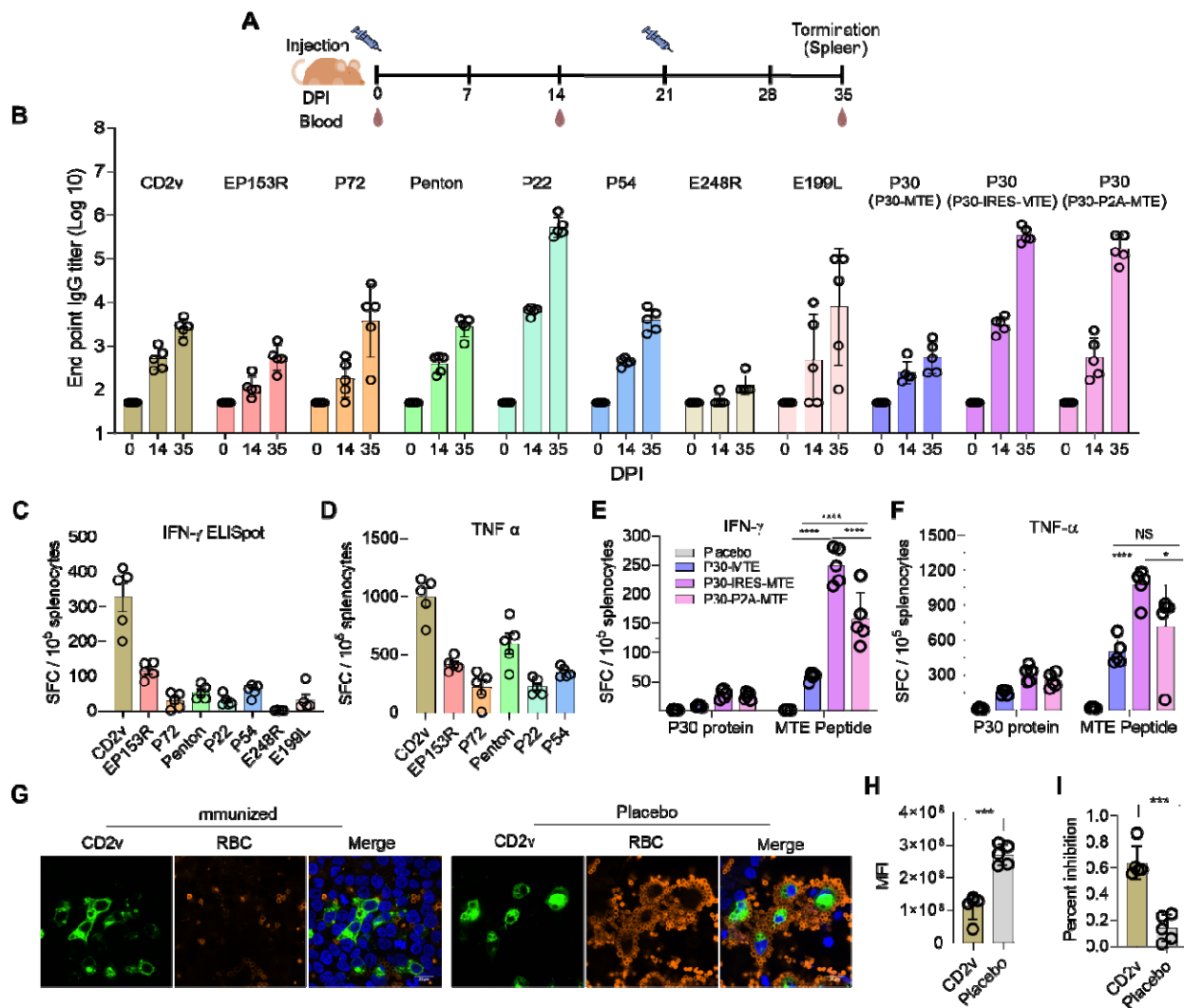


Fig. 2. Immunogenicity of candidate antigens in mice. (A) Experimental design. Mice (5 per group) were immunized with 5 μ g mRNA in LNP formulation at day 0 and day 21 or injected with PBS as control. Blood was collected before immunization and at 14 and 35 days post immunization (DPI) and used for assaying antigen-specific IgG levels by ELISA. Mice were euthanized at day 35 (14 days after boost) and spleen was collected for assaying antigen-specific T cell responses by ELISPOT. (B) Antigen-specific IgG responses normalized to day 0 (before immunization). For MTE groups, IgG titers specific for P30 were measured. The endpoint titer was defined as the reciprocal of the highest serum dilution that gives a reading above the cutoff which was determined by average OD₄₀₅ readings of placebo mice + 3 standard deviation. (C-D) Antigen-specific T cell responses to CD2v, EP153R, P72, Penton, P22, P54, E248R and E199L. 10⁵ splenocytes from each mouse were seeded on ELISpot plate and stimulated with 5 μ g/mL of purified proteins or P72 peptide. PMA/ionomycin cocktail and medium were included as positive and negative stimulation control, respectively. Dual-cytokine ELISpot was performed in 36 hours post stimulation. The number of IFN- γ (C) and TNF- α (D) spot-forming cells (SFC) per 10⁵ splenocytes was calculated and plotted. (E-F) Antigen-specific T cell responses to P30 (E) and MTE (F). ELISpot assay was done as in C-D except P30 protein and MTE peptide cocktail were used to stimulate splenocytes. (G-I) Hemoadsorption inhibition assay (HADIA) of CD2v antisera. Sera (n=5) from CD2v LNP-mRNA immunized mice were treated with

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receptor destroying enzyme (RDE) to remove unspecific factors inhibiting hemoadsorption, followed by incubation with CD2v-HA-expressing HEK 293T cells. 2% porcine RBC were added to the mixture and incubated for another 24 hours. Cells were fixed, permeabilized, and stained with a mouse anti-HA monoclonal antibody and rabbit anti-RBC polyclonal antibodies. AF594-labeled anti-rabbit and AF488-labeled anti-mouse IgG were added for secondary antibody staining. Cells were counterstained with DAPI before imaging using confocal microscope. Representative confocal images (**G**) and mean fluorescent intensity (**H**) and calculated percentage of inhibition (**I**). Each circle represents one mouse in B-F and H-I. One-way ANOVA for ELISpot assay and student's t-test for HADIA were used for statistical analysis. NS, no significance; *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.001$; ****, $P < 0.0001$.

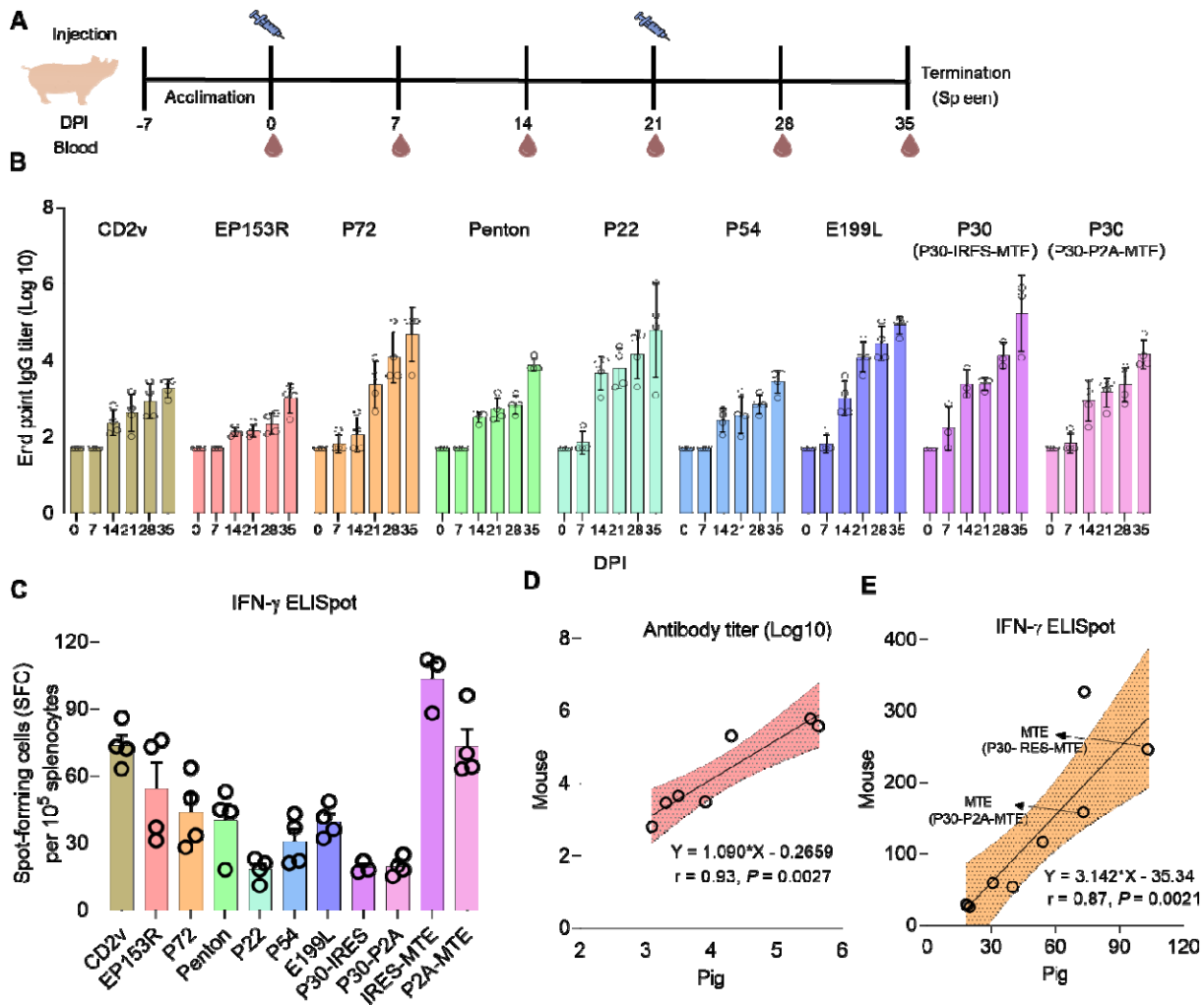


Fig. 3. Immunogenicity of candidate antigens in pigs. (A) Experimental design. Piglets (4 per group) were immunized with 30 μ g mRNA in LNP formulation at day 0 and day 21 or injected with PBS as control (2 pigs). Blood was collected before immunization and every 7 days post immunization (DPI) and used for assaying antigen-specific IgG levels by ELISA. Pigs were euthanized at day 35 (14 days after boost) and spleen was collected for assaying antigen-specific T cell responses by ELISPOT. (B) Antigen-specific IgG responses normalized to day 0 (before immunization). For MTE groups, IgG titers specific for P30 were measured. (C) Antigen-specific T cell responses. 10⁵ splenocytes from each pig were seeded on ELISpot plate and stimulated with 5 μ g/mL of purified proteins or peptides. PMA/ionomycin cocktail and medium were included as positive and negative stimulation control, respectively. IFN- γ ELISpot was performed in 36 hours post stimulation. The number of SFCs per 10⁵ splenocytes was calculated and shown. P30-IRES and P30-P2A refer to P30-specific IFN- γ ELISpot following P30-IRES-MTE and P30-P2A-MTE immunization, respectively. IRES-MTE and P2A-MTE refer to MTE-specific IFN- γ ELISpot following P30-IRES-MTE and P30-P2A-MTE immunization, respectively. (D-E) Pearson correlations for IgG responses (D) and IFN- γ response (E) evaluated in pigs (X-axis) and mice (Y-axis). Opened circles represent individual CD2v, EP153R, P72, Penton, P54, P22, E199L, P30 antigens. Arrows in (E) point to MTE. Equations for the linear regression and Pearson correlation coefficients along with P values are shown. Shading area shows the filled error bars.

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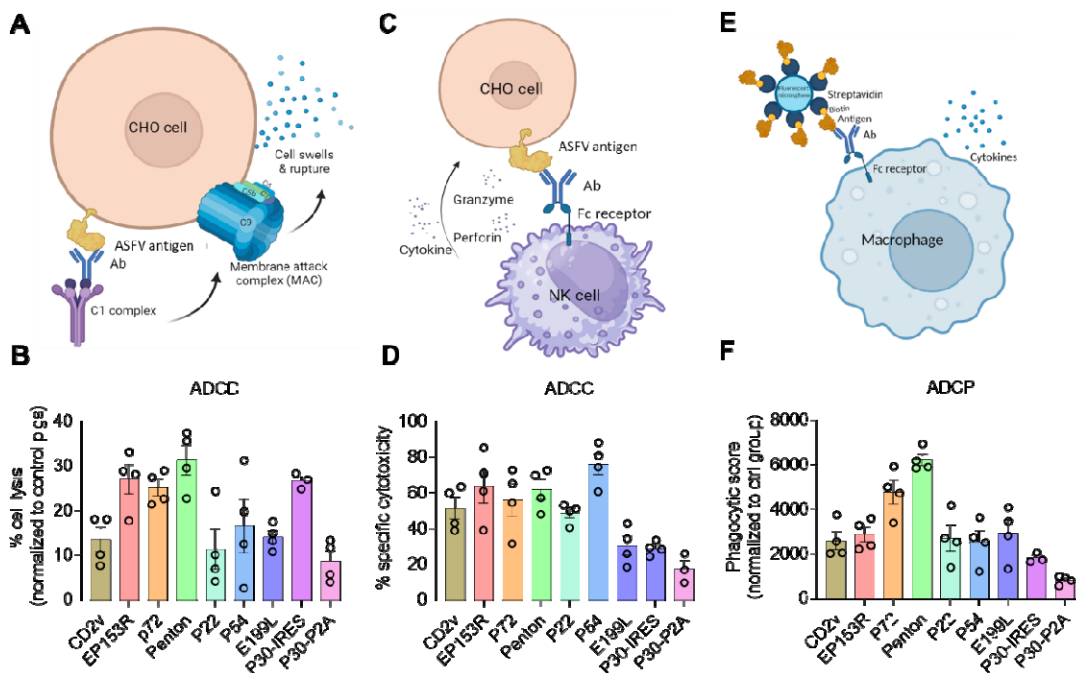


Fig. 4. Evaluation of antibody effector functions. (A) Schematic diagram of ADCD assay. CHO cells stably expressing each of the selected ASFV antigens were incubated with heat-inactivated serum from immunized pigs (antibody source), followed with serum from non-immunized pigs (complement source). Lysis of CHO cells were quantified by flow cytometry. (B) Comparison of percentages of lysis of CHO cells among sera from pigs immunized with the indicated ASFV antigens. (C) Schematic diagram of ADCC assay. CHO cells stably expressing each of the selected ASFV antigens were incubated with heat-inactivated serum from immunized pigs (antibody source), followed with incubation with total PBMC from non-immunized pigs (NK cell source). Lysis of CHO cells were quantified by a luminescent assay. (D) Comparison of percentages of lysis of CHO cells among sera from pigs immunized with the indicated ASFV antigens. (E) Schematic diagram of ADCP assay. Fluorescent beads conjugated with each of the selected ASFV antigens were incubated with heat-inactivated serum from immunized pigs (antibody source), followed with incubation 3D4/31 cells. Phagocytosis of beads by 3D4/31 was quantified by flow cytometry. (F) Comparison of phagocytic scores among sera from pigs immunized with the indicated ASFV antigens. P30-IRES and P30-P2A refer to P30-specific antibodies following P30-IRES-MTE and P30-P2A-MTE immunization, respectively. Each circle represents one pig.

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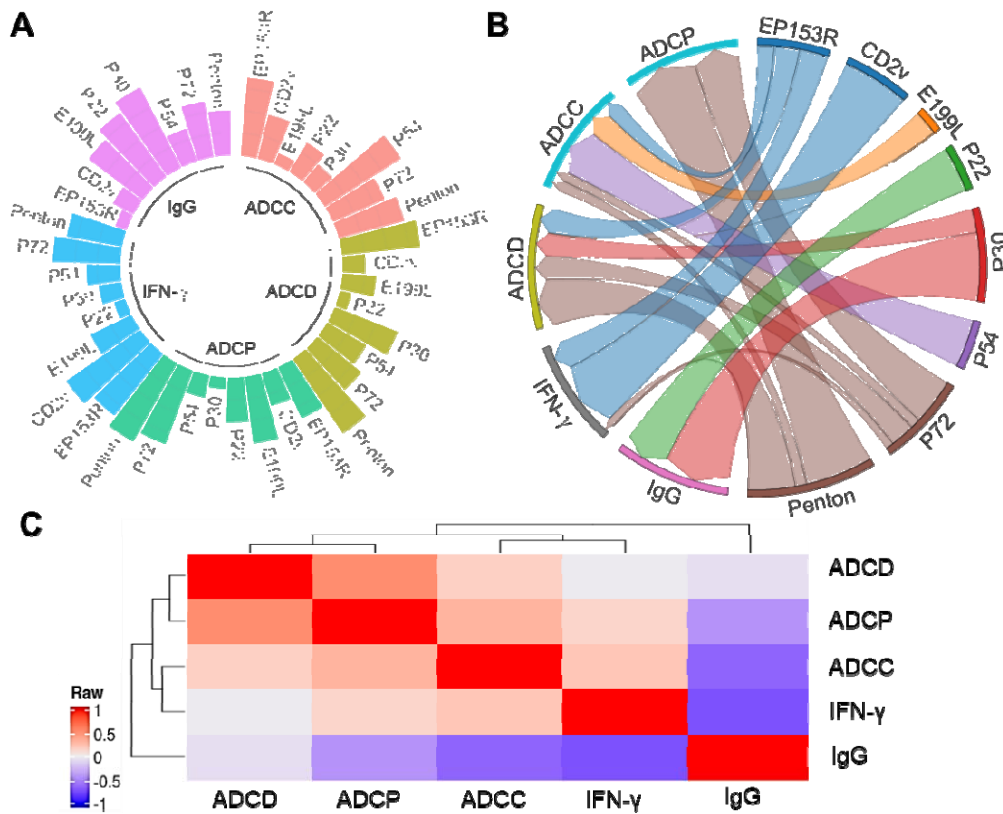
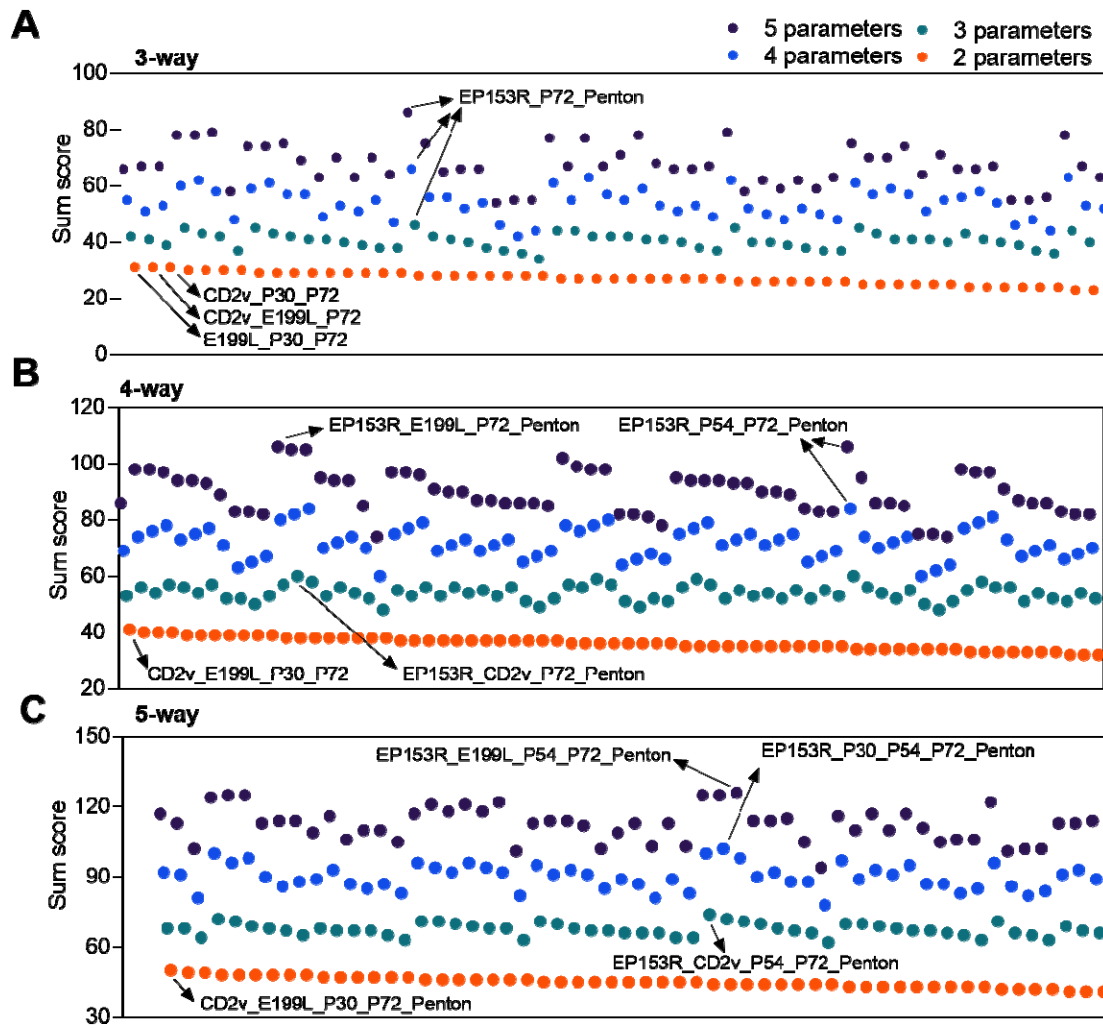
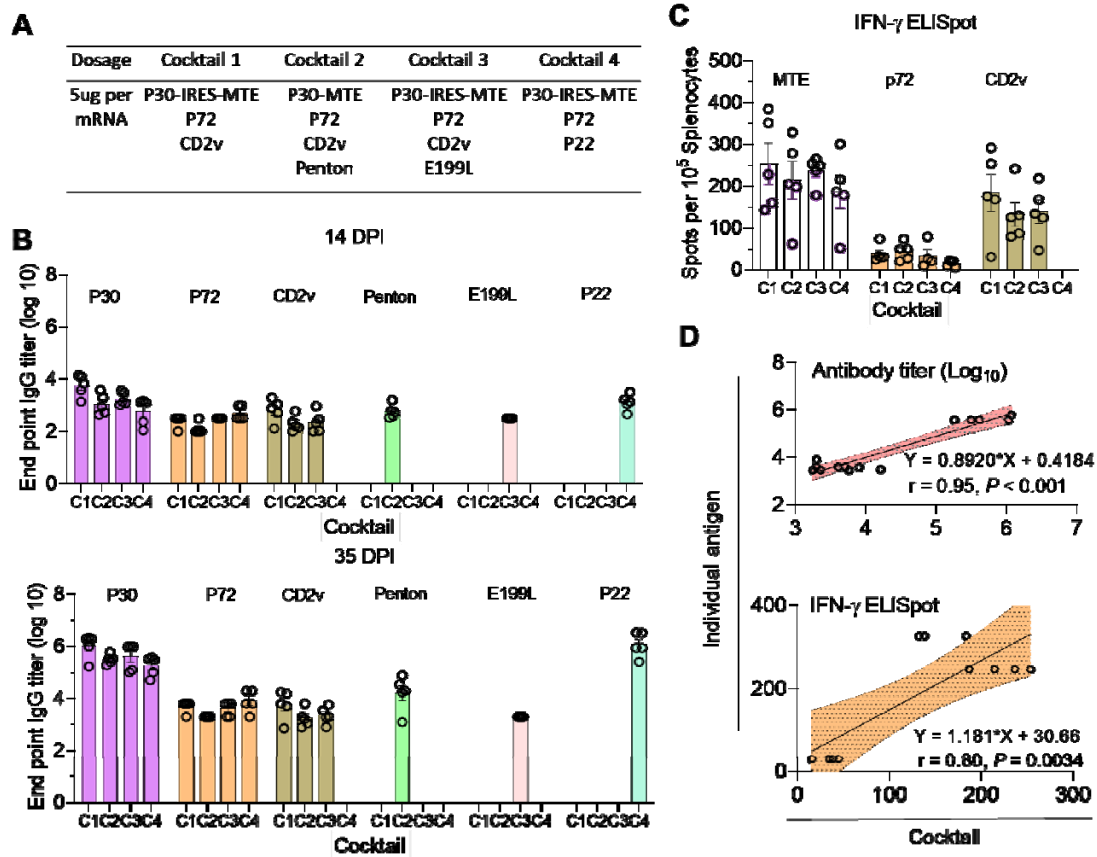


Fig.5. Computational analysis of immune profiles of specific antigens. (A) The polar bar plot of rank scores (1 to 8) of the indicated antigens in five immunological categories. The ggplot2 library in R is used to generate the Polar bar plot of rank scores of antigens. (B) The chord diagram of correlations between the indicated antigens and five immunological categories. Based on the average Z-scores of antigens in five immunological categories, the pyCirclize module in python3 is used to generate the chord diagram of correlations between antigens and five immunological categories, where the direction of connections is from antigens to immunological categories, and the width of the connections is proportional to the positive Z-scores. (C) Pearson correlation analysis of five immunological categories.

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103 **Fig. 6. Comparison of rank scores of antigen combinations for cocktail vaccines.** 3-
104 way (A), 4-way (B), and 5-way (C) antigen combinations based on rank sums of 5
105 immune parameters ("IgG", "IFN- γ ", "ADCD", "ADCC", "ADCP"), 4 parameters
106 ("IgG", "IFN- γ ", "ADCC", "ADCD"), 3 parameters ("IgG", "IFN- γ ", "ADCC"), and 2
107 parameters ("IgG", "IFN- γ ") colored in black, blue, green, and red, respectively.
108 Arrows point to the combinations with highest sum scores in different categories.



109 **Fig. 7. Induction of robust immune responses by candidate cocktail mRNA vaccines**
 110 **in mice.** (A) Composition of the four candidate cocktail vaccines. 5 μ g per mRNA
 111 was used to prepare LNP formulation. Four groups of mice (n=5 per group) were
 112 immunized at day 0 and 21. Blood samples were collected before immunization
 113 and 14 and 35 days after immunization for assaying antigen-specific IgG titers in
 114 the serum. Spleen was collected at day 35 after immunization for assaying IFN- γ
 115 secretion. (B) Comparison of antigen-specific IgG titers at day 14 and 35 after
 116 immunization for the indicated antigens. (C) Comparison of IFN- γ ELISpots in
 117 splenocytes following stimulation with MTE peptide cocktail, P72 peptides, and
 118 CD2v protein. (D) Pearson correlations for IgG (top) and T cell (bottom) responses
 119 induced by cocktail vaccination (X-axis) and individual antigen vaccination (Y-
 120 axis). Each circle represents one antigen: P30, P72, CD2v, Penton, E199L, P22
 121 (top), and MTE, P72, CD2v (bottom). Equations and correlation coefficients are
 122 shown. C1, C2, C3 and C4 represent cocktail 1, cocktail 2, cocktail 3 and cocktail
 123 4, respectively.

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Table 1. Comparison of vaccinia virus protective antigens with ASFV homologs

VACV ¹		ASFV		AA identity	Localization	Shared feature
Name	Size	Name	Size			
A33R	23-28 kDa	EP153R	18 kDa	18%	EEV ²	C-type lectin-like
B5R	42 kDa	CD2v	41 kDa	12%	EEV	Outer membrane
L1R	27 kDa	E248R	27 kDa	14%	IMV ³	Fusion protein
		E199L	24 kDa	8%		
A27L	14 kDa	P72	73 kDa	N/A [*]	IMV	Trimer
		Penton	28 kDa			Pentamer

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¹ Vaccinia virus (VACV)

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² Extracellular enveloped virus (EEV)

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³ Intracellular mature virion (IMV)

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* N/A: not applicable

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Table 2. T cell epitope sequences and their viral protein origins in MTE

Epitope Number	Protein name	Epitope Sequence	Reference
1	EP153R	KTLNLTKTYNHESNY	Burmakina, <i>et al.</i> , 2019; Oura, <i>et al.</i> , 2005; Fagbohun, <i>et al.</i> , 2022
2	EP153R	LTKTYNHESNYWVNY	
3	EP153R	YNHESNYWVNYSLIK	
4	EP153R	SNYWVNYSLIKNESV	
5	EP153R	GYKKQKHVSLL	
6	EP153R	KKQKHVSLLYICKS	
7	PP62	DFDPLVTFY	Herrera, <i>et al.</i> , 2021
8	PP62	GTDLYQSAM	
9	PP62	FINSTDFLYTAI	
10	PP62	LTDLVPTRL	
11	MGF100-1L	LQMAPGGSY	Bosch-Camós, <i>et al.</i> , 2021
12	MGF100-1L	LQMAPGGSYF	
13	MGF100-1L	QMAPGGSY	
14	MGF100-1L	QMAPGGSYF	
15	MGF100-1L	ITDNMTEEF	
16	A238L	DKDGNSALHYL	
17	K145R	AKIVEEGGEES	
18	MGF505-7R	NSTLVIRL	
19	P34	LTHGLRAEY	Zajac, <i>et al.</i> , 2022
20	P150	HIDKNIIQY	
21	P150	RVFSRLVIFY	
22	P37	KSMAAKIFI	
23	M448R	NTQPSHHVY	NetMHCpan-predicted MHC-I peptides
24	M448R	ALFPQYISY	
25	M448R	IIDHTTIQNY	
26	MGF505-7R	KLLEHVVKY	
27	MGF505-7R	LLDATLTRY	

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