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A new rapid and sensitive method for detecting chicken infectious anemia virus

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Since the chicken infectious anemia virus (CIAV) was discovered in 1979, which has been reported as an economically significant and immunosuppressive poultry disease in the world. A novel clinical detection method for the prevention and control of CIAV in the poultry sector is urgently needed. Here, we established a real-time recombinase-aided amplification assay (RAA) for CIAV on-site with a rapid, highly sensitive, strongly specific, low-cost, and simple operational molecular diagnosis detection method. The primers and probe were developed using the CIAV VP2 gene sequence, which has a 117-bp specific band. This assay, which could be carried out at 41°C and completed in 30min without cross-reactivity with other viruses, had the lowest detection limit of 10 copies of CIAV DNA molecules per reaction. Furthermore, the kappa value of this assay was 0.947, the sensitivity was 93.33%, and the specificity was 100% when compared to the real-time quantitative polymerase chain reaction assay (real-time qPCR). These results indicate that using a real-time RAA assay to detect CIAV on-site could be beneficial. In the future, the real-time RAA test may be a regular assay for the prevention and control of CIAV, as well as help the reduction of economic losses in the poultry business.

KEYWORDS

chicken infectious anemia virus, VP2, real-time recombinase-aided amplification assay, detection, real-time qPCR

Introduction

Chicken infectious anemia virus (CIAV) is an immunosuppressive virus, and its infection could cause aggravation of co-infections. CIAV is the only member of the genus gyrovirus of the family Anelloviridae with 20-dihedral symmetry and no envelope (Goryo et al., 1985; Natesan et al., 2006), and almost all of its ssDNA nucleic acids are 2,298 nt in

length (Ducatez et al., 2006). CIAV contains three overlapped ORFs, encoding three distinct viral proteins: major capsid and immunogenic protein (VP1, 51.6 kDa), non-structural scaffold protein (VP2, 24.0 kDa), and apoptin (VP3, 13.6 kDa) (Noteborn et al., 1991; Shao et al., 2021). Chickens are the host of CIAV, and all ages are susceptible, mainly affecting chickens of 2–3 weeks. CIAV invades bone marrow hematopoietic tissue and thymus, leading to bone marrow chlorosis and thymus atrophy, and clinical symptoms are aplastic anemia and systemic lymphoid tissue atrophy (Yuasa et al., 1976). CIAV can be transmitted vertically or horizontally, resulting in widely spread around the world, which is one of the major diseases in the poultry industry and causes huge economic losses (Goryo et al., 1985; Abdel-Mawgod et al., 2018).

Since the discovery of CIAV was first reported in young broilers in 1996 in China, it has been widespread in different provinces and has caused huge economic losses in the poultry industry (Zhou et al., 1996). A seroprevalence report on CIAV in the 1990s showed that its prevalence was about 42% in farms in five provinces of China (Zhou et al., 1996). In the live poultry market in southeastern China live poultry market, the prevalence of CIAV was as high as 87% during 2004-2005 (Ducatez et al., 2008). Furthermore, the investigation of 12 provinces in China from 2014 to 2015 showed that the CIAV positivity rate was 13.30% (Yao et al., 2019). Recently, a survey showed that 91 of 277 chickens collected in a molecular epidemiological survey in Guangdong, China from 2016 to 2017 were positive for CIAV (Tan et al., 2020). Moreover, a survey showed that the CIAV positive rate was 47.58% by detecting tissue samples of 330 in 11 provinces in northern Vietnam in 2019 (Van Dong et al., 2019).

After infection with CIAV, the immune system of the body was suppressed, which was easy to cause secondary infection or complications, thus making prevention and control more difficult and expanding indirect economic losses (Hagood et al., 2000). The previous report showed that the Newcastle disease virus attenuated vaccine was detected CIAV and fowl adenovirus (FAdV) (Brown Jordan et al., 2019; Su et al., 2019). Chickens inoculated with contaminated CIAV vaccines became a subclinical viral infection that expanded the area of infection (Li et al., 2017). In addition, cross-infection made prevention and control more difficult, and should be brought to the attention of the stock farmers.

Nowadays, we use the methods of pathogen isolation and molecular biological detection for CIAV in the laboratory (Miller et al., 2003; Song et al., 2018; Kannaki et al., 2021), and serological tests such as enzyme-linked immunosorbent assay (ELISA) are used in clinical (Bhatt et al., 2011; Kabir et al., 2021). However, pathogen isolation and molecular biological detection require skillful personnel and expensive equipment. The target of ELISA was antibodies, which was limited by the time and concentration of antibody production, as someone cannot accurately detect it. Thus, a low-cost, high-sensitivity, specific, portable, and simple operable method is imperative to detect CIAV on site.

The real-time recombinase-aided amplification assay (RAA) method is a new molecular biology technique

combining RAA with a portable fluorescence detection instrument to realize on-site detection (Wu et al., 2022a). The portable instrument has a built-in lithium battery, which can be used without plugging in grassroots farms with poor environmental conditions. The RAA method has been applied in food safety testing, environmental monitoring, aquatic disease detection, and animal and plant quarantine (Wang et al., 2019, 2021; Deng et al., 2021; He et al., 2021). They reported that RAA was applied in the detection of the Novel Coronavirus Disease 2019 (COVID-19) (Xue et al., 2020; Wang et al., 2020a; Shen et al., 2021). Previously, we established the application of real-time RT-RAA in avian leukosis virus subgroup J (ALV-J) and porcine epidemic diarrhea virus (PEDV) detection (Wu et al., 2022a,b). However, until now, there has been no report on CIAV detection by real-time RAA.

The RAA principle specifies that the recombinase and primer are tightly mixed at 35–40°C to form a polymer. With the help of single-stranded DNA binding protein and DNA polymerase, the primers search for complementary sequences on the template, and a new complementary DNA strand is formed (Chen et al., 2021). It is stable when the nucleic acid exonuclease is not activated, but when the probe begins to bind to the target DNA site, the exonuclease becomes active. When the nucleic acid exonuclease recognizes tetrahydrofuran (THF) on the probe, it cleaves both the reporter group and the burst group, releasing the reporter group. A blocker was removed as the probe ascended to the 3' end, and the probe continued to amplify until the 5' end. As a result, real-time fluorescence signals were gathered and plotted.

In this study, we used the VP2 gene sequence as the target for real-time RAA detection because it was the most conserved in CIAV, and the expression level of VP2 was higher than that of VP1 after infection in 12 h according to the previous reports (Douglas et al., 1995; Cheng et al., 2012; Lai et al., 2017).

Materials and methods

Virus and clinical specimens

The Marek's Disease Virus (MDV, GenBank: L37202) vaccine was purchased from Harbin Pharmaceutical Group Bio-vaccine Co., Ltd. (Harbin, China). Wen's Foodstuff Group Co, Ltd. (Yunfu, China) contributed to the Avian Reticuloendothelial Hyperplasia Virus (REV, not uploaded). In our laboratory, we retained the H9N2 subtype Avian Influenza Virus (H9N2, GenBank: MN064851), Infections Bursal Disease Virus (IBDV, GenBank: AF416621), Chicken Infectious Anemia Virus (CIAV, GenBank: JX260426), Newcastle Disease Virus (NDV, GenBank: JX458823), Infectious Laryngotracheitis Virus (ILTV, GenBank: JX458823), Avian Leukemia Virus (ALV, GenBank: MT175600) and Infectious Bronchitis Virus (IBV, GenBank: KR605489).

Wen's Foodstuff Group Co., Ltd. donated 42 clinical samples for this trial (Yunfu, China). Thymus tissue samples from chickens

TABLE 1 Real-time qPCR and PCR assays of primers sequences for CIAV.

Assay	Primer	Sequence (5'-3')	Localization
PCR	PCR-F	CAGTGAATCGGCGCTTAGCCG	394-414
	PCR-R	GATACCGCTGTCTCCTCCGA	933-952
Real-	qPCR-F	CGGACCATCAACGGTGTTCAGG	494-515
time	qPCR-R	GCAGCCACACAGCGATAGAGTG	590-611
qPCR			

suspected of being infected with CIAV were found from clinical samples and stored at -80° C in an ultralow temperature freezer.

Sample preparation

The nucleic acid of MDV, ALV, IBV, NDV, IBDV, H9N2, ILTV, REV, and CIAV (GenBank: JX260426) were extracted following the instructions of the AxyPrep Body Fluid Viral DNA/RNA Miniprep Kit (AP-MN-BF-VNA-250G, Axygen, New York, United States). The forty-two clinical samples were extracted using the steps specified by the HiPure Tissue DNA Mini Kit (D3121, Magen, Guangzhou, China). The nucleic acid was then kept in an ultralow-temperature freezer at a temperature of 80°C.

Reagents and instruments

The HiPure Tissue DNA Mini Kit was purchased from Guangzhou Magen Bio & Tech Co., Ltd. (D3121, Guangzhou, China). The AxyPrep Body Fluid Viral DNA/RNA Miniprep Kit was produced by Corning Incorporated (AP-MN-BF-VNA-250G, New York, United States). The freeze-dried powder for the real-time fluorescent RAA reaction was produced by Nanning Zhuangbo Bio & Tech Co., Ltd. (ZBA12001, Nanning, China). The EZ-press qPCR Kit was produced by Suzhou Yingze Biomedical Technology Co., Ltd. (EZB-qRT-R2, Suzhou, China). The pMDTM19-T Vector Cloning Kit was produced by Takara Biomedical Technology (Beijing) Co., Ltd. (6,013, Beijing, China). This study used a Bio-Rad Laboratories CFX96 real-time fluorescence quantitative PCR apparatus (Shanghai, China).

Generation of plasmid standard

The CIAV VP2 gene was amplified by PCR (Table 1). The primers were designed based on GD-1-12 (JX260426) and the product can be amplified at 559 bp. Recovery of PCR amplification products and cloned into pMD®19-T vectors (TaKaRa, China). After extraction, the concentration of this plasmid was calculated using the formula $(6.02 \times 10^{23}) \times (ng/\mu L \times 10^{-9})$ / (DNA length×

660), with the copy number of the CIAV VP2 gene being almost 2.92×10^{10} copies/µL.

Primers and probe

The nucleotide sequences of CIAV VP2 were downloaded from the database from the National Center for Biotechnology Information¹ (GenBank: AF285882, AF311892, AF311900, AF390038, AY040631, DQ991394, JX260426, KC414026, KF224934, KF224937, KJ872513, KU050680, KY486138, KY486141, KY888890, KY888903, KY888907, KY888910, KY888928, MK358456, M55918, MK687564, MN299313, MT268199, and MT795930), and we used the MEGAX64 (Mega Limited, Auckland, New Zealand) to align multiple sequences. Moreover, the Megalign of DNASTAR (DNASTAR, Inc., Madison, United States) was used to analyze homology and find the best region for designing the real-time RAA's primers and probes. We also designed primers for real-time qPCR on conserved genes of CIAV VP2. The primers and probes were synthesized by Sangon Biotech (Shanghai) Co., Ltd. (Shanghai, China).

Standard curve for real-time qPCR

The prepared CIAV VP2 gene plasmid was serially folded, and 6 dilution gradients of 10^7-10^2 copies/µL were used to construct a standard curve for real-time qPCR. Each dilution gradient was taken 2µl and repeated three times. The real-time qPCR reaction system was as follows: qPCR-F (10µM) 1.0µl, qPCR-R (10µM) 1.0µl (Table 1), ddH₂O 6.0µl, 2 x SYBR Green qPCR Master Mix 10.0µl, templates 2µl. The real-time qPCR reaction procedure was: 1 cycle of 95°C for 5 min; 40 cycles by 95°C for 10 s, 60°C for 30 s, and fluorescent signals were collected during the elongation step. The protocol of melting curves was produced by monitoring the real-time SYBR green signal from 65°C to 95°C, incrementing 0.5°C for 0.05 s.

The real-time RAA reaction system

The system of the real-time RAA per reaction was a total volume of 50 µl, including RT-RAA freeze-dried powder 1 tube, PEG buffer 25 µl, ddH₂O 12.9 µl, MgAc₂ buffer 2.5 µl, RAA-F (10 µM) 2.0 µl, RAA-R (10 µM) 2.0 µl, RAA-P (10 µM) 0.6 µl, and templates 5 µl. The real-time RAA amplification protocol was 1 cycle of 41°C for 60 s; 40 cycles of 41°C for 30 s and fluorescence signals were collected. After the reaction finished, the positive control had a smooth amplification curve, and the Ct value was \leq 39; the negative control has no

¹ https://www.ncbi.nlm.nih.gov

amplification curve, or the Ct value was>39, showing the assay has valid results.

RAA primers verification and screening

A total of 12 groups of real-time primers were F1R1, F1R2, F1R3, F1R4, F2R1, F2R2, F2R3, F1R4, F3R1, F3R2, F3R3, and F1R4 (Table 2), and the size of the amplified product of the primers was verified by PCR. Meanwhile, the sensitivity of 12 groups of primers, respectively, in the real-time RAA system was evaluated. Based on PCR and real-time RAA results, the greatest primers were chosen. The PCR system and procedure as follows: forward primer (10μ M) 1.0μ l, reverse primer (10μ M) 1.0μ l, ddH₂O 6.0μ l, 2xEsTaq Master Mix 10.0 μ l, templates 2 μ l; 95°C for 3 min; 95°C for 30 s, 58°C for 30 s, 72°C for 30 s, 35 cycles; 72°C for 5 min. 1.5% agarose gel electrophoresis was used to analyze the PCR products.

Sensitivity analysis of the real-time RAA assay

The CIAV VP2 standard plasmid was serially diluted 10 times, and six concentration gradients were utilized as templates: 10^5 copies/µL, 10^4 copies/µL, 10^3 copies/µL, 10^2 copies/µL, 10^1 copies/ µL, and 10^0 copies/µL. The sensitivity assay was then analyzed by repeating it five times. To achieve an accurate evaluation of sensitivity, the different dilution series (10^4 – 10^{-1} TCID50 CIAV DNA per reaction) were used as templates in both real-time qPCR and real-time RAA analyses five-time independently.

Specificity analysis of the real-time RAA assay

For the real-time RAA, the CIAV nucleic acid was performed as a positive control, while ddH₂O worked as a negative control.

TABLE 2 Real-time RAA assays of primers and probes for CIAV.

Primer/ probe	Sequence (5'-3')	Localization
F1	TTCACGGCCGTTGGAAACCCCTCACTG	527-553
F2	AAACCCCTCACTGCAGAGAGAGATCCGGATTG	541-570
F3	AACAAGTTCACGGCCGTTGGAAACCCC	521-547
R1	TTGTCCGCAGTTGCAGATCTTAGCGTGG	628-655
R2	TTCGAGGGAGGCTTGGGTTGATCGGTC	695-721
R3	AATTGTCCGCAGTTGCAGATCTTAGCGTG	629-657
R4	TTCGAGGGAGGCTTGGGTTGATCGGTCC	694-721
probe	ATCGCTGGAATTACAATCACTCTATCGCTG/	573-621
	i6FAMdT//THF//iBHQ1dT/	
	GGCTGCGCGAATGCTC[C3-spacer]	

FAM, 6-carboxyfluorescein; THF, tetrahydrofuran; BHQ, black hole quencher; C3-spacer, 3' phosphate blocker.

For specificity analysis of MDV, ALV, IBV, NDV, IBDV, H9N2, ILTV, and REV nucleic acids, $2 \mu l$ of templates were acquired and repeated three times.

Clinical sample evaluation of the real-time RAA assay

Real-time qPCR and real-time RAA were utilized to detect nucleic acids from 42 clinical samples suspected of being infected with CIAV. We compared and analyzed the results of real-time qPCR and RAA to evaluate the real-time RT-RAA method developed in this study.

Statistical analysis

Results of real-time RAA and real-time qPCR assays were measured with kappa and p-values at 95% reliability using IBM's SPSS software.

Results

Designing the real-time RAA primers and probe of CIAV

Based on the VP2 sequence of the GD-1-12 (JX260426) strain isolated in our laboratory, we performed multiple sequence alignment with the sequence published in the NCBI database using Megalign's castal W method. The homology values of CIAV VP2 sequences were over 99.4%, indicating great conservatism, as shown in Figure 1. We designed primers and probes for CIAV real-time RAA on this gene fragment depending on the results of homology analysis, as shown in Table 2. The sequences of the CIAV VP2 had not intersected with other viruses, according to NCBI's BLAST analysis of the amplified fragments of the primers and probe.

Verification and screening of real-time RAA primers for CIAV

PCR products were performed by 1.5% gel electrophoresis to verify the size and specificity of real-time RAA primers. Figure 2A showed the results. The size of amplified fragments from 12 different primers both met predictions and did no-primer-dimers, which has good specificity for CIAV. Figure 2B showed the results of a real-time RAA process using the nucleic acid of CIAV as the template. Under the same template concentration, F2R3 had a higher amplification efficiency than the other primers. Figure 2C showed the result of multiple sequence alignment of F2R3 and probe sequences.

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19		
1		99.7	99.7	99.8	99.4	99.8	99.4	99.8	99.4	99.8	99.5	99.4	99.7	99.8	99.5	99.8	99.8	99.4	99.7	1	AF285882
2	0.3		99.7	99.8	99.4	99.8	99.4	99.8	99.4	99.8	99.5	99.4	99.7	99.8	99.5	99.8	99.8	99.4	99.7	2	AF311892
3	0.3	0.3		99.8	99.4	99.8	99.4	99.8	99.4	99.8	99.5	99.7	99.7	99.8	99.5	99.8	99.8	99.4	99.7	3	AF311900
4	0.2	0.2	0.2		99.5	100.0	99.5	100.0	99.5	100.0	99.7	99.5	99.8	100.0	99.7	100.0	100.0	99.5	99.8	4	AF390038
5	0.6	0.6	0.6	0.5		99.5	99.0	99.5	99.0	99.5	99.2	99.0	99.4	99.5	99.2	99.5	99.5	99.4	99.4	5	AY040632
6	0.2	0.2	0.2	0.0	0.5		99.5	100.0	99.5	100.0	99.7	99.5	99.8	100.0	99.7	100.0	100.0	99.5	99.8	6	DQ991394
7	0.6	0.6	0.6	0.5	1.0	0.5		99.5	99.0	99.5	99.2	99.0	99.4	99.5	99.2	99.5	99.5	99.0	99.4	7	KC414026
8	0.2	0.2	0.2	0.0	0.5	0.0	0.5		99.5	100.0	99.7	99.5	99.8	100.0	99.7	100.0	100.0	99.5	99.8	8	KF224934
9	0.6	0.6	0.6	0.5	1.0	0.5	1.0	0.5		99.5	99.2	99.0	99.4	99.5	99.2	99.5	99.5	99.0	99.4	9	KF224937
10	0.2	0.2	0.2	0.0	0.5	0.0	0.5	0.0	0.5		99.7	99.5	99.8	100.0	99.7	100.0	100.0	99.5	99.8	10	MH269372
11	0.5	0.5	0.5	0.3	0.8	0.3	0.8	0.3	0.8	0.3		99.2	99.5	99.7	99.4	99.7	99.7	99.2	99.5	11	MT813075
12	0.6	0.6	0.3	0.5	1.0	0.5	1.0	0.5	1.0	0.5	0.8		99.4	99.5	99.2	99.5	99.5	99.0	99.4	12	MT813076
13	0.3	0.3	0.3	0.2	0.6	0.2	0.6	0.2	0.6	0.2	0.5	0.6		99.8	99.5	99.8	99.8	99.4	99.7	13	MT813077
14	0.2	0.2	0.2	0.0	0.5	0.0	0.5	0.0	0.5	0.0	0.3	0.5	0.2		99.7	100.0	100.0	99.5	99.8	14	MW817567
15	0.5	0.5	0.5	0.3	0.8	0.3	0.8	0.3	0.8	0.3	0.6	0.8	0.5	0.3		99.7	99.7	99.2	99.5	15	MW817568
16	0.2	0.2	0.2	0.0	0.5	0.0	0.5	0.0	0.5	0.0	0.3	0.5	0.2	0.0	0.3		100.0	99.5	99.8	16	MW817569
17	0.2	0.2	0.2	0.0	0.5	0.0	0.5	0.0	0.5	0.0	0.3	0.5	0.2	0.0	0.3	0.0		99.5	99.8	17	MW817570
18	0.6	0.6	0.6	0.5	0.6	0.5	1.0	0.5	1.0	0.5	0.8	1.0	0.6	0.5	0.8	0.5	0.5		99.4	18	M55918
19	0.3	0.3	0.3	0.2	0.6	0.2	0.6	0.2	0.6	0.2	0.5	0.6	0.3	0.2	0.5	0.2	0.2	0.6		19	GD-1-12
	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19		
RE 1																					

amplification assay.

F2R3 was chosen as the best primer for CIAV real-time RAA based on the above results.

Standard curve of real-time qPCR for CIAV

According to the results, the standard curve exhibits a good linear relationship with the correlation coefficient of R² value was 0.999, the amplification efficiency was 90.2%, and the melt temperature was 83.5° C. The logarithmic function relationship between cycle threshold and plasmid copy number was y = -3.580x + 36.753 and was shown in Figures 3A,B.

Sensitivity analysis of The real-time RAA assay for CIAV

As shown in Figure 4, we chose a typical result from the five independent real-time RAA assays and real-time qPCR. Figures 4A–C showed that the real-time RAA in 10^1 copies/µL of standard plasmid, and 10^0 TCID50/reaction of CIAV, and the real-time qPCR 10^1 TCID50/reaction had fluorescence signals among the five times results from the six concentrations gradients. Figures 4D–F showed probit regression analysis of the real-time RAA and qPCR, using SPSS software on data from the five runs of standard plasmid and CIAV DNA. As a result, the limited sensitivity of the real-time RAA we developed for CIAV was 10 copies of standard plasmid and 1 TCID50 of CIAV can be detected per reaction.

Specificity analysis of the real-time RAA assay for CIAV

Figure 5 showed a representative result from the three independent real-time RAA assays. The CIAV nucleic acid positive control had a single fluorescence signal curve. There was no fluorescence signal in the negative controls of ddH_2O , H9N2, MDV, ALV, ILTV, IBV, NDV, REV, and IBDV nucleic acid. As a result, we developed the real-time RAA method that could specifically identify CIAV while exhibiting no cross-reactivity with MDV, ALV, IBV, NDV, IBDV, H9N2, ILTV, or REV.

Clinical sample evaluation of the real-time RAA assay for CIAV

Table 3 showed that of the nucleic acids of the 42 clinical samples, 14 nucleic acids were positive amplification by realtime qPCR and real-time RAA assays. However, one of the nucleic acids showed positive amplification by real-time qPCR but negative amplification by real-time RAA. Real-time qPCR and real-time RAA both have a sensitivity of 93.33%. None of the 27 nucleic acids had a fluorescence signal for real-time qPCR and real-time RAA assays, indicating negative amplification with a specificity of 100%. This assay's kappa value of real-time RAA was 0.947 compared to the real-time qPCR assay, showing that the two methods were almost perfect.



control, i.e., ddH₂O. RAA, recombinase-aided amplification assay; CIAV, Chicken Infectious Anemia Virus.



Standard curve and melting curve of real-time qPCR for CIAV. (A) The logarithmic function relationship between cycle threshold and plasmid copy number was y=-3.580x+36.753, $R^2=0.999$, and efficiency was 90.2%. (B) The melt temperature of real-time qPCR for CIAV was 83.5°C. CIAV, Chicken Infectious Anemia Virus; qPCR, quantitative polymerase chain reaction assay.

Discussion

Researchers have developed many vaccines to prevent and control CIAV from an immunization perspective to reduce the

significant production losses carried on by clinical and subclinical infections in commercial poultry because of the prevalence of CIAV in producing countries' broilers (Barbour et al., 2002; Vaziry et al., 2011; Sawant et al., 2015). However, the CIAV antibody response remained low in most vaccinated



chicks and did not persist until day 18 post-immunization, and the vaccine strain persisted in the thymus and spleen of young chicks, resulting in a low humoral immune response and inducing thymic lymphoid cell disorders (Vaziry et al., 2011). Persistent infection was more likely to result from this subclinical infection phenomenon. For example, the dual infection of CIAV and infectious bursal disease virus (IBDV) had no obvious clinical symptoms, but the detection of the thymus showed persistent infection of CIAV (Vaziry et al., 2013). The CIAV co-infection with FAdV had obvious inclusion of body hepatitis- hydropericardium syndrome (Su et al., 2019). The reality that CIAV can infect a different species complicates matters further and makes it more challenging to prevent and control (Zhang et al., 2014; Fang et al., 2017). Therefore, an effective, accurate, and simple-tooperate detection method is required to better monitor and prevent CIAV outbreaks while also reducing the poultry industry's economic losses.

Researchers have developed a variety of detection methods for CIAV, each with its own set of benefits and drawbacks. As indicated in Table 4, we provide a brief overview of the different CIAV detection methods. The virus isolation period of CIAV is the longest, at least 14 days (Buscaglia et al., 1994). Before noticeable cytopathic alterations may be seen with a fluorescence microscope, MDCC-MSB1 cells must be infected with CIAV for more than 24 h, and there are issues for immunofluorescence assays (IFA) detect CIAV with non-specificity and insufficient sensitivity (Lucio et al., 1991; Buscaglia et al., 1994). Compared to other detection methods, virus isolation and IFA require skilled, highquality personnel, and sterile negative pressure environments, both of which are more expensive. The most commonly detected CIAV with two methods, PCR and qPCR, with the assay taking



TABLE 3 Comparison of the results of real-time RAA assay and qPCR method for clinical samples.

		Real-tir	ne qPCR	Performance characteristics					
		Positive	Negative	Sensitivity (%)	Specificity (%)	Kappa			
Real-	Positive	14	0	93.33		0.947			
time	Negative	1	27		100				
RAA	Total	15	27						

between 108 and 137 min (Todd et al., 1992; Kaffashi et al., 2021). However, PCR requires variable temperature equipment and electrophoresis (Zhang et al., 2020). It is possible to obtain accurate results in real-time with qPCR, but it requires expensive equipment and partitioned reaction systems to reduce the falsepositive rate (Yoo et al., 2021). LAMP needs four pairs of primers to detect CIAV at 63°C for 40 min, and the results can be visualized, making it convenient for clinical testing (Song et al., 2018). However, four pairs of primers with eight ssDNA in one tube are prone to form primer-dimers, increasing the LAMP detection of false-positive results (Wang et al., 2020b). In addition to being the easiest method of detection, the ELISA test also is commonly used in clinical to detect CIAV (Pallister et al., 1994; Tannock et al., 2003). Due to the low level of CIAV antibodies in the vaccinated chicks, no antibody can be detected. There are problems with non-specific binding and low sensitivity with an ELISA (Michalski et al., 1996).

In this study, we developed the real-time RAA that can overcome the deficiencies of the above method in the clinical detection of CIAV. When combined with recombinase at a constant temperature (39-42°C), the real-time RAA primers only further amplify once it matches the template with the completed complementary sequence, thus reducing error tolerance and improving detection accuracy. We designed 12 groups of primers for the highly conserved VP2 gene to select the optimal primers and performed double screening by PCR and real-time RAA. We used the screened F2R3 primers to evaluate its specificity with MDV, ALV, IBV, NDV, IBDV, H9N2, ILTV, and REV, and the results showed that there was no amplification reaction with these viruses, which further proved that the primers we designed can specifically detect CIAV. Sensitivity demonstrated that real-time RAA could detect a limitation of 10 copies per reaction of standard plasmid, and 1 TCID50 per reaction of CIAV, and the detection results of six concentration gradients in five repeated experiments had a 100% coincidence rate, which also showed that our established real-time RAA had high sensitivity and reproducibility. In other words, real-time RAA can be used helpfully for clinical diagnosis, whether it is infected by CIAV in the clinical

Methods	Sensitivity	Instrument requirements	Assay time	On-site	References
Real-time RAA	10 copies	Portable fluorescence reporter	30 min	YES	This study
PCR	0.1 pg	PCR; Electrophoresis System;	137 min	NO	Todd et al., 1992
		Transilluminator			
qPCR	21 copies	Fluorescence PCR	108 min	NO	Kaffashi et al., 2021
LAMP	50 copies	Portable fluorescence reporter	40 min	YES	Song et al., 2018
ELISA	-	ELIASA	135 min	YES	Li, 2021
IFA	-	Fluorescence microscope	24 h	NO	Lamichhane et al., 1992
Virus isolation	-	Laboratory of cell biology	At least 14 day	NO	Buscaglia et al., 1994

TABLE 4 Different detection methods for CIAV.

symptom or early stages. Noteworthy, the real-time RAA detection results of 42 clinical samples compared with qPCR, the performance characteristics of kappa value was 0.947, sensitivity was 93.33%, and specificity was 100%, indicating the accuracy of the results was consistent with qPCR. What is more superior is that real-time RAA can be completed within 30 min at 41°C and combined with a portable fluorescence detector to move detection in a complex environment to complete accurate detection leave out the laboratory.

The real-time RAA method for rapid detection of CIAV has the characteristics of simple operation, high sensitivity, strong specificity, and short detection time that can better prevent and control CIAV in clinical. The real-time RAA has excellent detection performance and can be well implemented in early diagnosis, long-term monitoring, clinical mobile detection, and cross-host transmission of CIAV.

Conclusion

This study successfully established and applied a real-time RAA method to detect CIAV within 30 min at 41°C. In the poultry industry, real-time RT-RAA offers high performance in sensitivity, specificity, repeatability, and low cost, which could be key for preventing and controlling CIAV.

Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

Ethics statement

Institutional and National Guidelines for the use and care of laboratory animals were closely followed. The use of animals in this study was approved by the South China Agricultural University Committee for Animal Experiments (approval ID: SYXK2019-0136).

Author contributions

XW conducted the experiments, analyzed the data, and wrote the manuscript. JK, ZY, HS, HZ, HH, JW, MC, YZ, and YH conducted the experiments. FC analyzed part of the data. YL, ZW, and JL checked and finalized the manuscript. QX and XZ designed the experiments, modified the manuscript, and supervised the whole work. All authors have read and approved the final manuscript.

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Conflict of interest

JL and ZW were employed by Wen's Foodstuffs Group Co., Ltd.

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

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References

Abdel-Mawgod, S., Adel, A., Arafa, A. S., and Hussein, H. A. (2018). Full genome sequences of chicken anemia virus demonstrate mutations associated with pathogenicity in two different field isolates in Egypt. *Virus* 29, 333–341. doi: 10.1007/s13337-018-0467-z

Barbour, E. K., Farran, M. T., Hamadeh, S. K., Bouljihad, M., Faroon, O., and Kreydiyyeh, S. (2002). Comparative impact of live chicken infectious anaemia virus vaccine versus natural exposure in meat chicken breeders on immunity to infectivity by CIA and inclusion body hepatitis viruses in their offspring. *Vet. Res. Commun.* 26, 397–405. doi: 10.1023/a:1016251013681

Bhatt, P., Shukla, S. K., Mahendran, M., Dhama, K., Chawak, M. M., and Kataria, J. M. (2011). Prevalence of chicken infectious anaemia virus (CIAV) in commercial poultry flocks of northern India: a serological survey. *Transbound. Emerg. Dis.* 58, 458–460. doi: 10.1111/j.1865-1682.2011.01215.x

Brown Jordan, A., Blake, L., Bisnath, J., Ramgattie, C., Carrington, C. V., and Oura, C. A. L. (2019). Identification of four serotypes of fowl adenovirus in clinically affected commercial poultry co-infected with chicken infectious anaemia virus in Trinidad and Tobago. *Transbound. Emerg. Dis.* 66, 1341–1348. doi: 10.1111/tbed.13162

Buscaglia, C., Crosetti, C. F., and Nervi, P. (1994). Identification of chicken infectious anaemia, isolation of the virus and reproduction of the disease in Argentina. *Avian Pathol.* 23, 297–304. doi: 10.1080/03079459408418997

Chen, W., Fan, J., Li, Z., Zhang, Y., Qin, Y., Wu, K., et al. (2021). Development of Recombinase aided amplification combined with disposable nucleic acid test strip for rapid detection of porcine circovirus type 2. *Front. Vet. Sci.* 8:676294. doi: 10.3389/fvets.2021.676294

Cheng, J. H., Sheu, S. C., Lien, Y. Y., Lee, M. S., Chen, H. J., Su, W. H., et al. (2012). Identification of the NLS and NES motifs of VP2 from chicken anemia virus and the interaction of VP2 with mini-chromosome maintenance protein 3. *BMC Vet. Res.* 8:15. doi: 10.1186/1746-6148-8-15

Deng, Y., Liu, Y. H., Chen, W. Q., Zhang, Y. L., Jiang, T. T., Li, S. H., et al. (2021). Establishment of a fluorescent recombinase-aided isothermal amplification assay for nucleic acid detection of *Paraginiumus skrjabini* and preliminary evaluation of its detection efficiency. *Zhongguo Xue Xi Chong Bing Fang Zhi Za Zhi* 33, 464–469. doi: 10.16250/j.32.1374.2021218

Douglas, A. J., Phenix, K., Mawhinney, K. A., Todd, D., Mackie, D. P., and Curran, W. L. (1995). Identification of a 24 kDa protein expressed by chicken anaemia virus. *J. Gen. Virol.* 76, 1557–1562. doi: 10.1099/0022-1317-76-7-1557

Ducatez, M. F., Chen, H., Guan, Y., and Muller, C. P. (2008). Molecular epidemiology of chicken anemia virus (CAV) in southeastern Chinese live birds markets. *Avian Dis.* 52, 68–73. doi: 10.1637/8049-070407-Reg

Ducatez, M. F., Owoade, A. A., Abiola, J. O., and Muller, C. P. (2006). Molecular epidemiology of chicken anemia virus in Nigeria. *Arch. Virol.* 151, 97–111. doi: 10.1007/s00705-005-0609-7

Fang, L., Li, Y., Wang, Y., Fu, J., Cui, S., Li, X., et al. (2017). Genetic analysis of two chicken infectious anemia virus variants-related gyrovirus in stray mice and dogs: the first report in China, 2015. *Biomed. Res. Int.* 2017:6707868. doi: 10.1155/2017/6707868

Goryo, M., Sugimura, H., Matsumoto, S., Umemura, T., and Itakura, C. (1985). Isolation of an agent inducing chicken anaemia. *Avian Pathol.* 14, 483–496. doi: 10.1080/03079458508436251

Hagood, L. T., Kelly, T. F., Wright, J. C., and Hoerr, F. J. (2000). Evaluation of chicken infectious anemia virus and associated risk factors with disease and production losses in broilers. *Avian Dis.* 44, 803–808. doi: 10.2307/1593052

He, Z., Tong, Z., Tan, B., He, X., Zhang, T., Guo, Y., et al. (2021). Rapid detection of DNA methylation with a novel real-time fluorescence recombinase-aided amplification assay. *J. Biomed. Nanotechnol.* 17, 1364–1370. doi: 10.1166/jbn. 2021.3111

Kabir, M. A. A., Saha, S., Hossain, M. G., Khan, K. A., Islam, M. A., and Rahman, L. (2021). Serological survey on the prevalence of chicken infectious anemia virus in broiler breeder and layer farms in some selected areas of Bangladesh. *J. Adv. Vet. Anim. Res.* 8, 323–329. doi: 10.5455/javar.2021.h518

Kaffashi, A., Mahmoudzadeh, M., and Ataei Kachooei, S. (2021). Development of TaqMan real-time polymerase chain reaction assay for detection of chicken anemia virus in Newcastle disease vaccines. *Arch. Razi. Inst.* 76, 421–427. doi: 10.22092/ari.2020.342677.1474

Kannaki, T. R., Priyanka, E., Subbiah, M., and Haunshi, S. (2021). Development and validation of high throughput real-time polymerase chain reaction assay for quantitative detection of chicken infectious anemia virus. *Virus* 32, 343–346. doi: 10.1007/s13337-020-00648-1

Lai, G. H., Lien, Y. Y., Lin, M. K., Cheng, J. H., Tzen, J. T., Sun, F. C., et al. (2017). VP2 of chicken anaemia virus interacts with apoptin for down-regulation of

apoptosis through de-phosphorylated threonine 108 on apoptin. *Sci. Rep.* 7:14799. doi: 10.1038/s41598-017-14558-8

Lamichhane, C. M., Snyder, D. B., Girschick, T., Goodwin, M. A., and Miller, S. L. (1992). Development and comparison of serologic methods for diagnosing chicken anemia virus infection. *Avian Dis.* 36, 725–729. doi: 10.2307/1591773

Li, J. (2021). Isolation and identification of chicken infectious anemia virus, preparation of DNA qualitative standard and establishment and preliminary application of antibody indirect ELISA. Master Thesis Yangzhou University.

Li, Y., Hu, Y., Cui, S., Fu, J., Wang, Y., Cui, Z., et al. (2017). Molecular characterization of chicken infectious anemia virus from contaminated live-virus vaccines. *Poult. Sci.* 96, 1045–1051. doi: 10.3382/ps/pew406

Lucio, B., Schat, K. A., and Taylor, S. (1991). Direct binding of protein a, protein G, and anti-IgG conjugates to chicken infectious anemia virus. *Avian Dis.* 35, 180–185. doi: 10.2307/1591311

Michalski, W. P., O'Rourke, D., and Bagust, T. J. (1996). Chicken anaemia virus antibody ELISA: problems with non-specific reactions. *Avian Pathol.* 25, 245–254. doi: 10.1080/03079459608419139

Miller, M. M., Ealey, K. A., Oswald, W. B., and Schat, K. A. (2003). Detection of chicken anemia virus DNA in embryonal tissues and eggshell membranes. *Avian Dis.* 47, 662–671. doi: 10.1637/7007

Natesan, S., Kataria, J. M., Dhama, K., Rahul, S., and Bhardwaj, N. (2006). Biological and molecular characterization of chicken anaemia virus isolates of Indian origin. *Virus Res.* 118, 78–86. doi: 10.1016/j.virusres.2005.11.017

Noteborn, M. H., de Boer, G. F., van Roozelaar, D. J., Karreman, C., Kranenburg, O., Vos, J. G., et al. (1991). Characterization of cloned chicken anemia virus DNA that contains all elements for the infectious replication cycle. *J. Virol.* 65, 3131–3139. doi: 10.1128/JVI.65.6.3131-3139.1991

Pallister, J., Fahey, K. J., and Sheppard, M. (1994). Cloning and sequencing of the chicken anaemia virus (CAV) ORF-3 gene, and the development of an ELISA for the detection of serum antibody to CAV. *Vet. Microbiol.* 39, 167–178. doi: 10.1016/0378-1135(94)90097-3

Sawant, P. M., Dhama, K., Rawool, D. B., Wani, M. Y., Tiwari, R., Singh, S. D., et al. (2015). Development of a DNA vaccine for chicken infectious anemia and its immunogenicity studies using high mobility group box 1 protein as a novel immunoadjuvant indicated induction of promising protective immune responses. *Vaccine* 33, 333–340. doi: 10.1016/j.vaccine.2014.11.020

Shao, H., Li, J., Yuan, H., Ji, L., Zhang, J., Jin, W., et al. (2021). Isolation and molecular characteristics of a CIAV isolate from pigeons. *China. Front. Vet. Sci.* 8:669154. doi: 10.3389/fvets.2021.669154

Shen, X., Wang, J., Li, J., He, A., and Liu, H. (2021). Ma X (2021) field validation of a rapid recombinase aided amplification assay for SARS-CoV-2 RNA at customs – Zhejiang Province, China, January 2021. *China CDC Wkly* 3, 973–976. doi: 10.46234/ccdcw2021.236

Song, H., Bae, Y., Park, S., Kwon, H., Lee, H., and Joh, S. (2018). Loop-mediated isothermal amplification assay for detection of four immunosuppressive viruses in chicken. *J. Virol. Methods* 256, 6–11. doi: 10.1016/j.jviromet.2018.02.013

Su, Q., Meng, F., Li, Y., Zhang, Y., Zhang, Z., Cui, Z., et al. (2019). Chicken infectious anemia virus helps fowl adenovirus break the protection of maternal antibody and cause inclusion body hepatitis-hydropericardium syndrome in layers after using co-contaminated Newcastle disease virus-attenuated vaccine. *Poult. Sci.* 98, 621–628. doi: 10.3382/ps/pey153

Tan, C., Wang, Z., Lei, X., Lu, J., Yan, Z., Qin, J., et al. (2020). Epidemiology, molecular characterization, and recombination analysis of chicken anemia virus in Guangdong province. *China. Arch. Virol.* 165, 1409–1417. doi: 10.1007/s00705-020-04604-8

Tannock, G. A., Tan, J., Mawhinney, K. A., Todd, D., O'Rourke, D., and Bagust, T. J. (2003). A modified blocking ELISA for the detection of antibody to chicken anaemia virus using an Australian strain. *Aust. Vet. J.* 81, 428–430. doi: 10.1111/j.1751-0813. 2003.tb11556.x

Todd, D., Mawhinney, K. A., and McNulty, M. S. (1992). Detection and differentiation of chicken anemia virus isolates by using the polymerase chain reaction. *J. Clin. Microbiol.* 30, 1661–1666. doi: 10.1128/jcm.30.7.1661-1666. 1992

Van Dong, H., Tran, G. T. H., Van Nguyen, G., Dao, T. D., Bui, V. N., Huynh, L. T. M., et al. (2019). Chicken anemia virus in northern Vietnam: molecular characterization reveals multiple genotypes and evidence of recombination. *Virus Genes* 55, 643–653. doi: 10.1007/s11262-019-01686-8

Vaziry, A., Silim, A., Bleau, C., Frenette, D., and Lamontagne, L. (2011). Chicken infectious anaemia vaccinal strain persists in the spleen and thymus of young chicks and induces thymic lymphoid cell disorders. *Avian Pathol.* 40, 377–385. doi: 10.1080/03079457.2011.586330

Vaziry, A., Silim, A., Bleau, C., Frenette, D., and Lamontagne, L. (2013). Dual infections with low virulent chicken infectious anaemia virus (lvCIAV) and intermediate infectious bursal disease virus (iIBDV) in young chicks increase lvCIAV in thymus and bursa while decreasing lymphocyte disorders induced by iIBDV. *Avian Pathol.* 42, 88–99. doi: 10.1080/03079457.2013.766306

Wang, S., Cai, G., Duan, H., Qi, W., and Lin, J. (2021). Automatic and multichannel detection of bacteria on a slidable centrifugal disc based on FTA card nucleic acid extraction and recombinase aided amplification. *Lab Chip* 22, 80–89. doi: 10.1039/d1lc00915j

Wang, J., Cai, K., He, X., Shen, X., Wang, J., Liu, J., et al. (2020a). Multiple-centre clinical evaluation of an ultrafast single-tube assay for SARS-CoV-2 RNA. *Clin. Microbiol. Infect.* 26, 1076–1081. doi: 10.1016/j.cmi.2020.05.007

Wang, Y., Sun, J., Guo, X., Zhang, D., Cui, Y., Li, W., et al. (2020b). TaqMan-based real-time polymerase chain reaction assay for specific detection of bocavirus-1 in domestic cats. *Mol. Cell. Probes* 53:101647. doi: 10.1016/j.mcp.2020.101647

Wang, R. H., Zhang, H., Zhang, Y., Li, X. N., Shen, X. X., Qi, J. J., et al. (2019). Development and evaluation of recombinase-aided amplification assays incorporating competitive internal controls for detection of human adenovirus serotypes 3 and 7. *Virol. J.* 16:86. doi: 10.1186/s12985-019-1178-9

Wu, X., Liu, Y., Gao, L., Yan, Z., Zhao, Q., Chen, F., et al. (2022a). Development and application of a reverse-transcription recombinase-aided amplification assay for porcine epidemic diarrhea virus. *Viruses* 14:591. doi: 10.3390/v14030591

Wu, X. H., Yao, Z. Q., Zhao, Q. Q., Chen, S., Hu, Z. Z., Xie, Z., et al. (2022b). Development and application of a reverse-transcription recombinase-aided

amplification assay for subgroup J avian leukosis virus. *Poult. Sci.* 101:101743. doi: 10.1016/j.psj.2022.101743

Xue, G., Li, S., Zhang, W., Du, B., Cui, J., Yan, C., et al. (2020). Reversetranscription recombinase-aided amplification assay for rapid detection of the 2019 novel coronavirus (SARS-CoV-2). *Anal. Chem.* 92, 9699–9705. doi: 10.1021/acs. analchem.0c01032

Yao, S., Tuo, T., Gao, X., Han, C., Yan, N., Liu, A., et al. (2019). Molecular epidemiology of chicken anaemia virus in sick chickens in China from 2014 to 2015. *PLoS One* 14:e0210696. doi: 10.1371/journal.pone.0210696

Yoo, H. M., Kim, I. H., and Kim, S. (2021). Nucleic acid testing of SARS-CoV-2. Int. J. Mol. Sci. 22:6150. doi: 10.3390/ijms22116150

Yuasa, N., Yoshida, I., and Taniguchi, T. (1976). Isolation of a reticuloendotheliosis virus from chickens inoculated with Marek's disease vaccine. *Natl. Inst. Anim. Health Q.* 16, 141–151. PMID: 189220

Zhang, X., Liu, Y., Ji, J., Chen, F., Sun, B., Xue, C., et al. (2014). Identification of a chicken anemia virus variant-related gyrovirus in stray cats in China, 2012. *Biomed. Res. Int.* 2014:313252. doi: 10.1155/2014/313252

Zhang, X., Yang, Q., Zhang, Q., Jiang, X., Wang, X., Li, Y., et al. (2020). Rapid detection of cytochrome cd1-containing nitrite reductase encoding gene nirS of denitrifying bacteria with loop-mediated isothermal amplification assay. *Sci. Rep.* 10:16484. doi: 10.1038/s41598-020-73304-9

Zhou, W., Yang, B., Shen, B., Han, S., and Zhou, J. (1996). A serologic survey of antibody against chicken infectious anemia virus by indirect immunofluorescent assay in domestic poultry in China. *Avian Dis.* 40, 358–360. doi: 10.2307/1592232