

Electronic Supplementary Material

Porcine Immunoglobulin Fc Fused P30/P54 Protein of African Swine Fever Virus Displaying on Surface of *S. cerevisiae* Elicit Strong Antibody Production in Swine

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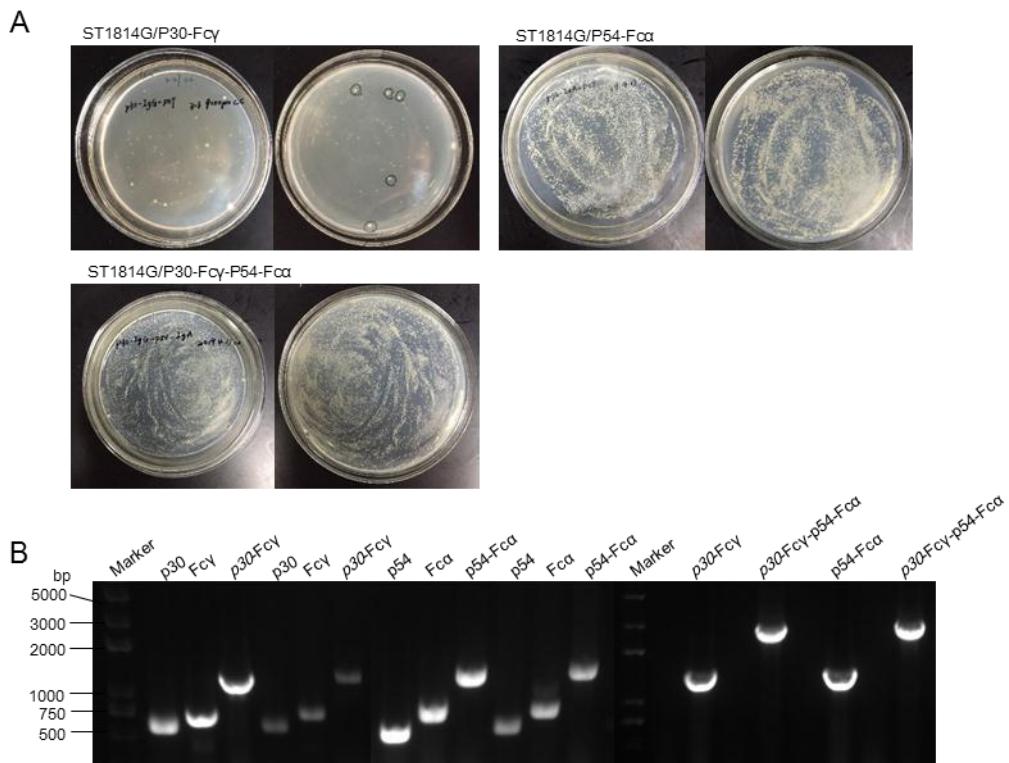


Fig. S1 Screening of the recombinant *S. cerevisiae* strains. **A** The *S. cerevisiae* transformants screened by SD-LEU auxotrophic medium. The p30-F γ and p54-F α expression plasmids first linearized with endonuclease *Bsa*I were transformed into *S. cerevisiae* strain ST1814G by LiAc transformation method in LEU-deficient plates to obtain transformant units. The larger diameter strains were selected and named ST1814G/P30-F γ , ST1814G/P54-F α , ST1814G/P30-F γ -P54-F α , which were verified as subsequent experimental materials. **B** A gel electrophoresis pattern of PCR identification of positive clones using genomic DNA of the three recombinant yeast strains. The genomic DNA of them was used as a template, and PCR amplification was carried out using specific primers. The length of p30-F γ , p54-F α and p30-F γ -p54-F α gene fragment was 1.2 kb, 1.3 kb, and 3 kb respectively, which was consistent with expectation. Lane 1, 4: p30; Lane 2, 5: F γ ; Lane 3, 6: p30-F γ ; Lane 7, 10: p54; Lane 8, 11: F α ; Lane 9, 12: p54-F α ; Lane 13: p30-F γ ; Lane 15: p54-F α ; Lane 14, 16: p30-F γ -p54-F α .

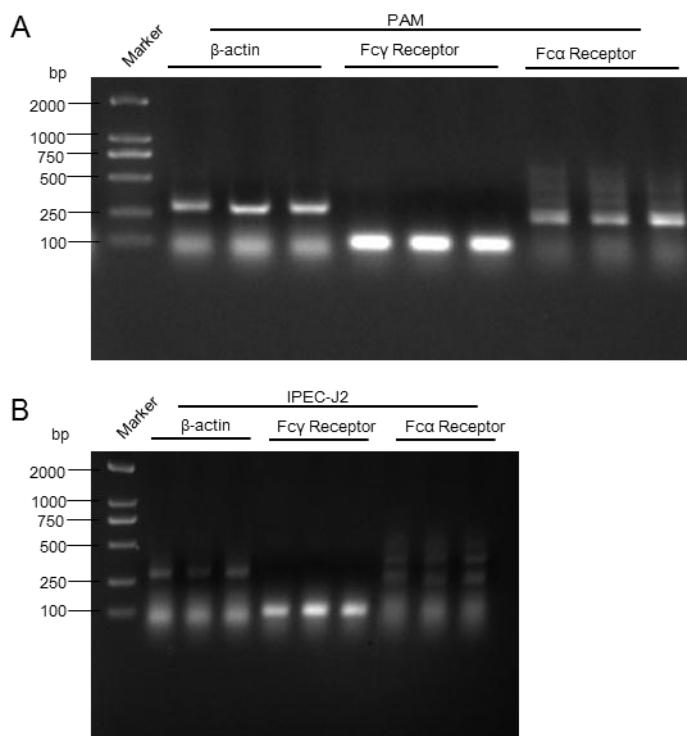


Fig. S2 Detection of Fc γ receptor and Fc α receptor of PAM and IPEC-J2 cells. **A** Detection of Fc γ receptor and Fc α receptor of PAM cells. **B** Detection of Fc γ receptor and Fc α receptor of IPEC-J2 cells.

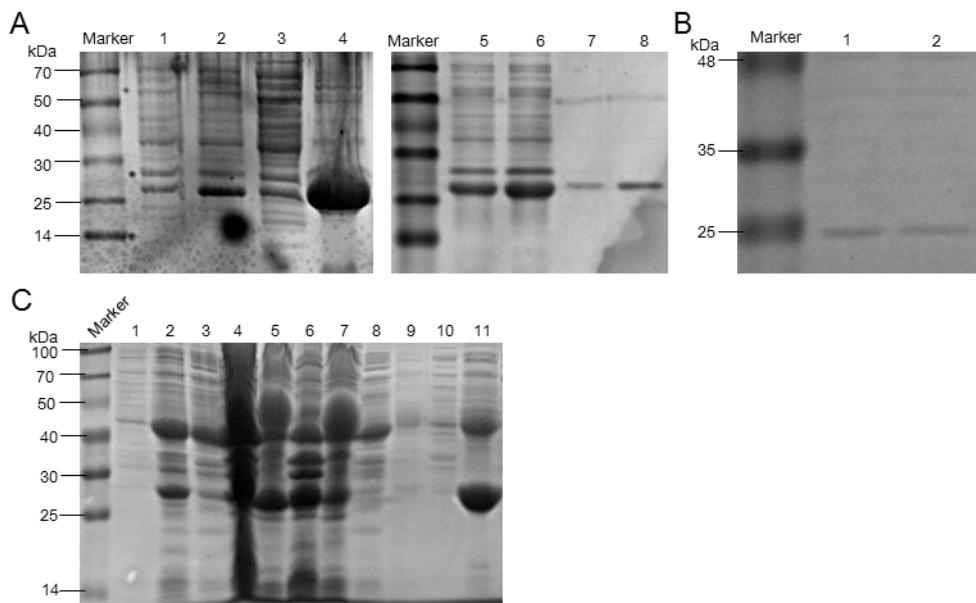


Fig. S3 Protein purification results of P30, P54, K145R. **A** Protein purification results of P30. The molecular weight of P30 is about 26 kDa. Lane Description: lane 1: Before induction; lane 2: After induction; lane 3: Supernatant after centrifugation; lane 4: Precipitation after centrifugation; lane 5: Centrifugal supernatant after precipitation of urea; lane 6: Centrifugal precipitation after precipitation of urea; lane 7: After inclusion bodies (200 mmol/L imidazole); lane 8: After inclusion bodies (250 mmol/L imidazole). **B** Protein purification results of P54. The molecular weight of P30 is about 25 kDa. Lane Description: Lane 1: After elution (200 mmol/L imidazole); Lane 2: After elution (250 mmol/L imidazole). **C** Protein purification results of K145R. The molecular weight of K145R is about 20 kDa. Lane Description: Lane 1: Before induction; lane 2: After induction; Lane 3: supernatant after centrifugation; Lane 4: Precipitation after centrifugation; Lane 5: Centrifugal supernatant after precipitation of urea; Lane 6: Centrifugal precipitation after precipitation of urea; Lane 7: After sedimentation; Lane 8: After hanging the column; Lane 9: Precipitation washing; Lane 10: Cleaning up; Lane 11: After inclusion bodies (250 mmol/L imidazole).

Table S1. Primers for overlapping PCR.

Primer name	Genbank number	Sequence of primer (5'-3')
IgG1-F	AK405786.1	ATGGAGTTCGGCTGAAC
IgG1-R		TTACCCCTGAGTCTGGAGATG
IgA1-F	AB194101.1	AGTAGCAGATGCCCTCTGCCTCC
IgA1-R		GTGTCCTGAAACCAGCCCCAAAAT
p30-Fcy-overlap-F ₁	MK757460.1	GACGATAAAGGTACCAAGGATCCATGAAAATGGAGGTACCTTCAAAAA
p30-Fcy-overlap-R ₁		GCATATGGGACATGTTGCATTTTTTTAAAGTTAATAACCATG
p30-Fcy-overlap-F ₂	MK757460.1	TCATGGTTATTAAACTTTAAAAAAAATGCAAACATGTCCCATATGC
p30-Fcy-overlap-R ₂		GAATTCCACCACACTGGATCCTTACCCCTGAGTCTGGAGATGG
p54-Fca-overlap-F ₁	MK276918.1	GACGATAAAGGTACCAAGGATCCATGGATTCTGAATTTCACCG
p54-Fca-overlap-R ₁		GAGAGGTGCTGCACTTGCACTAAGGAGTTCTAGGTCTTATGC
p54-Fca-overlap-R ₂	MK276918.1	GCATAAAGACCTAGAAAACCTTGATGCAAGTGCAGCACCTCTC
p54-Fca-overlap-R ₂		GAATTCCACCACACTGGATCCACCCGCCAGGCGGTC
pET-28a-p30-F	MK757460.1	CAGCAAATGGGTCGGGATCCATGAAAATGGAGGTACCTT
pET-28a-p30-R		ACGGAGCTCGAATTGGGATCCTTTTTAAAGTTAATAACCA
pET-28a-p54-F	MK276918.1	CAGCAAATGGGTCGGGATCCATGGATTCTGAATTTCACCC
pET-28a-p54-R		ACGGAGCTCGAATTGGGATCCCAAGGAGTTCTAGGTCTTATGCG
pET-28a-k145r-F	MK628478.1	CAGCAAATGGGTCGGGATCCATGGATCATTATCTTAAA
pET-28a-k145r-R		ACGGAGCTCGAATTGGGATCC GGATTCTCTCCTTC

Table S2. Primers for qRT-PCR.

Primer name	Genbank number	Sequence of primer (5'-3')
ACT1-F	L00026.1	ACTTTCAACGTTCCAGCCTTC
ACT1-R		CGTAAATTGGAACGACGACGTGAGTA
p30-F	MK757460.1	ATATTGTGAAATCTGCTCGT
p30-R		TCATGAATGTTCTCCGAAG
p54-F	MK276918.1	TTCTCATTGCTATCGTGGTCTTA
p54-R		AAGACCACGGATAGCAATGAGAAT
β-actin-F	DQ452569.1	GAATCCTGCGGCATCCACGA
β-actin-R		CTCGTCGTACTCCTGCTTGCT
Fcγ R-F	AK405786.1	AGCGAAGTCCTCCATGCCAGT
Fcγ R-R		CAGCCCCTTCCTCATCCTCCT
Fcα R-F	AB194101.1	TCCTACCATACTGCCACACC
Fcα R-R		TCATACAAGCCTGTCACCAC