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Evaluation of clinical applicability of reverse transcription-loop-mediated isothermal amplification assay for detection and subtyping of Influenza A viruses



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ARTICLE INFO	ABSTRACT
Keywords: Influenza A viruses Molecular detection Pandemic influenza RT-LAMP	 Background: Influenza A viruses (IAVs) have always remain a serious concern for the global economy and public health. A rapid, specific and sensitive detection method is always needed to control the influenza in its early stages by timely intervention of therapy and early clinical management. Objectives: To develop RT-LAMP assays for detection of influenza A viruses, their further subtyping into seasonal (H1N1, H3N2) and novel pandemic H1N1 viruses and to evaluate clinical applicability of optimized RT-LAMP assays on patients' samples. Study design: In this study, we optimized RT-LAMP assay to detect IAVs by using primers against matrix gene and subtyping of IAVs was done by using primers against hemagglutinin gene. Optimized RT-LAMP assays were applied on clinical samples from patients having influenza like illness and results were compared with conventional one-step RT-PCR and real-time RT-PCR. Results: RT-LAMP assays successfully detected and differentiated IAVs into H1N1, H3N2 and pdm09/H1N1 subtypes. One hundred and sixty seven clinical swab samples from influenza a virus. Out of 30 samples, 21, 7 and 2 were found positive for pdm09/H1N1, H3N2 and seasonal H1 respectively. Conventional one-step RT-PCR detected a total of 27 (16.2%) samples for influenza A and further subtyping showed 20 and 7 samples positive for pdm09/H1N1 and H3N2 virus respectively whereas none was found positive for seasonal H1N1. RT-LAMP assay demonstrated higher sensitivity (93.8%) than conventional RT-PCR (84.4%) for influenza A viruses detection in clinical samples. Conclusions: RT-LAMP assay is rapid, sensitive, specific and cost effective method for detection of influenza A viruses than conventional one-step RT-PCR and it can serve as a good alternate for diagnosis and surveillance tradies than conventional one-step RT-PCR and it can serve as a good alternate for diagnosis and surveillance tradies than conventional one-step RT-PCR

1. Background

Influenza A viruses (IAVs) pose a serious threat to the world and are responsible for severe morbidity and mortality among human population globally. Seasonal outbreaks, epidemics and pandemics have been witnessed for IAVs and have resulted in immense loss to public health and economy (Wright and Neumann, 2013). The devastating potential of Influenza is shown by the facts that annually there are estimated 3-5 million cases of severe illness and 300,000-500,000 deaths around the globe (WHO, 2016). Influenza viruses belong to the family Orthomyxoviridae, characterized by segmented, negative-sense, single stranded

RNA genome (Shaw and Palese, 2013). The family contains seven genera viz., Influenzavirus A, Influenzavirus B, Influenzavirus C, Influenzavirus D, Isavirus, Quaranjavirus and Thogotovirus (ICTV, 2016). Out of the four genera of influenza viruses, influenza A and B are the main cause of seasonal epidemics while influenza C causes mild reparatory illness and influenza D affects cattle and not known to cause any infection in humans (Hause et al., 2014; CDC, 2016; Ferguson et al., 2016).

Influenza or "Flu" is a respiratory illness characterized by high grade fever, myalgia, headache, malaise, sore throat, rhinitis and cough (Cox and Subbarao, 1999). Generally influenza is a self limiting disease

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Abbreviations: IAV, Influenza A viruses; LAMP, loop mediated isothermal amplification; RT, reverse transcription; RT-PCR, reverse transcription polymerase chain reaction; rRT-PCR, real-time RT-PCR: WHO, World Health Organization

but may become life-threatening in high risk group individuals including young children, pregnant women, aged people and people with compromised immunity (Cox and Subbarao, 1999; Cox and Subbarao, 2000). Influenza vaccines are updated annually according to the circulating strains and this information is provided by the surveillance system of WHO's Global Influenza Surveillance and Response System (GISRS) for both the northern and southern hemispheres (WHO, 2012; Webster and Govorkova, 2014). Antivirals treatment proved effective against influenza if given within 24-48 h of the onset of disease (Moscona, 2005). Hence, early and accurate diagnosis of influenza plays a crucial role for timely intervention of therapy, clinical management and thus controlling the spread of disease. Clinical symptoms of influenza and other respiratory infections can be very similar. making it hard to differentiate. Only laboratory diagnosis can provide an accurate influenza diagnosis. These include conventional virus isolation, antigen/antibody detection and molecular methods like reverse transcription polymerase chain reaction (RT-PCR) (Poddar, 2002) and real time RT-PCR (rRT-PCR) (Spackman and Suarez, 2008; Carr et al., 2009; Wang and Taubenberger, 2010). Virus isolation and serological assays are time consuming and take days to weeks to get results making them unsuitable during epidemics and also these methods require highly sophisticated biosafety laboratories. Molecular methods such as RT-PCR and rRT-PCR are although rapid, sensitive and specific but need costly equipments and trained laboratory persons making these assays hard to use in resource-limited laboratories of developing countries.

Loop-mediated isothermal amplification (LAMP) is a specific, efficient and rapid technique that is similar to PCR amplification but the target DNA amplification is under isothermal conditions. This method makes use of a DNA polymerase with strand displacement activity and a set of four specially designed primers that recognize a total of six distinct sequences on the target DNA (Notomi et al., 2000). The method is applicable to amplification of RNA templates when combined with a reverse transcription reaction (RT-LAMP). RT-LAMP assay has been successfully applied for the detection of various RNA viruses including chikungunya (Parida et al., 2007; Lu et al., 2012), dengue (Lu et al., 2012; Parida et al., 2005), Japanese encephalitis (Parida et al., 2006; Toriniwa and Komiya, 2006), severe acute respiratory syndrome (Thai et al., 2004) and West Nile virus (Parida et al., 2004). Influenza viruses and their novel subtypes like H5, H7, H9 and H10, have been effectively detected by RT-LAMP assay (Poon et al., 2005; Ito et al., 2006; Imai et al., 2006; Chen et al., 2008; Zhang et al., 2013; Luo et al., 2015). The specificity of RT-LAMP is extremely high due to use of four primers that recognize six distinct regions on the target DNA. It is faster than PCR because of single temperature is needed for amplification and also cost effective as costly equipments are not required. The amplification results can be viewed directly by using fluorescent dye like SYBR green, reducing the time of the assay.

2. Objectives

The objectives of the current study were to (1) optimize RT-LAMP assay for detection of influenza A viruses and their subtypes (H1N1, H3N2 and pdm09/H1N1); (2) determine sensitivity and specificity of RT-LAMP assay; (3) clinical evaluation of RT-LAMP assay and conventional one-step RT-PCR in comparison to WHO recommended rRT-PCR taken as standard.

3. Study design

3.1. Reference strains of influenza A viruses

Reference strains of Influenza A viruses were kindly provided by the department of Microbiology, All India Institute of Medical Sciences (AIIMS), New Delhi, India. These strains include; A/California/7/2009 (H1N1) pdm09-like virus; A/Texas/50/2012 (H3N2)-like virus; and A/ Malaysia/2089302/2009(H1N1) virus. These strains were aliquoted

and stored at -80 °C until further used.

3.2. RNA extraction from reference strains

Viral RNA was extracted from an aliquot of each reference strain of influenza viruses, using the GeneJET Viral DNA/RNA Purification Kit (Thermo Fisher Scientific, USA) by following the manufacturer's instructions. RNA was eluted in a final volume of 50 μ l in elution buffer (provided with kit) and divided into aliquots of 5 μ l each and stored at - 80 °C until used. RNA extraction was performed in class II biosafety cabinets.

3.3. RT-LAMP primers

For detection of Influenza A viruses, published primers were taken from matrix (M) gene (Poon et al., 2005). For subtyping of IAVs, primers were selected from hemagglutinin (HA) gene. Published primers were used to detect pandemic influenza H1N1 virus (pdm09/H1N1) (Nakauchi et al., 2011). For seasonal H1N1 and H3N2 virus, HA gene sequence of the respective reference strain was retrieved from GenBank database (GeneBank accession numbers CY118255 and KC892952) and primers were designed by using Primer Explorer version 4 (https:// primerexplorer.jp/e/). Four primers consisting of two internal primers (Forward internal primer or FIP and backward internal primer or BIP) and two external primers (F3 and B3) were designed for the study. All primers were chemically synthesized by Sigma-Aldrich (Sigma-Aldrich, Bangalore, India) and were of HPLC grade purity (Table 1).

3.4. RT-LAMP assay

Twenty five microliters of RT-LAMP reaction cocktail was prepared in 0.2 ml tubes using 0.2 µM of each outer primers (F3 and B3); 1.6 µM of each inner primers (FIP and BIP); 1.4 mM of each deoxynucleotides (dNTPs); 20 mM Tris-HCl (pH 8.8); 10 mM (NH₄)₂SO₄; 10 mM KCl; 8 mM MgSO₄; 0.8 M betaine; 0.1% Tween 20; 8 U of Bst DNA polymerase (New England Biolabs); 0.5 U of AMV reverse transcriptase (Invitrogen, USA) and 1 µl of SYBR green I (Thermo Fisher Scientific, USA). Two microliters of each reference strain's RNA was added to their respective tubes (containing strain specific primers) making final volume 25 µl. No template or negative controls were also included in the assay. The reaction tubes were incubated at 63 °C for 1 h either in a water bath or a heating block. The tubes were heated at 80 °C for 5 min to stop the reaction. Amplified products were visualized both by electrophoresis using 2.5% agarose gel and by visual detection of green fluorescence due to added SYBR green I stain (Thermo Fisher Scientific, USA) under ultra violet (UV) light.

3.5. Conventional one-step RT-PCR

Conventional one-step RT-PCR was performed on RNA samples extracted from the reference strains of IAVs. For detection of IAVs, primer pair was taken from the matrix (M) gene while subtyping primer pairs were taken from hemagglutinin (HA) gene. For diagnosis of pandemic influenza virus, primer pair was selected from the conserved region of HA gene (GeneBank accession number KU933485) using primer blast software (https://www.ncbi.nlm.nih.gov/tools/primerblast/). Details of primer sets were given in Table 2. One-step RT-PCR reaction was carried out in final reaction volume of 25 µl by using Verso 1-Step RT-PCR Kit (Thermo Fisher Scientific, USA) by following manufacturer's instructions. Briefly, reaction mixture was comprised of 12.5 µl of 2X RT-PCR buffers; 1 µl of RT enhancer; 0.2 µM of each forward and reverse primer; 0.5 µl of enzyme mix and 2 µl of extracted RNA. The protocol for optimization of one-step RT-PCR was as follows: 50 °C for 30 min, 95 °C for 15 min, followed by 40 cycles of 95 °C for 30 s, 52 °C–58 °C for 30 s, 72 °C for 45 s and final extension at 72 °C for 10 min. PCR products were separated by electrophoresis using 2%

Table 1

Details of primers used in RT-LAMP assay.

Primer	Sequence $(5' \rightarrow 3')$	Genome location	Primer length (bp)	Reference	
Influenza A/M-gene					
Inf A-F3	TGGTGCACTTGCCAGTTG	375–392	18	Poon et al., 2005	
Inf A-B3	CCAGCCATCTGTTCCATAGC	593-574	20		
Inf A-FIP	TGCTGTGAATCAGCAATCTGTTACAGGA	F1c, 488-467	42		
(F1c + F2)	TGGGAACAGTGACC	F2, 410–429			
Inf A-BIP	AGACAAATGGCTACTACCACCCGTAGT	B1c, 499–519	40		
(B1c + B2)	GCTAGCCAGCACC	B2, 567-549			
A/pdm09H1/HA-gene (par	demic Influenza A)				
pdmH1-F3	AGCTAAGAGAGCAATT	170–185	16	Nakauchi et al., 2011	
pdmH1-B3	TTTCCCTTTATCATTAATGTAGGATTTG	384-357	28		
pdmH1-FIP	ACCTTTGTTCGAGTCATGATTTGGTTCTC	F1c, 264-244	51		
(F1c + F2)	AGTGTCATCATTTGAAAGGTTT	F2, 189–213			
pdmH1-BIP	TAACGGCAGCATGTCCTCATTGTATGAA	B1c, 266–285	48		
(B1c + B2)	TTTCCTTTTTTAACTAGCCA	B2, 345-319			
A/H1/HA-gene (seasonal Influenza A/H1)					
H1-F3	CTGTTATGCACATTTACAGCT	39–59	21	This study	
H1-B3	TTCCTAAGATCCACCCGG	258-241	18		
H1-FIP	ACTGTGTCAACAGTGTCGGTTTTGCAGA	F1c, 124-105	44		
(F1c + F2)	CACAATATGTATAGGC	F2, 65–86			
H1-BIP	ACACACTCTGTCAACCTGCTTTTCCAAT	B1c, 147–167	41		
(B1c + B2)	TGTAGTGGGGCTA	B2, 225-208			
A/H3/HA-gene (seasonal I	nfluenza A/H3)				
H3-F3	ATAGTAAAACCGGGAGACAT	754–773	20	This study	
H3-B3	TGCTTAACATATCTGGGACA	980-961	20		
H3-FIP	TGGGTGCATCTGATCTCATTATTGATTTT	F1c, 868-844	48		
(F1c + F2)	AGGGAATCTAATTGCTCCT	F2, 792–810			
H3-BIP	GCAAGTCTGAATGCATCACTCCTTTTTG	B1c, 878–899	48		
(B1c + B2)	ATCCTGTTTACATTTTGGAA	B2, 949-928			

agarose gel and visualized by UV transilluminator.

3.6. Sensitivity and specificity of RT-LAMP

Influenza virus A/California/7/2009 (H1N1) strain with known virus titre as plaque forming units per microliters (PFU/ μ l) was used to compare sensitivities of RT-LAMP and conventional RT-PCR with WHO recommended Taqman real-time RT-PCR (rRT-PCR). Ten-fold serial dilutions were made from virus strain in separate tubes ranging from 10² to 10⁻⁴ PFU per reaction using molecular grade water. RNA was extracted from each dilution and used separately for RT-LAMP, conventional RT-PCR and rRT-PCR. Matrix gene specific primers and 2 μ l of RNA from each dilution were used in all assays. Likewise, sensitivities of RT-LAMP, conventional RT-PCR and rRT-PCR and rRT-PCR were also assessed for subtyping of influenza viruses using HA gene specific primers on H1N1 and H3N2 reference strains. For specificity evaluation, each individual RT-LAMP assay was applied on influenza A viruses (pdm09/H1N1, A/H1N1 and A/H3N2), influenza B virus along with other additional

Table 2

Details of primers used in conventional one-step RT-PCR assay.

respiratory viruses including respiratory syncytial virus (RSV) and human metapneumovirus (hMPV), and cross-reactivity of the assay was checked.

3.7. Clinical samples

A total of 167 clinical samples were collected from patients with influenza like illness (high grade fever (> 38.5 °C), cough, sore throat and myalgia) from Post Graduate Institute of Medical Sciences (PGIMS), Rohtak, Haryana, India from November 2014 to December 2016. One throat and one nasal swab was collected in 2 ml of viral transport medium (Hanks' balanced salt solution supplemented with 2% BSA, 100 U/ml penicillin and 50 µg/ml streptomycin) from each patient and transported to the laboratory under cold conditions at 4 °C. The samples were processed and RNA was extracted in class II biosafety cabinets.

Primer	Sequence $(5' \rightarrow 3')$	Primer length (bp)	Amplicon size (bp)	Reference
Influenza A/M-gene Inf A-F Inf A-R	AGTCTTCTAACCGAGGTCGAAACG TGGACAAAGCGTCTACGCTGCA	24 22	239	WHO, 2014
Influenza A/H1/HA-gene H1-F H1-R	AATTTGCTATGGCTGACGGG GCTATAATGTGAAGACACTACAGA	20 24	183	Poddar, 2002
Influenza A/H3/HA-gene H3-F H3-R	GCGACAGTCCTCATCAGATCCT CATTGTTTGGCATAGTCACGTTC	22 23	360	Poddar, 2002
Influenza A/pH1/HA-gene pdmH1-F pdmH1-R	ACACTAGTAGAGCCGGGAGA CAATCCTGTGGCCAGTCTCA	20 20	261	This Study

Fig. 1. Detection of influenza A viruses by RT-LAMP and conventional

RT-PCR using matrix (M) gene primers. (a) examination of RT-LAMP products in 2.5% agarose gel electrophoresis and direct visualization of green fluorescence in positive amplification reactions in 0.2 ml PCR tubes using SYBR green dye under UV light (b); lane M, 100 bp DNA marker: lane/tube 1. no template control: lane/tube 2. influenza H1N1; lane/tube 3, influenza H3N2; lane/tube 4, influenza pdm09/ H1N1. (c) Agarose gel electrophoresis profile of conventional one-step RT-PCR amplified products showing a band of 239 bp. All lanes have similar order as mentioned above except lane M containing 50 bp DNA marker. (For interpretation of the references to colour in this figure

legend, the reader is referred to the web version of this article.)



with WHO recommended Taqman real-time RT-PCR (rRT-PCR) considering the later as standard. Standardized RT-LAMP and conventional Sensitivity of RT-LAMP assays were assessed for detection of influone-step RT-PCR were applied on clinical samples and amplified products were analyzed by gel electrophoresis. Taqman rRT-PCR assay was applied on extracted RNAs from clinical samples for detection of influenza A viruses and further subtyping into H1N1, H3N2 and pdm09/ H1N1 subtypes. Tagman rRT-PCR assay was performed in a 25 µl reaction volume using WHO recommended protocol and primers/probes 4.1. Optimization of RT-LAMP and conventional one-step RT-PCR assay

All three reference strains of influenza A virus were detected by matrix gene specific RT-LAMP reaction (Fig. 1a). On further subtyping, each respective subtype was also detected by HA gene specific RT-LAMP assay (Fig. 2a). Optimum amplification temperature and incubation time were found to be 63 °C and 1 h respectively. Amplification results were visualized by both gel electrophoresis and green fluorescence detection under UV light (Fig. 1b and 2b). Conventional one-step RT-PCR was able to detect all three influenza A virus strains using matrix gene primers. Optimum annealing temperature and time was found to be 55 °C for 30 s. Amplicons of 239 bp were visualized using 2% agarose gel electrophoresis (Fig. 1c). For subtyping conventional RT-PCR, annealing at 55 °C for 30 s was found to be optimum for both pdm09/H1N1 and H3N2 subtypes whereas for seasonal H1N1 subtype, annealing temperature of 54 °C for 30 s was optimized. Amplicons of 261 bp, 183 bp and 360 bp were detected for pdm09/H1N1,

3.8. Detection and subtyping of influenza A viruses in clinical samples

sequences (WHO, 2014).

4. Results

Clinical samples were evaluated by using optimized RT-LAMP and conventional one-step RT-PCR and detection results were compared

> enza A viruses using M gene primers and for detection of three subtypes i.e. H1N1, H3N2 and pdm09/H1N1, using HA gene primers (Table 3). The results were compared with that of conventional one-step RT-PCR and rRT-PCR. The detection limit of M gene specific RT-LAMP was found to be ten times higher (0.01 PFU/reaction) than conventional one-step RT-PCR (0.1 PFU/reaction) (Fig. 3a) and ten times lower than rRT-PCR (0.001 PFU/reaction) (Table 3). The detection limits of HA gene specific RT-LAMP assays for pdm09/H1N1, H3N2 and H1N1 were 0.01, 0.1 and 0.1 PFU per reaction respectively. The detection limits of conventional one-step RT-PCR for pdm09/H1N1, H3N2 and H1N1 were 1, 0.1 and 1 PFU per reaction respectively, making it 10–100 times less sensitive than RT-LAMP. RT-LAMP demonstrated similar sensitivities as that of rRT-PCR for the detection of pdm09/H1N1 and H3N2 while 10 times less sensitivity for detection of H1N1 subtype (Table 3). Specificity of each RT-LAMP primer set was evaluated against different influenza viruses and other respiratory viruses. All the RT-LAMP primer sets were found to be highly specific for their respective influenza viruses as there was no amplification with other influenza viruses or different respiratory viruses. As shown in Fig. 3b, pdm09/H1N1 RT-LAMP (HA gene) was found to be highly specific for detecting pandemic IAVs as only pdm09/H1N1 reaction showed positive amplification while there was no amplification with other viruses.

4.3. Detection and subtyping of influenza A viruses in clinical samples

All clinical samples were analyzed for presence of IAVs and their subtypes (seasonal H1N1, H3N2 and pdm09/H1N1) using WHO

H1N1 and H3N2 respectively, after 2% agarose gel electrophoresis (Fig. 2c).

4.2. Sensitivity and specificity of RT-LAMP assay



Fig. 2. Detection of influenza A viruses subtypes using RT-LAMP and conventional one-step RT-PCR using HA gene primers. (a) agarose (2.5%) gel electrophoresis profile of RT-LAMP assays products (b) visual detection of green fluorescence in 0.2 ml PCR tubes using SYBR green dye under UV light. Lane M, 100 bp DNA marker; lane/tube 2, influenza H1N1: lane/tube 4, influenza H3N2: lane/tube 6, influenza pdm09/H1N1; lane/tube 1, 3, 5 are no template controls for H1N1, H3N2 and pdm09/H1N1 respectively. (c) agarose (2.5%) gel electrophoresis profile of conventional one-step RT-PCR reaction products. Lane M, 100 bp DNA marker; lane 1, no template control; lane 2, influenza A virus detection using M gene primers showing a band of 239 bp; lane 3, influenza A/pdm09/H1N1 showing a band of 261 bp; lane 4, influenza A seasonal H1N1 showing a band of 183 bp; lane 5, influenza A H3N2 showing a band of 360 bp. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

recommended Taqman real-time RT-PCR, RT-LAMP assay and conventional RT-PCR assay. Real-time RT-PCR results showed that out of 167 swab samples, 32 (19.2%) were positive for influenza A viruses. Further subtyping of 32 (n = 32) positive samples by real-time RT-PCR showed that 23 (71.9%) samples were positive for pdm09/H1N1, 7 (21.9%) were positive for H3N2 and 2 (6.2%) were positive for H1N1 virus. Matrix gene specific RT-LAMP assay detected 30 (17.9%) samples positive for influenza A viruses which were further sub-typed using HA gene specific RT-LAMP. Results showed that 21 out of 30 (70%) samples were positive for pdm09/H1N1, 7 (23.3%) samples were positive for H3N2, and 2 (6.7%) samples were positive for H1N1 virus. These results showed the comparable sensitivity and specificity of real-time RT-PCR with that of RT-LAMP assay (Table 4). Conventional RT-PCR detected 27 (16.2%) samples positive for influenza A viruses out of which 20 (74.1%) were found positive for pdm09/H1N1 and 7 (25.9%) were positive for H3N2 virus where as no sample was detected positive for H1N1 virus. All samples which were found to be negative by realtime RT-PCR also come negative by both RT-LAMP and conventional RT-PCR assay.

5. Discussion

Influenza surveillance has importance not only in diagnosis of circulating strains of the virus but also to upgrade the available vaccine on annual basis to provide protection against influenza. Control of novel influenza A viruses mainly depends of early and accurate identification of virus strain followed by early intervention of antiviral therapy. So, a rapid, sensitive, specific and cost effective method of detection and subtyping of influenza A viruses is always seems fruitful in this scenario. Virus isolation using egg embryo culture or cell culture is considered as 'gold standard' for influenza virus detection but the whole process is labour intensive and lengthy taking 3-7 days for obtaining results (Ellis and Zambon, 2002). Serological methods are also not very reliable for influenza surveillance as these are time consuming and require analysis of serum samples from both acute phase and recovering phase. Rapid detection kit based assays are although fast but are not very consistent, have low sensitivity and may produce false results (Wang and Taubenberger, 2010; Ellis and Zambon, 2002; Kim and Poudel, 2013). Molecular detection methods have emerged as very

Table 3

Limit of detection of RT-LAMP, conventional one-step RT-PCR and real-time RT-PCR assay using virus titre as plaque forming units (PFU).

Influenza strain	Target gene	RT-LAMP (PFU/reaction)	Conventional RT-PCR (PFU/reaction)	WHO rRT-PCR (PFU/reaction)
A/California/7/2009 (H1N1) pdm09-like virus	M	0.01	0.1	0.001
A/California/7/2009 (H1N1) pdm09-like virus	HA	0.01	1	0.01
A/Texas/50/2012 (H3N2)-like virus	HA	0.1	0.1	0.1
A/Malaysia/2089302/2009 (H1N1)-like virus	HA	0.1	1	0.01



Fig. 3. Sensitivity and specificity of RT-LAMP assay. (a) Sensitivity comparison of RT-LAMP assay (upper panel) and conventional RT-PCR (lower panel) for detection of IAVs using M gene primers, was performed using ten-fold serial dilutions of influenza A/California/7/ 2009(H1N1) strain with concentrations ranging from 100 to 0.0001 PFU/reaction tube. Detection limit of RT-LAMP assav was 0.01 PFU/ reaction (upper panel) while conventional RT-PCR (lower panel) showed a detection limit of 0.1 PFU/reaction making RT-LAMP assay ten times more sensitive than conventional RT-PCR. Lane M, 100 bp DNA marker. (b) Influenza pdm09/H1N1 specific RT-LAMP reaction virus using HA gene primers was applied on other respiratory viruses to check the specificity of the reaction. Agarose electrophoresis profile of the RT-LAMP reaction products; lane M, 100 bp DNA maker; lane 1, no template control; lane 2, influenza A pdm09/H1N1; lane 3, influenza A/H1N1; lane 4, influenza A/H3N2; lane 5, influenza B virus; lane 6, respiratory syncytial virus (RSV); lane 7, human metapneumovirus (hMPV).

Table 4

Identification of influenza A viruses and subtypes (H1N1, H3N2 and pdm09/H1N1) in clinical samples from Haryana, India, using RT-LAMP and conventional one-step RT-PCR, confirmed by WHO recommended rRT-PCR. Sensitivity and specificity were calculated by taking rRT-PCR as standard. The numbers in parentheses indicate the real-time RT-PCR results. ND: not determined.

	Detection and subtyping of influenza A viruses					
	Influenza A virus	Influenza A/H1N1	Influenza A/H3N2	Influenza A/pdm09/H1N1	Negative	Total
RT-LAMP assay Conventional RT-PCR Sensitivity/specificity of RT-LAMP assay (%) Sensitivity/specificity of conventional RT-PCR (%)	30 (32) 27 (32) 93.8/100 84.4/100	2 (2) 0 (2) ND ND	7 (7) 7 (7) 100/100 100/100	21 (23) 20 (23) 91.3/100 87/100	137 (135) 140 (135)	167 167

useful tools in comparison to time consuming conventional virus isolation and serological methods. Molecular methods based on conventional RT-PCR and real-time RT-PCR have been proved useful in diagnosis and surveillance of influenza viruses, but most of these assays require highly sophisticated instruments and are not feasible to conduct in resource-limited settings of developing countries.

Therefore, present study was focused on RT-LAMP assay for detection of influenza A viruses and further subtyping into seasonal influenza (H1 and H3) and novel pandemic influenza A virus (pdm09/H1N1). RT-LAMP assay has been successfully applied to detect influenza A viruses like swine flu (H1N1), pdmH1N1, avian influenza H5, H7, H9 and H10 subtypes (Poon et al., 2005; Ito et al., 2006; Imai et al., 2006; Chen et al., 2008; Luo et al., 2015; Kubo et al., 2010; Parida et al., 2011; Bao et al., 2012). In this study, we used matrix gene based primers for detection of influenza A viruses (Poon et al., 2005) and hemagglutinin gene based primers for their further subtyping. Matrix gene based RT-LAMP assay was able to amplify all three reference strains of influenza A virus and hemagglutinin gene based RT-LAMP assay successfully detected the respective subtypes. No cross-reactivity was observed for other related viruses and influenza subtypes making the reaction highly specific.

Sensitivity of a detection assay is an important deciding factor when it comes to applicability of the assay. In this study we have compared RT-LAMP with conventional RT-PCR and Taqman real-time RT-PCR. In our study, detection limit of Taqman rRT-PCR was found to be ten times more than RT-LAMP for influenza A and its subtype H1N1 whereas for detection of pdm09/H1N1 and H3N2 both assays showed similar values. Sensitivity comparison of RT-LAMP with conventional RT-PCR showed that RT-LAMP was 10–100 times more sensitive than conventional RT-PCR. Previous studies have also established that RT-LAMP is more sensitive than conventional RT-PCR and has comparable sensitivity with real-time RT-PCR for detection of influenza viruses (Poon et al., 2005; Kubo et al., 2010; Parida et al., 2011; Bao et al., 2012; Sharma and Kaushik, 2016).

In the present study, a total of 167 clinical samples were analyzed for the presence of IAVs and their further subtyping was done. We have

used RT-LAMP assay, conventional RT-PCR and rRT-PCR assay in parallel reactions for diagnosis of influenza in clinical samples. Real-time RT-PCR reaction was performed using WHO recommended primers (WHO, 2014) and was used as standard to compare RT-LAMP and conventional RT-PCR assay. Real-time RT-PCR detected 32 samples positive for influenza A viruses, which on further subtyping showed 2, 7 and 23 samples for influenza H1N1, H3N2 and pdm09/H1N1 respectively. Sensitivity of conventional RT-PCR for detection of influenza A viruses and subtype pdm09/H1N1 was 84.4% (27/32) and 86.9% (20/ 23) respectively. Whereas RT-LAMP assay showed sensitivity of 93.8% (30/32) for influenza A viruses and 91.3% (21/23) for subtype pdm09/ H1N1 detection, indicating higher sensitivity than conventional RT-PCR. Both these assays detected H3 subtype with equal sensitivity of 100% (7/7). For seasonal H1 subtype, sensitivity and specificity could not be determined due to small number of positive samples. The results showed that RT-LAMP assay can be applied successfully for the diagnostic and surveillance studies of influenza viruses.

Detection time is a crucial factor of any diagnostic assay. RT-LAMP assays were completed in 1 h and results were shown after 30 min of agarose gel electrophoresis. So RT-LAMP assay gave efficient results within one and a half hours of sample processing (RNA extraction) making this assay fast when compared with conventional RT-PCR and real time RT-PCR which gave results within 2-3 h (Gu et al., 2010). Previous studies have reported that by using two accessory loop primers the amplification time can be reduced from 60 min to 30-40 min (Luo et al., 2015; Kubo et al., 2010; Gu et al., 2010; Imai et al., 2007; Peng et al., 2011). For the sake of simplicity and making our reaction more cost effective we stuck to the basic protocol of RT-LAMP assay and use only four primers for amplification which didn't compromise with sensitivity of the assay. RT-LAMP assay can be modified to detect samples in real-time by using turbidity measurement by turbidimeter as turbidity of reaction increases due to release of pyrophosphates during amplification. Moreover, addition of a florescent dye like calcein or SYBR green in the reaction can be used for direct visualization of the amplification products under UV light thus eliminating the need of gel electrophoresis making the reaction more rapid and less prone to crosscontaminations (Luo et al., 2015; Kubo et al., 2010; Bao et al., 2012; Gu et al., 2010; Peng et al., 2011) .We used SYBR green I dye (Sigma-Aldrich, USA) for direct visualization under UV light and both visual detection and agarose gel electrophoresis produced similar results. This showed that RT-LAMP assay can be applied as a point-of-care test during influenza outbreaks due to its simplicity, high sensitivity and less turn-around time.

In conclusion, this study showed that RT-LAMP assay can be used for detection and surveillance studies during the time of influenza outbreaks as a good alternative for conventional RT-PCR and real-time RT-PCR. RT-LAMP assay is highly cost effective and easy to perform as sophisticated instruments like a thermocycler and fluorescent data analyzing computers (as in real-time RT-PCR) are not needed and the whole reaction can be done in a controlled temperature water bath or dry heating block. The results of the study clearly showed that RT-LAMP assay can be applied successfully to the clinical samples and the results were compatible with that of the WHO's recommended real-time RT-PCR. The RT-LAMP assay is a simple, rapid, specific, sensitive and cost effective method for detection and subtyping of influenza A viruses and may prove useful in the resource-poor laboratories of the developing countries during influenza outbreaks.

Contributions

VS has designed the study, performed the experiments, analyses and writing, DC participated in designing the study, SK designed the study, performed analyses and writing.

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Ethical clearance

This study was approved by the Human Ethical Committee (letter no. PHY/13/362), Maharshi Dayanand University, Rohtak, Haryana, India and detailed informed consent was taken from each patient before taking swab samples.

Competing interests

None declared.

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