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# Development of a Rapid and Sensitive Method for Detection of African Swine Fever Virus Using Loop-Mediated Isothermal Amplification

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# ABSTRACT

A loop-mediated isothermal amplification (LAMP) assay was developed for rapid, sensitive and specific detection of African swine fever virus (ASFV). A set of LAMP primers was designed based on the sequence of the ASFV gene K205R. Reaction temperature and time were optimized to 64 °C and 60 min, respectively. LAMP products were detected by agarose gel electrophoresis or visually with the addition of fluorescent dye. The detection limit of the LAMP assay was approximately 6 copies of the target gene per microliter, 100 times more sensitive than conventional PCR. LAMP is a simple and inexpensive molecular assay format for ASFV detection. To date, African swine fever has not been reported in China. LAMP can be used to monitor ASFV spread into China, thereby reducing the threat of ASF.

Keywords: African swine fever virus; LAMP; K205R gene; molecular biology.

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#### **INTRODUCTION**

African swine fever virus (ASFV) is an enveloped, complex, icosahedral virus with a double-stranded DNA genome of 170 to 190 kb in length (<u>Takamatsu et al., 2013</u>). It has been classified as the sole member of the new family Asfarviridae(<u>Sanchez et al., 2013</u>). This virus infects warthogs and bush pigs through bites from the argasid tick vector, as well as domesticated pigs and wild boar. It causes acute hemorrhagic fever and leads to high morbidity and mortality with consequent economic losses. There is no available vaccine for ASF (<u>Karalyan et al., 2012</u>).

Montgomery was the first to describe ASF, found in Kenya in 1921 (Gil et al., 2008). The virus spread from infected warthogs to domesticated pigs, causing disease with 100% mortality(Jezewska et al., 2011). In 1957, ASFV was reported in Lisbon, Portugal(Wieland et al., 2011). Following these reports, the disease was found on the Iberian Peninsula, and sporadic outbreaks occurred in France, Belgium and other European countries in the 1980s (Sampoli Benitez et al., 2013; Uttenthal et al., 2013). In the late 1970s, ASFV crossed the Atlantic Ocean and outbreaks were reported on some Caribbean islands, including Cuba and the Dominican Republic (de Glanville et al., 2014). The most recent outbreak of ASFV outside of Africa was in Georgia, at the beginning of 2007, and it has since spread to Azerbaijan and Russia, neighboring Armenia(Diaz et al., 2012; Uttenthal et al., 2013). So far, there have been no reports of ASF in China, but due to its geographical proximity, a plan of defense is urgently needed, making a rapid and sensitive detection method for ASFV of critical importance.

A variety of laboratory assays can be used to detect ASFV, including PCR (Howey et al., 2013) and real-time PCR (Oura, 2013), two assays that have been adopted for routine diagnosis. Although PCR is a highly sensitive method for the detection of ASFV, it relies on precision thermocycling which requires expensive instrumentation. Advances in molecular biology have seen the establishment of the loop-mediated isothermal amplification (LAMP) assay, originally described by Notomi et al (Notomi et al., 2000). LAMP allows rapid

amplification of target DNA sequences in a highly specific manner under isothermal conditions. In this study, with the DNA polymerase *Bst* and specific internal and external primers enabled the identification of six independent regions of the target gene, we described the development of a rapid LAMP assay for the detection of ASFV, and compare this assay to PCR tests.

#### MATERIALS AND METHODS

#### **Preparation of DNA template and samples**

The *K205R* gene of ASFV was synthesized by Invitrogen in Shanghai, China (GenBank accession no. NC\_001659.1). The recombinant plasmid pMD18T-K205R was kept at -70 °C until use as the template in ASFV LAMP and conventional PCR assays.

#### Primer design

LAMP primers were designed based on ASFV *K205R*. Two sets of primers recognized six different regions of the target gene—inner primers FIP and BIP, and outer primers F3 and B3. FIP consists of F1c, which is complementary to the F1 and F2 sequences. BIP consists of B1c, which is complementary to the B1 and F2 sequences (Fig. 1). A pair of specific PCR primers was also designed. The LAMP and PCR primer sequences are shown in Table 1.

	F3	F2
5'	GCCATTATCGCCCAA	CTTGAGATTCTGATGATAAATGGCACTCCACT 3'
5'	CGGTAATAGCGGGTT	GAACTCTAAGACTACTATTTACCGTGAGGTGA 3'
5'	TC CGGCAAAAAAAA	CAACGATTAAGGAGGCTATGCCCCCTACCTTAC 3'
5'	AGGC CGTTTTTT	GTTGCTAATTCCTCCGATACGGGGGATGGAAGT 3' F1c
	B1c	
5'	TCAAACACGAACAAT	GATCAAACGAGTCCTCCCGCCTCAGGCAAAA 3'
5'	AGTTTGTGCTTGTTA	CTAGTTTGCTCAGGAGGGGGGGAGTCCGTTTT 3'
5'	CAAGTGAAACACCTA	AAAAAAATCCCACGAATGCAATGTTCTTCACG 3'
5'	GTTCACTTTGTGGAT	TTTTTTTAGGGTGCTTACGTTACAAGAAGTGC 3'
	B2	B3
Fig	ure 1: Location	n and sequences of ASFV K205R
use	d to design th	ne inner and outer primers. The

sequences of the primer sites are underlined.

Table 1: Oligonucleotide primer sets used for	LAMP
and conventional PCR	

Prim er nam e	Positio n <sup>a</sup>	Sequence $(5' \rightarrow 3')$
FIP BIP F3 B3 PCR -F PCR -R <sup>a</sup> Gen00	F1:463 93- 46416 F2:463 53- 46375 B1:464 17- 46439 B2:464 73- 46497 46334- 46351 46500- 46519 46355- 46355- 46356 46635- 46656	GCATAGCCTCCTTAATCG TTGTT- AGATTCTGATGATAAATG GCACT CCTACCTTCATCAAACAC GAACA- GGATTTTTTTTAGGTGTTT CACTTG GCCATTATCGCCCAACTT CGTGAAGAACATTGCATT CG ATTCTGATGATGATAAATGGC ACTC GTTCTCCACATGTAAAGA CCCT

"Genome position accordingtoASFV strain BA71V (GenBankID:NC\_001659.1).

#### **Protocols for LAMP and PCR assays**

The LAMP assayreaction volume was  $25\mu$ l, consisting of  $2.5\mu$ l 10× ThermoPolreaction buffer (20 mM Tris–HCl, 10 mMKCl, 10 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>, and 0.1% w/v Triton X-100; New EnglandBiolabs Inc.,Beverly, MA, USA), 8 U of *Bst*DNA polymerase (largefragment; New EnglandBiolabs), 0.2  $\mu$ Mofeachouter primer and 2  $\mu$ Mofeachinner primer, 1 M betaine(Sigma-Aldrich, St. Louis, MO, USA), 0.6 mMdNTPs (2.5mMeach; TIANGEN Biotech), and4 mM MgCl<sub>2</sub> (25mM; TIANGEN Biotech); 1  $\mu$ IASFV plasmidwasthenadded. The reactionwasrun in a gradientPCR

thermocycler(T100<sup>TM</sup>ThermalCycler, Bio-Rad. Hercules, CA, USA). LAMP assayreactiontemperatures of 60, 61, 62, 63, 64and 65 °C, and reaction times of 30, 40, 50, and 60 minweretested. The amplificationproductswereanalyzedbyelectropho resison 1.5% agarose for а gel optimalreactionconditions.

The PCR wascarried out in a 25-µlreaction volume containing 0.2 µMeachofprimers PCR-F

and PCR-R, 1 µl DNA templateand 12.5 µl  $2 \times$ Taq PCR Mastermix(TIANGEN Biotech), consistingofTaq DNA enzyme,dNTPs, MgCl<sub>2</sub>, andthereaction buffer. The reactionswererun in a gradient PCR thermocycler (T100<sup>TM</sup>ThermalCycler) andtemperaturesof 56, 57, 58, 59, 60 and 61 °Cweretested. The amplificationproductswereanalyzedbyelectropho resison a 1.5% agarose gel to determine optimalconditions.

#### Analysis of LAMP products

additiontoelectrophoreticanalysis, LAMP In productswerealsodetectedvisually. LAMP productswereidentifiable as а largeamountofwhiteprecipitateformedduringthe LAMP reaction. In addition, 100-fold dilutedSYBR Green Istain (BioTekeBiotechCo., Ltd., Beijing, China) wasaddedtothereaction andthesolutionwasobserved tube for color changeunderultraviolet light. Green fluorescencewasobserved in positive reactions.

#### Sensitivity of LAMP and PCR assays

The ASFV plasmidsweresubjected serial 10fold dilutions, to a final concentration of  $6 \times 10^{0}$  to  $6 \times 10^{4}$  copies/µl. The detection limit of the LAMP assay was compared with that of the PCR assay using identical concentrations of the same DNA templates. The optimized methods were used to detect the sensitivit y of LAMP and PCR, with a reaction volume of 25 µl.

#### RESULTS

#### **Optimization of reaction conditions**

The LAMP productsappearedontheagarose gel in a ladder-likepatternatdifferenttemperatures (60, 61, 62, 63, 64 and 65 °C) and different times (30, 40, 50, and 60 min). Little differencewasfound in theelectrophoreticpatternsat 60 and 65 °C (Fig. 2a), and the amount of amplification product was large sta

fter 60 min reaction (Fig. 2b). Thus, 64 °C for 60 min wasselected as theoptimalreactioncondition for ASFV LAMP assay, andthecomponentsandaddedvolumes for the LAMP reaction are given in Table 2. Agarose gel electrophoresis of the ASFV PCR products °C indicated 59 as theoptimalreactiontemperature for thisassay (Fig. 2c).



PCR Figure 2:Optimizationof and ASFV detection. The LAMP LAMP for andtheproductswereexaminedbyagarose reactionwasrununderdifferenttemperaturesand times, gel electrophoresis. (a)Lanes 1-6: LAMP wascarried out at 60, 61, 62, 63, 64 and 65 °C, respectively. (b) Lanes 1-4: LAMP wascarried out for 30, 40, 50 and 60 min, respectively. (c) Lane 1: negative control; lanes 2-7: PCR wascarried out at 56, 57, 58, 59, 60 and 61 °C, respectively. Lane M: DNA Marker (DL 2000).

Table 2: LAMP reaction conditions

Components	Volume (µl)
Bst DNA polymerase (8 U)	1
ThermoPol buffer ( $10\times$ )	2.5
dNTPs (2.5 mM each)	6
MgCl <sub>2</sub> (25 mM)	4
Betaine (1 M)	2.5
Outer primers (5 µM each)	1
Inner primers (50 µM each)	1
Template	1
Deionized water	4
Total	25

Visual detection of ASFV LAMP

As alreadynoted, agarose gel electrophoresisofthe LAMP productsshown a ladder-likepatternwithfragmentsofdifferentsizes (Fig. 3a). In addition, the ASFV LAMP productswerevisuallydetectedbytheirturbidity in thereaction tubes. In ASFV-positive samples, turbiditycouldbeseenwiththenakedeyeunder natural light (Fig. 3b), whereastherewas no turbiditywiththe negative samples. Uponadditionof 1 µl SYBR Green I (100-fold dilution), greenfluorescencecouldbeobserved in the positive tubes underultraviolet light, and no fluorescencewasseen in the negative ones (Fig. 3c).



**Figure 3:**Analysis of LAMP products. (a) LAMP products detected by electrophoretic analysis. (b) Visual observation under natural light. (c) Visual observation following the addition of SYBR Green I under ultraviolet light. Lane M: DNA marker (DL 2000); 1: negative control reaction; 2: positive reaction.

# Detection limits of LAMP and conventional PCR

LAMP and conventional PCR were carried out and their respective detection limits compared using 10-fold serial dilutions ( $6 \times 10^{0}$  to  $6 \times 10^{4}$  copies/µl) of ASFV recombinant plasmid as the template. We could detect 6 copies/ $\mu$ l DNA by LAMP (Fig. 4a), whereas the minimum detectable amount for PCR was  $6 \times 10^2$  copies/ $\mu$ l (Fig. 4b). Thus, the LAMP assay was approximately 100 times more sensitive than the PCR assay.



**Figure 4:**Sensitivities of LAMP (a) and PCR (b) assays for ASFV detection, using 10-fold serial dilutions of the same template. Lane M: DNA Marker (DL 2000); lane 1: negative control reaction; lanes 2–6:  $6 \times 10^4$ ,  $6 \times 10^3$ ,  $6 \times 10^2$ ,  $6 \times 10^1$ , and  $6 \times 10^0$  copies/µl, respectively.

## DISCUSSION

To date, therehasbeen no recordedoccurrenceof ASF in China, but in the context of globalization, with the development of import, export trade and frequent movement of personnel, ASFV wouldbea threat for China, dealing а devastatingblowtopig-breedingindustryof China. There is no efficient treatment measure or vaccine for thecontrolof

ASFnow.themaincontrolmeasureremainsquarantin e. A simple, rapid, and sensitive assay for thedetection of ASFV is essential. In this study, a novel LAMP diagnosticprotocolwasestablishedtodetectthisvirus. Therehasbeen quite a bit ofresearchon P72, P73andP54of ASFV. However. there are ASFV K205R. K205Ris fewstudieson а conservedandspecific ASFV gene thatappears in earlyinfectionandismostrapidlyamplified(Cobbold et al., 2001; Gutierrez-Castaneda et al., 2008). In addition, PK205R canbeexpressed when ASFV just invades. Thesemake it a powerfulindicator for thedetection of ASFV. In ourstudy, the target gene wasamplified wellonly in 60 min, but it needs at least 2–3 h for conventional PCR.

LAMP wasverysensitive for ASFV detection, approximately 100-fold more sensitivethanconventional PCR usedwiththesame DNA template. The minimumdetectableamountof LAMP copies/µl, was 6 andthusthisassaycandetectthevirus in earlyasymptomaticinfectedsamples. The high sensitivitvof LAMP can serve for diseasesurveillance, andmakes it suitable for use in theChineseEntry-ExitInspectionandQuarantine

Bureau toprevent ASFV spread into China. In addition, LAMP productsappear as whiteprecipitatestothenakedeye, as explainedby Mori et al. (Mori et al., 2001). Visual detectionofLAMP

productscanalsobeperformedusing a fluorescentchromogenwithnucleicacidstain(<u>Cardos</u> <u>o et al., 2010; Suebsing et al., 2013</u>). Therefore, thenovel LAMP assay for ASFV describedhereinis more practical for routine ASFV detectionthanconventional PCR.

In conclusion, a rapid and sensitive LAMP assay for the detection of ASFV was developed and validated. The LAMP reaction took approximately 1 h in a PCR thermocycler, or in a water bath at 64 °C. The detection limit of the ASFV LAMP assay was as low as 6 copies per µl DNA, 100 times more sensitive than that of conventional PCR. LAMP products could be detected by agarose gel electrophoresis, but also visually. The LAMP assay is a rapid, simple, specific, sensitive and reliable diagnostic protocol that can be applied under field conditions, even at China's trade borders, to prevent ASFV spread into China. This technique for ASFV detection holds great potential, pending further refinement.

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# REFERENCES

- 1. Cardoso TC, Ferrari HF, Bregano LC, Silva-Frade C, Rosa AC, Andrade AL. Visual detection of turkey coronavirus RNA in tissues and feces by reverse-transcription loop-mediated isothermal amplification (RT-LAMP) with hydroxynaphthol blue dye. Molecular and cellular probes 2010; 24: 415-417.
- 2. Cobbold C, Windsor M, Wileman T. A virally encoded chaperone specialized for folding of the major capsid protein of African swine fever virus. Journal of virology 2001; 75: 7221-7229.
- 3. de Glanville WA, Vial L, Costard S, Wieland B, Pfeiffer DU. Spatial multi-criteria decision analysis to predict suitability for African swine fever endemicity in Africa. BMC veterinary research 2014; 10: 9.
- 4. Diaz AV, Netherton CL, Dixon LK, Wilson AJ. African swine fever virus strain Georgia 2007/1 in Ornithodoros erraticus ticks. Emerging infectious diseases 2012; 18: 1026-1028.
- Gil S, Sepulveda N, Albina E, Leitao A, Martins C. The low-virulent African swine fever virus (ASFV/NH/P68) induces enhanced expression and production of relevant regulatory cytokines (IFNalpha, TNFalpha and IL12p40) on porcine macrophages in comparison to the highly virulent ASFV/L60. Archives of virology 2008; 153: 1845-1854.
- 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. Archives of virology 2008; 153: 2303-2306.

- Howey EB, O'Donnell V, de Carvalho Ferreira HC, Borca MV, Arzt J. Pathogenesis of highly virulent African swine fever virus in domestic pigs exposed via intraoropharyngeal, intranasopharyngeal, and intramuscular inoculation, and by direct contact with infected pigs. Virus research 2013; 178: 328-339.
- Jezewska MJ, Szymanski MR, Bujalowski W. Interactions of the DNA polymerase X from African Swine Fever Virus with the ssDNA. Properties of the total DNA-binding site and the strong DNAbinding subsite. Biophysical chemistry 2011; 158: 26-37.
- Karalyan Z, Zakaryan H, Arzumanyan H, Sargsyan K, Voskanyan H, Hakobyan L, Abroyan L, Avetisyan A, Karalova E. Pathology of porcine peripheral white blood cells during infection with African swine fever virus. BMC veterinary research 2012; 8: 18.
- 10. Mori Y, Nagamine K, Tomita N, Notomi T. Detection of loop-mediated isothermal amplification reaction by turbidity derived from magnesium pyrophosphate formation. Biochemical and biophysical research communications 2001; 289: 150-154.
- Notomi T, Okayama H, Masubuchi H, Yonekawa T, Watanabe K, Amino N, Hase T. Loop-mediated isothermal amplification of DNA. Nucleic acids research 2000; 28: E63.
- Oura C. African swine fever virus: on the move and dangerous. The Veterinary record 2013; 173: 243-245.
- Sampoli Benitez B, Barbati ZR, Arora K, Bogdanovic J, Schlick T. How DNA polymerase X preferentially accommodates incoming dATP opposite 8-oxoguanine on the template. Biophysical journal 2013; 105: 2559-2568.

- 14. Sanchez EG, Quintas A, Nogal M, Castello A, Revilla Y. African swine fever virus controls the host transcription and cellular machinery of protein synthesis. Virus research 2013; 173: 58-75.
- 15. Suebsing R, Prombun P, Kiatpathomchai W. Reverse transcription loop-mediated isothermal amplification (RT-LAMP) combined with colorimetric gold nanoparticle (AuNP) probe assay for visual detection of Penaeus vannamei nodavirus (PvNV). Letters in applied microbiology 2013; 56: 428-435.
- Takamatsu HH, Denyer MS, Lacasta A, Stirling CM, Argilaguet JM, Netherton CL, Oura CA, Martins C, Rodriguez F. Cellular immunity in ASFV responses. Virus research 2013; 173: 110-121.
- Uttenthal A, Braae UC, Ngowi HA, Rasmussen TB, Nielsen J, Johansen MV. ASFV in Tanzania: asymptomatic pigs harbor virus of molecular similarity to Georgia 2007. Veterinary microbiology 2013; 165: 173-176.
- 18. Wieland B, Dhollander S, Salman M, Koenen F. Qualitative risk assessment in a data-scarce environment: a model to assess the impact of control measures on spread of African Swine Fever. Preventive veterinary medicine 2011;99: 4-14.

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