

METHODOLOGY Open Access

Visual detection of H3 subtype avian influenza viruses by reverse transcription loop-mediated isothermal amplification assay

Yi Peng¹, Zhixun Xie^{1*}, Jiabo Liu¹, Yaoshan Pang¹, Xianwen Deng¹, Zhiqin Xie¹, Liji Xie¹, Qing Fan¹, Jiaxun Feng³ and Mazhar I Khan^{2*}

Abstract

Background: Recent epidemiological investigation of different HA subtypes of avian influenza viruses (AlVs) shows that the H3 subtype is the most predominant among low pathogenic AlVs (LPAlVs), and the seasonal variations in isolation of H3 subtype AlVs are consistent with that of human H3 subtype influenza viruses. Consequently, the development of a rapid, simple, sensitive detection method for H3 subtype AlVs is required. The loop-mediated isothermal amplification (LAMP) assay is a simple, rapid, sensitive and cost-effective nucleic acid amplification method that does not require any specialized equipment.

Results: A reverse transcription loop-mediated isothermal amplification (RT-LAMP) assay was developed to detect the H3 subtype AlVs visually. Specific primer sets target the sequences of the hemagglutinin (HA) gene of H3 subtype AlVs were designed, and assay reaction conditions were optimized. The established assay was performed in a water bath for 50 minutes, and the amplification result was visualized directly as well as under ultraviolet (UV) light reflections. The detection limit of the RT-LAMP assay was 0.1pg total RNA of virus, which was one hundred-fold higher than that of RT-PCR. The results on specificity indicated that the assay had no cross-reactions with other subtype AlVs or avian respiratory pathogens. Furthermore, a total of 176 clinical samples collected from birds at the various live-bird markets (LBMs) were subjected to the H3-subtype-specific RT-LAMP (H3-RT-LAMP). Thirty-eight H3 subtype AlVs were identified from the 176 clinical samples that were consistent with that of virus isolation.

Conclusions: The newly developed H3-RT-LAMP assay is simple, sensitive, rapid and can identify H3 subtype AlVs visually. Consequently, it will be a very useful screening assay for the surveillance of H3 subtype AlVs in underequipped laboratories as well as in field conditions.

Keywords: Loop-mediated isothermal amplification, H3 subtype, avian influenza

Background

Influenza A viruses are classified into subtypes consisting of 16 hemagglutinin (HA) and 9 neuraminidase (NA) based on the antigenic differences of the HA and NA proteins, which are surface glycoproteins found on the viral envelope [1,3]. While the viruses with all subtypes can be detected in wild aquatic waterfowl, only few of the subtypes' influenza A viruses can infect mammalian

species [1]. Avian influenza viruses (AIVs) can be divided into two distinct groups based on their virulence: the highly virulent viruses, including H5-and H7-subtype AIVs, are called highly pathogenic avian influenza (HPAI) and cause high mortality and morbidity; all other subtypes are categorized as low pathogenic avian influenza (LPAI), which cause mild or no symptoms in birds [2.4.5]

It has been shown that LPAIVs from birds can be a potential source of reassortant human influenza A viruses. In 1957, the virus that caused the influenza pandemic was found to posses three genes from subtype H2N2 from an avian virus and all remaining genes from a circulating

²Department of Pathobiology and Veterinary Science, University of Connecticut, 61 North Eagleville Road Storrs, CT 06269-3089, USA Full list of author information is available at the end of the article



^{*} Correspondence: xiezhixun@126.com; mazhar.khan@uconn.edu

¹Department of Biotechnology, Guangxi Veterinary Research Institute, 51 You Ai Road, Nanning, Guangxi 530001, China

human H1N1 virus [1]. Moreover, the Hong Kong influenza virus (H3N2) in 1968 was a reassortant with avian (H3) PB1 and HA genes and six other genes from human (H2N2) virus [6].

Currently, there is no evidence that H5N1 HPAIVs can transmit widely between humans; the human-pandemic influenza strains are all generated by gene reassortment between human influenza viruses, avian influenza viruses and swine influenza viruses [7]. The influenza A viruses spreading in humans recently include seasonal H3N2, H1N1 and pandemic influenza A H1N1 viruses. It's notable that the results of a recent epidemiological investigation into different HA subtype AIVs shows that the H3 subtype is the predominant subtype among LPAIVs, and the seasonal variations in isolation of H3 subtype AIVs are consistent with that of human H3 subtype influenza viruses [8]. Some research predicts that H3 subtype AIVs will get the ability of infecting human directly through gene reassortment [9,10]. Consequently, it is important to enhance surveillance for H3 subtype AIVs infections and to develop a simple, rapid and sensitive detection method.

Loop-mediated isothermal amplification (LAMP) is a new nucleic acid amplification method that uses a set of six primers that target the eight site of a conserved gene, yielding a large white pyrophosphate ion by-product. The result can be viewed by adding a fluorescent metal indicator to the reaction mix before amplification [11-13]. Another significant advantage is that the assay can be performed in a water bath within 30-60 minutes. Therefore, LAMP is simple, rapid and specific enough to detect pathogenic microorganisms in field conditions, and currently the assay is being used widely for the detection of various viral pathogens [14-22].

The purpose of the study reported here was specifically to enhance the surveillance of the H3 subtype of AIVs by developing and optimizing the LAMP technique for the rapid detection of H3 subtype AIVs in live birds.

Methods

Viral strains and DNA/RNA extraction

The reference strains of AIVs, clinical samples and other avian respiratory pathogens used in this study are listed in Table 1 and 2. The Genomic DNA/RNA was extracted from 200 ul samples using a DNA/RNA Miniprep Kit (Axygen Biosciences, Hangzhou, China), according to the protocol suggested by the manufacturer. The DNA and RNA were eluted with 40 ul elution buffer and were stored at -70°C immediately, until use.

Design of primers for the RT-LAMP assay

According to the sequences of the H3 subtype AIVs' hemagglutinin (HA) gene available in GenBank (accession no. JN003630) and the sequences of viruses isolated in China, several primer sets of RT-LAMP assays were

designed using the LAMP primer design software Primer Explorer V4 http://primerexplorer.jp/elamp4.0.0/index. html. Finally, an optimal set was chosen after many comparable experiments to ensure the highest sensitivity and specificity of the RT-LAMP assay. The RT-LAMP primer set comprising two outer primers (forward primer F3 and backward primer B3), two inner primers (forward inner primer FIP and backward inner primer BIP), and two loop primers (forward loop primer LF and backward loop primer LB) recognized eight sites on the target sequence specific to the HA gene. The details for the primers are shown in Table 3 and Figure 1.

Optimization of the RT-LAMP conditions

The RT-LAMP assay was carried out in a conventional water bath with 25 µl of reaction mixture (Table 4). The concentration of reactions component comprised primers (synthesized by Invitrogen), MgSO₄ (Sigma), Betaine (Sigma), deoxynucleoside triphosphate (dNTP), RNA template, RevertAid™ M-MuLV Reverse Transcriptase (MBI Fermentas) and Bst DNA polymerase (New England Biolabs) were optimized. To visualize the reaction, 25 µmol/L Calcein (International Laboratory, USA) and 0.5 mmol/L MnCl₂ (International Laboratory, USA) had been added previously to the reaction mixture, as suggested by the related reports [13]. The amplification reaction was performed in a water bath at 59°C, 60°C, 61°C, 62°C, 63°C, 64°C, and 65°C for 30, 45, and 60 min respectively to find the optimal temperature and time. Then the reaction was terminated by heating at 80°C for 5 min. All of the experiments were repeated three times.

RT-PCR

Prior to this study, there was no official primer or procedure for RT-PCR to detect H3 subtyping of AIVs. The outer primers of RT-LAMP (forward primer F3 and backward primer B3) that were specific to target sequence and had high priming efficiency were applied in RT-PCR assay of H3 subtype AIVs. The reaction condition of RT-PCR was optimized to get the highest sensitivity of amplification. Finally, the RT-PCR assay was carried out in a 50 ul reaction volume containing 10 mM of each dNTP, 5 ul of 5 × RT Reaction Buffer, 200 U of RevertAidTM M-MuLV Reverse Transcriptase, 5 ul of 10 × Taq Buffer, 5 U of Taq DNA polymerase (MBI Fermentas), 10 uM each of B3 and F3 primers (Table 3), and 1ul of extracted RNA, DEPC-treated water up to 50 ul.

The reaction was performed in a Thermal Cycler (BIO-RAD) at 42°C for 60 min and at 94°C for 5 min; followed by 30 cycles of 94°C for 40 s, 55°C for 30 s, and 72°C for 30 s; and a final extension for 10 min at 72°C. Then the amplified products were analyzed using 1% agarose gel electrophoresis.

Table 1 Clinic cloacal swab samples used to evaluate the feasibility of RT-LAMP assay

Number	Sources and identification	Virus isolation	RT-PCR	RT-LAMF
1	A/Duck/Guangxi/N42/2009(H3N2)	+	+	+
2	A/Duck/Guangxi/M20/2009(H3N2)	+	+	+
3	A/Duck/Guangxi/04D3/2009(H3)	+	+	+
4	A/Duck/Guangxi/04D8/2009(H3)	+	+	+
5	A/Duck/Guangxi/017D5/2009(H3)	+	+	+
6	A/Duck/Guangxi/LZD11/2009(H3)	+	+	+
7	A/Duck/Guangxi/LZD15/2009(H3)	+	+	+
8	A/Duck/Guangxi/LZD23/2009(H3)	+	+	+
9	A/Duck/Guangxi/LZD27/2009(H3)	+	+	+
10	A/Duck/Guangxi/LZD28/2009(H3)	+	+	+
11	A/Duck/Guangxi/042D16/2009(H3)	+	+	+
12	A/Duck/Guangxi/046D6/2009(H3)	+	+	+
13	A/Duck/Guangxi/046D15/2009(H3)	+	+	+
14	A/Duck/Guangxi/048D617/2010(H3)	+	-	+
15	A/Duck/Guangxi/047D20/2010(H3)	+	+	+
16	A/Duck/Guangxi/047D15/2010(H3)	+	+	+
17	A/Duck/Guangxi/047D16/2010(H3)	+	+	+
18	A/Duck/Guangxi/049D7/2010(H3)	+	+	+
19	A/Duck/Guangxi/054D2/2010(H3)	+	+	+
20	A/Duck/Guangxi/054D3/2010(H3)	+	+	+
21	A/Duck/Guangxi/054D4/2010(H3)	+	+	+
22	A/Duck/Guangxi/054D7/2010(H3)	+	+	+
23	A/Duck/Guangxi/054D16/2010(H3)	+	+	+
24	A/Duck/Guangxi/054D19/2010(H3)	+	+	+
25	A/Duck/Guangxi/054D20/2010(H3)	+	+	+
26	A/Chicken/Guangxi/055C2/2010(H3)	+	+	+
27	A/Duck/Guangxi/057D19/2010(H3)	+	+	+
28	A/Duck/Guangxi/057D6/2010(H3)	+	+	+
29	A/Duck/Guangxi/WYD1/2010(H3)	+	+	+
30	A/Duck/Guangxi/066D16/2010(H3)	+	+	+
31	A/Duck/Guangxi/068D8/2010(H3)	+	+	+
32	A/Chicken/Guangxi/068C3/2010(H3)	+	+	+
33	A/Duck/Guangxi/070D8/2010(H3)	+	-	+
34	A/Duck/Guangxi/070D14/2010(H3)	+	_	+
35	A/Duck/Guangxi/072D5/2010(H3)	+	+	
36	A/Duck/Guangxi/072D4/2010(H3)	+	+	+
37	A/Duck/Guangxi/072D2/2010(H3)	+	+	+
38	A/Duck/Guangxi/072D7/2010(H3)	+	+	+
39	A/Chicken/Guangxi/LF/2007	т -	т -	т -
40	A/Duck/Guangxi/RX/2009	-	_	_
41	A/Chicken/Guangxi/015C7/2009	-	-	-
		-	-	-
42	A/Duck/Guangxi/015D7/2009	-	-	-
43	A/Duck/Guangxi/011D3/2009	-	-	-
44	A/Duck/Guangxi/011D10/2009	-	-	-
45	A/Duck/Guangxi/016D10/2009	-	-	-
46	A/Duck/Guangxi/016D8/2009	-	-	-
47	A/Duck/Guangxi/022D1/2009	-	-	-
48	A/Chicken/Guangxi/037C10/2009	-	-	-
49	A/Duck/Guangxi/LAD9/2009	-	-	-
50	A/francolin/Nanning/018B-3/2010	-	-	-
51	A/francolin/Nanning/020B-7/2010	-	-	-

Table 1 Clinic cloacal swab samples used to evaluate the feasibility of RT-LAMP assay (Continued)

	• · · · · · · · · · · · · · · · · · · ·	•	, , , , , , , , , , , , , , , , , , , ,	
52	A/francolin/Nanning/022B12/2010	-	-	-
53	A/francolin/Nanning/022B13/2010	-	-	-
54	A/francolin/Nanning/022B7/2010	-	-	-
55	A/Chicken/Guangxi/DXC4/2010	-	-	-
56	A/Chicken/Guangxi/066C10/2010	-	-	-
57	A/Chicken/Guangxi/067C4/2010	-	-	-
58	A/Chicken/Guangxi/067C1/2010	-	-	-
59	A/Duck/Guangxi/068D18/2010	-	-	-
60	A/Duck/Guangxi/01D5/2009	-	-	-
61	A/Duck/Guangxi/02D6/2009	-	-	-
62	A/Duck/Guangxi/02D7/2009	-	-	-
63	A/Duck/Guangxi/04D5/2009	-	-	-
64	A/Duck/Guangxi/010D9/2009	-	-	-
65	A/Duck/Guangxi/011C5/2009	-	-	-
66	A/Duck/Guangxi/011C17/2009	-	-	-
67	A/Duck/Guangxi/011D5/2009	-	-	-
68	A/Duck/Guangxi/014D10/2009	-	-	-
69	A/Duck/Guangxi/014D15/2009	-	-	-
70	A/Duck/Guangxi/014D18/2009	-	-	-
71	A/Duck/Guangxi/016D7/2009	-	-	-
72	A/Duck/Guangxi/017C2/2009	-	-	-
73	A/Duck/Guangxi/017C12/2009	-	-	-
74	A/Duck/Guangxi/019C13/2009	-	-	-
75	A/Duck/Guangxi/026D10/2009	-	-	-
76	A/Duck/Guangxi/024C2/2009	-	-	-
77	A/Duck/Guangxi/024D2/2009	-	-	-
78	A/Duck/Guangxi/028D6/2009	-	-	-
79	A/Goose/Guangxi/027G10/2009	-	-	-
80	A/Duck/Guangxi/025D6/2009	-	-	-
81	A/Duck/Guangxi/026D21/2009	-	-	-
82	A/Duck/Guangxi/027D2/2009	-	-	-
83	A/Duck/Guangxi/033C1/2009	-	-	-
84	A/Duck/Guangxi/033C2/2009	-	-	-
85	A/Duck/Guangxi/033C3/2009	-	-	-
86	A/Duck/Guangxi/033C4/2009	-	-	-
87	A/Duck/Guangxi/033C5/2009	-	-	-
88	A/Duck/Guangxi/033C7/2009	-	-	-
89	A/Duck/Guangxi/033C9/2009	-	-	-
90	A/Duck/Guangxi/033C10/2009	-	-	-
91	A/Duck/Guangxi/031G2/2009	-	-	-
92	A/Duck/Guangxi/031G11/2009	-	-	-
93	A/Duck/Guangxi/031G12/2009	-	-	-
94	A/Duck/Guangxi/042D1/2009	-	-	-
95	A/Duck/Guangxi/042D3/2009	-	-	-
96	A/Duck/Guangxi/014D4/2009	-	-	-
97	A/Duck/Guangxi/042D9/2009	-	-	-
98	A/Duck/Guangxi/042D10/2009	-	-	-
99	A/Duck/Guangxi/042D19/2009	-	-	-
100	A/Duck/Guangxi/044D6/2009	-	-	-
101	A/Duck/Guangxi/044D7/2009	-	-	-
102	A/Duck/Guangxi/044D8/2009	-	-	-
103	A/Duck/Guangxi/044D10/2009	-	-	-

Table 1 Clinic cloacal swab samples used to evaluate the feasibility of RT-LAMP assay (Continued)

		•	, , , , , , , , , , , , , , , , , , , ,	
104	A/Duck/Guangxi/044D12/2009	-	-	-
105	A/Duck/Guangxi/044D17/2009	-	-	-
106	A/Duck/Guangxi/044C4/2009	-	-	-
107	A/Duck/Guangxi/044C7/2009	-	-	-
108	A/Duck/Guangxi/041D10/2009	-	-	-
109	A/Duck/Guangxi/041D11/2009	-	-	-
110	A/Duck/Guangxi/041D13/2009	-	-	-
111	A/Duck/Guangxi/048D15/2010	-	-	-
112	A/Duck/Guangxi/048D18/2010	-	-	-
113	A/Duck/Guangxi/048D2/2010	-	-	-
114	A/Duck/Guangxi/048D16/2010	-	-	-
115	A/Duck/Guangxi/PXD7/2010	-	-	-
116	A/Duck/Guangxi/PXD5/2010	-	-	-
117	A/Duck/Guangxi/PXD15/2010	-	-	-
118	A/Duck/Guangxi/052D16/2010	-	-	-
119	A/Duck/Guangxi/052D14/2010	-	-	-
120	A/Duck/Guangxi/052D8/2010	-	-	-
121	A/Duck/Guangxi/052D2/2010	-	-	-
122	A/Duck/Guangxi/052D1/2010	-	-	-
123	A/Duck/Guangxi/049D1/2010	-	-	-
124	A/Duck/Guangxi/049D2/2010	-	-	-
125	A/Duck/Guangxi/049D32010	-	-	-
126	A/Duck/Guangxi/049D4/2010	-	-	-
127	A/Duck/Guangxi/049D5/2010	-	-	-
128	A/Duck/Guangxi/049D8/2010	-	-	-
129	A/Duck/Guangxi/049D12/2010	-	-	-
130	A/Duck/Guangxi/049D15/2010	-	-	-
131	A/Duck/Guangxi/049D17/2010	-	-	-
132	A/Duck/Guangxi/049D18/2010	-	-	-
133	A/Duck/Guangxi/055D4/2010	-	-	-
134	A/Duck/Guangxi/055D5/2010	-	-	-
135	A/Duck/Guangxi/055D6/2010	-	-	-
136	A/Duck/Guangxi/055D7/2010	-	-	-
137	A/Duck/Guangxi/055D13/2010	-	-	-
138	A/Duck/Guangxi/055D20/2010	-	-	-
139	A/Duck/Guangxi/058D1/2010	-	-	-
140	A/Duck/Guangxi/058D6/2010	-	-	-
141	A/Duck/Guangxi/058D8/2010	-	-	-
142	A/Duck/Guangxi/058D12/2010	-	-	-
143	A/Duck/Guangxi/058D13/2010	-	-	-
144	A/Duck/Guangxi/058D15/2010	-	-	-
145	A/Duck/Guangxi/062D2/2010	-	-	-
146	A/Duck/Guangxi/062D3/2010	-	-	-
147	A/Duck/Guangxi/062D4/2010	-	-	-
148	A/Duck/Guangxi/062D5/2010	-	-	-
149	A/Duck/Guangxi/062D7/2010	-	-	-
150	A/Duck/Guangxi/062D8/2010	-	-	-
151	A/Duck/Guangxi/062D9/2010	-	-	-
152	A/Duck/Guangxi/062D10/2010	-	-	-
153	A/Duck/Guangxi/062D11/2010	-	-	-
154	A/Duck/Guangxi/062D12/2010	-	-	-
155	A/Duck/Guangxi/062D13/2010	-	-	-

Table 1 Clinic cloacal swab samples used to evaluate the feasibility of RT-LAMP assay (Continued)

156	A/Duck/Guangxi/062D14/2010	-	-	-
157	A/Duck/Guangxi/062D15/2010	-	-	-
158	A/Duck/Guangxi/062D17/2010	-	-	-
159	A/Duck/Guangxi/062D18/2010	-	-	-
160	A/Duck/Guangxi/062D19/2010	-	-	-
161	A/Duck/Guangxi/066D2/2010	-	-	-
162	A/Duck/Guangxi/066D5/2010	-	-	-
163	A/Duck/Guangxi/066D7/2010	-	-	-
164	A/Duck/Guangxi/066D10/2010	-	-	-
165	A/Duck/Guangxi/066D20/2010	-	-	-
166	A/Duck/Guangxi/BSD20/2010	-	-	-
167	A/Duck/Guangxi/BSD12/2010	-	-	-
168	A/Duck/Guangxi/BSD18/2010	-	-	-
169	A/Duck/Guangxi/BSD23/2010	-	-	-
170	A/Duck/Guangxi/BSD38/2010	-	-	-
171	A/Duck/Guangxi/070D5/2010	-	-	-
172	A/Duck/Guangxi/072D14/2010	-	-	-
173	A/Duck/Guangxi/072D16/2010	-	-	-
174	A/Duck/Guangxi/072D13/2010	-	-	-
175	A/Duck/Guangxi/072D19/2010	-	-	-
176	A/Duck/Guangxi/072D1/2010	-	-	-

Table 2 Virus strains used for specificity of H3-RT-LAMP assay

Number	Virus strain	H3-RT-LAMP
1	A/Duck/Guangxi/030D/2009(H1N1)	-
2	A/Mallard/Alberta/77(H2N3)	-
3	A/Mallard/Alberta/85(H3N6)	+
4	A/Duck/Guangxi/N42/2009(H3N2)	+
5	A/Duck/Guangxi/027D/2009 (H4)	-
6	A/Turkey/GA/209092/02(H5N2)	-
7	A/Turkey/CA/35621/84(H5N3)	-
8	A/waterfowl/GA/269452-56/03(H5N7)	-
9	A/Turkey/MA/40550/87-Bel42(H5N1)	-
10	A/Turkey/WI/68(H5N9)	-
11	A/Turkey/Ontario/63(H6N8)	-
12	A/Chicken/NY/273874/03(H7N2)	-
13	A/Turkey/Ontario/6118/67(H8N4)	-
14	A/Duck/Guangxi/RX/09(H9N2)	-
15	A/Turkey/MN/24838-590/79(H10N7)	-
16	A/Chicken/Guangxi/43C/09(H11)	-
17	A/Duck/Alberta/60/76(H12N5)	-
18	A/Gull/MD/704/77(H13N6)	-
19	A/Duck/Australia/341/83(H15N8)	-
20	Newcastle disease virus (Lasota)	-
21	Infectious bronchitis virus (M41)	-
22	Infectious Laryngotracheitis virus	-
23	Mycoplasma gallisepticum (S6)	-

Analysis of RT-LAMP assay products

For the visual inspection of the products of the RT-LAMP assay, white magnesium pyrophosphate precipitations in the tube were centrifuged at 8,000 X g for 5 min in a Biofuge centrifuge (Primo R, HERAEUS). Meanwhile, color changes of the reaction mixture were observed directly under daylight and UV light.

Agarose gel electrophoresis analysis and DNA sequencing Five μ l of the RT-LAMP assay product was analyzed using 1.5% agarose gel electrophoresis. To test the specificity of the RT-LAMP assay product, the RT-LAMP amplicons were first digested with EcoRI and EcoRV restriction enzymes using the suggested protocol and then analyzed with 1.5% agarose gel electrophoresis. The products digested by restriction enzymes were purified using an AxyPrep TM DNA Gel Extraction Kit and were sequenced by Invitrogen Company.

Sensitivity and specificity of RT-LAMP assay

The detection limit of RT-LAMP was determined by testing serial 10-fold dilutions of total RNA of A/Duck/ Guangxi/N42/2009 (H3N2) respectively, and the same concentration samples were comparatively detected by RT-PCR. All of the experiments were repeated three times. The concentration range of total RNA in the diluted sample was 1ng/tube to 1fg/tube, measured by ultraviolet spectrophotometer (Beckman UV-800).

Table 3 Sequences of primers designed for RT-LAMP assay

Primer name	Sequence(5'-3') ^a	Genome position
FIP	GGATTATAGTCTGTTGGCTTCTCC-GAATTC-TGTATGTTCAAGCCTCA	658-720
BIP	GGCCAATCTGGCAGAATAAGC-GATATC-CCATTACTATTGATTACCAGTACGT	750-823
F3	CACAAATCAAGAACAA	635-654
В3	CCGAGGAGCGATTAGGTT	825-842
LF	GGTAGAGACTGTGACTCT	678-695
LB	ATCTATTGGACAGTAGTCAAACCTG	771-795

^a The restriction enzyme site sequences of *EcoR I* and *EcoR V* were added in FIP and BIP primer respectively. Primers of F3, B3 were applied to the RT-PCR assay of H3 subtype AIVs.

To evaluate the specificity of the primer set used for the RT-LAMP assay, DNA/RNA extracted from different subtype AIVs and from Newcastle disease virus (NDV), infectious bronchitis virus (IBV), infectious laryngotracheitis virus (ILTV), *Mycoplasma gallisepticum* (MG) were detected by RT-LAMP assay. The amplification results were analyzed using 1.5% agarose gel electrophoresis and the color changes of reaction mixture were inspected under daylight and UV light.

Detection of clinical specimens by RT-LAMP assay

A total of 176 cloacal swabs were collected from poultry at LBMs. The clinical specimens were prepared in a viral transport medium, which was made up of 0.05 M phosphate buffered saline (PBS) containing antibiotics of penicillin (10000 units/ml), streptomycin (10 mg/ml), gentamycin (10 mg/ml), kanamycin (10 mg/ml) and 5% (v/v) fetal bovine serum. Cloacal swabs samples were injected into 9-to 11-day-old embryonated specific-pathogen-free (SPF) chicken eggs as previously described [5]. Allantoid fluids were recovered 48-72 hrs after incubation for virus detection and titration as described [23]. All clinical specimens were tested by RT-LAMP, RT-PCR, and virus isolation, respectively.

632 F3 AAGCACAAATCAAGAA	F2 CAAACCAATCTGTATGTTCAAG	LFc CCTCAGGAAGAGTC
F1c ACAGTCTCTACCAGGAC	GAAGCCAACAGACTATAATCCC	AAACATTGGATCTAG
Blc	LB CAATCTGGCAGAATAAGCATCTA	
B2c	ВЗс	
AACCIGGGGACGIACIO	GGTAATCAATAGTAATGG <mark>A</mark> AACG	CIAAICGCICCICGG
GGCTATTTCAAAACACG 900	GTATTGGGAAAAGCTCAATAATG	AGATCAGATGCACCT
ATTG		

Figure 1 Positions of RT-LAMP primers on HA gene of H3 subtype AIVs. Locations of primers binding sequences are underlined and boxed. The referenced sequence A/Duck/GuangXi/ N42/2009(H3N2) can be obtained in GenBank (accession number JN003630).

Results

The optimal protocol of RT-LAMP assay and inspection of products

The RT-LAMP reaction was optimized in a 25 ul total reaction mixture and performed in a conventional waterbath for 45 min at 63°C and then for 5 min at 80°C to terminate the reaction. The details of optimize reaction solution of RT-LAMP assay are shown in Table 4.

The positive results of RT-LAMP assay showed typical ladder pattern by 1.5% agarose gel electrophoresis (Figure 2-A). Furthermore, a large amount of DNA was synthesized, yielding a large white magnesium pyrophosphate by-product, so the white magnesium pyrophosphate on the bottom of the tube could be inspected after centrifugation. Alternatively, a color change in the reaction solution could be seen with the naked eye: the solution changed from orange to green for a positive reaction and remained orange for the negative reaction (Figure 2-C). In addition, the positive sample showed a strong green fluorescence under UV light (Figure 2-D). The products of the RT-LAMP assay digested by *EcoR* I

Table 4 The reaction system of RT-LAMP assay

Reaction cor	nponent	Reaction volume of	
		H3-RT-LAMP assay	
$10 \times \text{Thermopol Buffer}$		2.5 μL	
MgSC)4	5 mmol/L	
Betain	e	1 mmol/L	
F3		0.2 μmol/L	
	В3	0.2 μmol/L	
primer	FIP	1.6 µmol/L	
	BIP	1.6 µmol/L	
	LF	0.8 µmol/L	
	LB	0.8 µmol/L	
dNTP		1.4 mmol/L	
Calcein		25 μmol/L	
MnCl ₂		0.5 mmol/L	
M-MuLV RTase		200U	
Bst DNA pol	ymerase	8U	
Template	RNA	2 μL	
Nuclease-free water		Up to 25 μL	

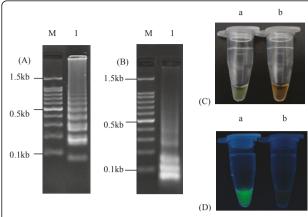


Figure 2 Detection of H3 subtype AIVs by RT-LAMP assay. (A) Products of RT-LAMP assay confirmed by 1.5% agarose gel electrophoresis. (B) Products of RT-LAMP assay digested by *EcoR* I and *EcoR* V restriction enzyme. (C) Result of RT-LAMP assay under daylight. (D) Result of RT-LAMP assay under UV light. M:100 bp. a: positive sample. b: negative sample.

and *EcoR* V restriction enzymes displayed predicted fragments (Figure 2-B). These fragments were purified and then confirmed by sequencing analysis (date not shown).

Specificity and sensitivity of RT-LAMP assay for detection of H3 Subtype AIVs

The viral DNA/RNA extracted from the different subtype AIVs and NDV, IBV, ILTV, MG were detected using the RT-LAMP assay to evaluate the specificity. The results of the negative sample showed no typical ladder pattern by 1.5% agarose gel electrophoresis and also had no color change detectable under daylight or UV light, which indicated that RT-LAMP assay had no cross reaction with other viruses (Table 2). The detection limit of RT-LAMP assay was 0.1 pg total RNA of virus, which was 100-fold higher than that of RT-PCR. The results of the visual detection of sensitivity tests are shown in Figure (3A, B and 3C).

Evaluation of RT-LAMP assay with clinical Samples

A total of 176 random cloacal swab samples were collected from poultry at various LBMs and were tested by RT-LAMP, RT-PCR, and virus isolation respectively (Table 1). The results of virus isolation showed that there were 38 positive samples of H3 subtype AIVs among the 176 cloacal swab samples, which was consistent with that of RT-LAMP assay; however, three positive samples were missed by RT-PCR. The results of statistical analysis showed that comparative detection of H3 subtype AIVs from cloacal swab samples by RT-LAMP was not statistically significant from RT-PCR and virus isolation (Table 5).

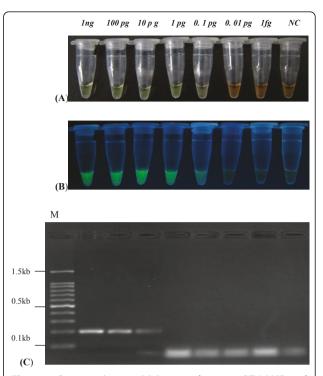


Figure 3 Comparative sensitivity tests between RT-LAMP and RT-PCR assays for detection of H3 subtype AIVs (the detection range of virus total RNA was 1ng/tube to 1fg/tube). (A) The result of sensitivity test for RT-LAMP assay under daylight. (B) The result of sensitivity test for RT-LAMP assay under UV light. (C) The result of sensitivity test for RT-PCR. M: 100 bp. NC: negative control.

Discussion

The H3 subtype AIVs can provide genes for human influenza virus through gene reassortment, which raises great concerns in terms of its potential threat to human health [6]. Consequently, the development of a rapid, simple, sensitive detection method for H3 subtype AIVs is required. So far, there are several PCR-based methods being used to detect AIVs, but they all need precision instruments to amplify the nucleic acid of target sample; therefore, they can't be applied in field conditions [24,25]. The loop-mediated isothermal amplification (LAMP) assay is a nucleic acid amplification method

Table 5 Comparative detection of cloacal swab samples by virus isolation, RT-PCR and RT-LAMP

Results	Virus isolation ^a	RT-PCR ^b	RT-LAMP ^c
Positive	38	35	38
Negative	138	141	138
Total	176	176	176
Sig.(P value)	a-b:0.696	b-a:0.696	c-a:1.000
	a-c:1.000	b-c:0.696	c-b:0.696

^aVirus isolation, ^bRT-PCR, ^cRT-LAMP.

The significant differences among the three detection methods were analyzed by SPSS software. The default significance level is 0.05.

developed by Notomi et al. that does not require any specialized equipment [11] and can be performed in a water bath or a heating block at an isothermal temperature between 60°C to 65°C within 30 to 60 min. In contrast, the conventional PCR method takes at least one hour.

The LAMP assay relies on autocycling strand displacement DNA synthesis performed by Bst DNA polymerase with a high degree of strand displacement activity, which leads to an excellent sensitivity. The primer set of LAMP assay comprise two outer primers (forward primer F3 and backward primer B3), two inner primers (forward inner primer FIP and backward inner primer BIP), and two loop primers (forward loop primer LF and backward loop primer LB) and it recognizes eight sites on the target sequence specific to HA gene, so it's specificity is very high. The viral RNA can be detected by the one-step RT-LAMP assay. When this is performed with reverse transcriptase and LAMP reaction mixture, the reverse transcription and DNA amplification can be accomplished at a constant temperature within 60 min.

During amplification, a large amount of white magnesium pyrophosphate is produced, which can be inspected directly with the naked eye. Alternatively, the result of the RT-LAMP assay can be visualized by adding fluorescence reagent (Calcein and MnCl₂) into the reaction mixture to observe the change of the reaction color under day light or UV light [11-13]. Before the amplification reaction, calcein is combined with manganese ion to get the quenching effect, so the reaction solution is orange. Along with LAMP reaction processing, pyrophosphate ions remove manganese ions from calcein, resulting in greater fluorescence, which indicates the presence of the target gene [12].

In the present study, we first developed a rapid and sensitive RT-LAMP assay to visually detect H3 subtype AIVs. According to the sequences of the HA gene of H3 subtype AIVs available in GenBank and the sequences of H3 subtype AIVs isolated in China, we designed several sets of primers and finally chose the optimal set after many comparative experiments to ensure the high specificity and sensitivity of the RT-LAMP assay.

The proportions of the reagents in the reaction mixture were optimized to develop a stable RT-LAMP assay with high sensitivity. The detection limit of the RT-LAMP assay was 0.1 pg total RNA of virus, which was 100-fold higher than that of RT-PCR. The results of the specificity test showed that the assay had no cross reaction with other subtype AIVs and avian respiratory pathogens. Furthermore, a total of 176 cloacal swab samples collected from LBMs were tested using RT-LAMP, RT-PCR, and virus isolation respectively. The results showed that the clinical sensitivity of the RT-LAMP assay was consistent with virus isolation.

Although the RT-LAMP assay has many advantages over the similar nucleic acid amplification method, there is still a problem to note. Because of the high sensitivity of the RT-LAMP assay, a micro amount of RNA contamination in reagents, environment and instruments, such as the pipette, can result in a false positive. Moreover, the RT-LAMP assay has great amplification efficiency so the reaction product can form aerosol to contaminate the surroundings when opening the tube. To avoid the contamination in this study, the fluorescence reagent (Calcein and MnCl₂) were added to RT-LAMP reaction mixture before amplification in order to inspect the result of RT-LAMP assay directly with the naked eye.

Conclusions

In this study, the established RT-LAMP assay with high sensitivity was performed in a water bath within only 50 min, and the amplification results were visualized by adding fluorescence reagent. In summary, the newly developed assay can be used as one important method for detecting H3 subtype AIVs in field conditions with no need for specialized equipment.

Acknowledgements

This work was supported by grants from Bai Qian Wan Talents Engineering Foundation (945200603) and the Guangxi Science and Technology Bureau (10100014-5).

Author details

¹Department of Biotechnology, Guangxi Veterinary Research Institute, 51 You Ai Road, Nanning, Guangxi 530001, China. ²Department of Pathobiology and Veterinary Science, University of Connecticut, 61 North Eagleville Road Storrs, CT 06269-3089, USA. ³Guangxi Key Laboratory of Subtropical Bioresources Conservation and Utilization, College of Life Science and Technology, Guangxi University, 100 Daxue Road, Nanning, Guangxi 530004, China.

Authors' contributions

YP, ZXX and MIK designed the experiments. QF and LJX prepared the RNA samples. YP designed the primers and optimized conditions of the RT-LAMP assay. YP, JBL, YSP, XWD and ZQX carried out the experiments shown in Figure 2 and 3 and in Table 1, 4, 5. JXF performed the data analysis. YP and MIK wrote the manuscript. All authors read and approved the final manuscript.

Competing interests

The authors declare that they have no competing interests.

Received: 1 April 2011 Accepted: 5 July 2011 Published: 5 July 2011

References

- Webster RG, Bean WJ, Gorman OT, Chambers TM, Kawaoka Y: Evolution and ecology of influenza A viruses. Microbiol Rev 1992, 56(1):152-179.
- Horimoto T, Kawaoka Y: Pandemic threat posed by avian influenza A viruses. Clin Microbiol Rev 2001, 14(1):129-149.
- Fouchier RAM, Munster V, Wallensten A, Bestebroer TM, Herfst S, Smith D, Rimmelzwaan GF, Olsen B, Osterhaus ADME: Characterization of a novel influenza A virus hemagglutinin subtype (H16) obtained from blackheaded gulls. J Virol 2005, 79(5):2814-2822.
- Alexander DJ: A review of avian influenza in different bird species. Vet Microbiol 2000, 74(1-2):3-13.
- Edwards S: OIE laboratory standards for avian influenza. Dev Bio (Basel) 2006, 124:159-162.

- Scholtissek C, Rohde W, Von-Hoyningen V, Rott R: On the origin of the human influenza virus subtype H2N2 and H3N2. Virology 1978, 87(1):13-20.
- Karasin Al, Carman S, Olsen CW: Identification of human H1N2 and human-swine reassortant H1N2 and H1N1 influenza A viruses among pigs in Ontario, Canada (2003 to 2005). J Clin Microbiol 2006, 44(3):1123-1126.
- Peng Y, Zhang W, Xue F, Wang WB, Li YF, Meng CC, Zhang WJ, Liu XF: Etiological examination on the low pathogenicity avian influenza viruses with different HA subtypes from poultry isolated in eastern China from 2006 to 2008. Chin J Zoonoses 2009, 25(2):119-121.
- Song MS, Oh TK, Moon HJ, Yoo DW, Lee EH, Lee JS, Kim CJ, Yoo GJ, Kim H, Choi YK: Ecology of H3 avian influenza viruses in Korea and assessment of their pathogenic potentials. J Gen Virol 2008, 89:949-957.
- Campitelli L, Fabiani C, Puzelli S, Fioretti A, Foni E, De Marco MA, Krauss S, Webster RG, Donatelli I: H3N2 influenza viruses from domestic chickens in Italy:an increasing role for chickens in the ecology of influenza? J Gen Virol 2002, 83:413-420.
- Notomi T, Okayama H, Masubuchi H, Yonekawa T, Watanabe K, Amino N, Hase T: Loop-mediated isothermal amplification of DNA. Nucleic Acids Res 2000, 28(12):e63.
- Mori Y, Nagamine K, Tomita N, Notomi T: Detection of loop-mediated isothermal amplification reaction by turbidity derived from magnesium pyrophosphate formation. Biochem Biophys Res Commun 2001, 280(1):150-154
- Tomita N, Mori Y, Kanda H, Notomi T: Loop-mediated isothermal amplification (LAMP) of gene sequences and simple visual detection of products. Nature protocol 2008, 3(5):877-882.
- Rovira A, Abrahante J, Murtaugh M, Mun oz-Zanzi C: Reverse transcription loop-mediated isothermal amplification for the detection of Porcine reproductive and respiratory syndrome virus. J Vet Diagn Invest 2009, 21(3):350-354.
- Pham HM, Nakajima C, Ohashi K, Onuma M: Loop-Mediated Isothermal Amplification for Rapid Detection of Newcastle Disease Virus. J Clin Microbiol 2005, 43(4):1646-1650.
- Xu JT, Zhang ZM, Yin YB, Cui SJ, Xu SZ, Guo YY, Li JD, Wang JL, Liu XC, Han LM: Development of reverse-transcription loop-mediated isothermal amplification for the detection of infectious bursal disease virus. J Virol Methods 2009, 162:267-271.
- Poon LLM, Leung CSW, Chan KH, Lee JHC, Yuen KY, Guan Y, Peiris JSM: Detection of human influenza A viruses by loop-mediated isothermal amplification. J Clin Microbiol 2005, 43(1):427-430.
- Ito M, Watanabe M, Nakagawa N, Okuno Y: Rapid detection and typing of influenza A and B by loop-mediated isothermal amplification: comparison with immunochromatography and virus isolation. J Virol Methods 2006. 135:272-275.
- Imai M, Ninomiya A, Minekawa H, Notomi T, Ishizaki T, Tashiro M, Odagiri T: Development of H5-RT-LAMP (loop-mediated isothermal amplification) system for rapid diagnosis of H5 avian influenza virus infection. Vaccine 2006, 24:6679-6682.
- Imai M, Ninomiya A, Minekawa H, Notomi T, Ishizaki T, Tu PV, Tien NTK, Tashiro M, Odagiri T: Rapid diagnosis of H5N1 avian influenza virus infection by newly developed influenza H5 hemagglutinin gene-specific loop-mediated isothermal amplification method. J Virol Methods 2007, 141:173-180.
- Ma XJ, Shu YL, Nie K, Qin M, Wang DY, Gao RB, Wang M, Wen LY, Han F, Zhou SM, et al: Visual detection of pandemic influenza A H1N1 Virus 2009 by reverse-transcription loop-mediated isothermal amplification with hydroxynaphthol blue dye. J Virol Methods 2010, 167:214-217.
- Kubo T, Agoh M, Q-Mai Le, Fukushima K, Nishimura H, Yamaguchi A, Hirano M, Yoshikawa A, Hasebe Fi, Kohno S, et al: Development of a reverse transcription-loop-mediated isothermal amplification assay for detection of pandemic (H1N1) 2009 virus as a novel molecular method for diagnosis of pandemic influenza in resource-limited settings. J Clin Microbiol 2010, 48(3):728-735.
- 23. Swayne ED, Glisson RJ, Jackwood MW, Pearson EJ, Reed MW: A laboratory manual for the isolation and identification of avian pathogens. American Association of Avian Pathologist University of Pennsylvania 1998.
- 24. Das A, Spackman E, Senne D, Pedersen J, Suarez DL: Development of an internal positive control for rapid diagnosis of avian influenza virus

- infections by real-time reverse transcription-PCR with lyophilized reagents. *J Clin Microbiol* 2006, **44(9)**:3065-3073.
- Tsukamoto K, Ashizawa H, Nakanishi K, Kaji N, Suzuki K, Okamatsu M, Yamaguchi S, Mase M: Subtyping of avian influenza viruses H1 to H15 on the basis of hemagglutinin genes by PCR assay and molecular determination of pathogenic potential. J Clin Microbiol 2008, 46(9):3048-3055.

doi:10.1186/1743-422X-8-337

Cite this article as: Peng et al.: Visual detection of H3 subtype avian influenza viruses by reverse transcription loop-mediated isothermal amplification assay. Virology Journal 2011 8:337.

Submit your next manuscript to BioMed Central and take full advantage of:

- Convenient online submission
- Thorough peer review
- No space constraints or color figure charges
- Immediate publication on acceptance
- Inclusion in PubMed, CAS, Scopus and Google Scholar
- Research which is freely available for redistribution

Submit your manuscript at www.biomedcentral.com/submit

