

**Isothermal amplification
techniques for FBP detection.
A focus on LAMP application for
the detection of *Toxoplasma
gondii* in different food matrices.**

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Detection of parasite in food: technical and conceptual challenges

➤ A major shortcoming in food parasitology is the **lack of well standardized protocols to recover, detect, and identify sparsely distributed parasites in food and in the entire food chain**

- Toxoplasma tissue cysts in meat
- Anisakis larvae in fish flesh
- Protozoan (oo)cysts in water and vegetable



➤ Parasites load in foodstuff and water is different from the load in clinical samples (i.e. Cryptosporidium oocyst in stool vs vegetables)

➤ Large number of **different matrices to analyze**

➤ Unlike most bacteria, parasites cannot be enriched by culturing; therefore, **an optimal recovery process for parasite from suspected foods is critical.**

➤ Diagnostics are still based on macro- and microscopic inspection, and require highly qualified personnel.

➤ Nucleic acid based test improve sensitivity and specificity while also reduce detection time

➤ **Necessity to detect very low number of organisms**

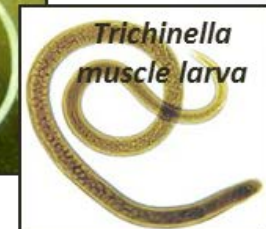
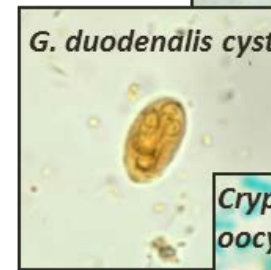
Diagnostic solutions: qPCR, PCR, Microarray, Loop-mediated isothermal amplification (LAMP)

➤ **Necessity to distinguish between human and non-human pathogens and/or trace the source of infection (outbreak investigations)**

Diagnostic solutions: PCR (polymorphic loci), qPCR, Microarray, Single-strand conformation polymorphism (SSCP), Nucleic acid sequence based amplification (NASBA)

➤ **Necessity to distinguish between viable (infectious) and non-viable organisms**

Diagnostic solutions: RT-PCR, RT-LAMP, Nucleic acid sequence based amplification (NASBA), FISH, but also the use of vital dyes and bioassays



Commonly used nucleic acid-based assays

➤ Polymerase Chain Reaction (PCR)

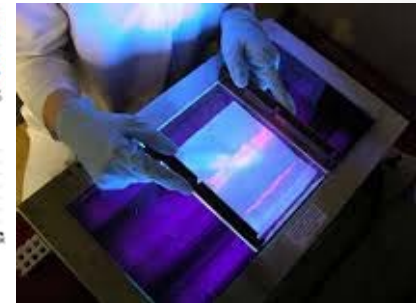
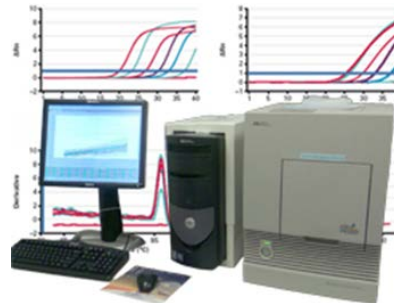
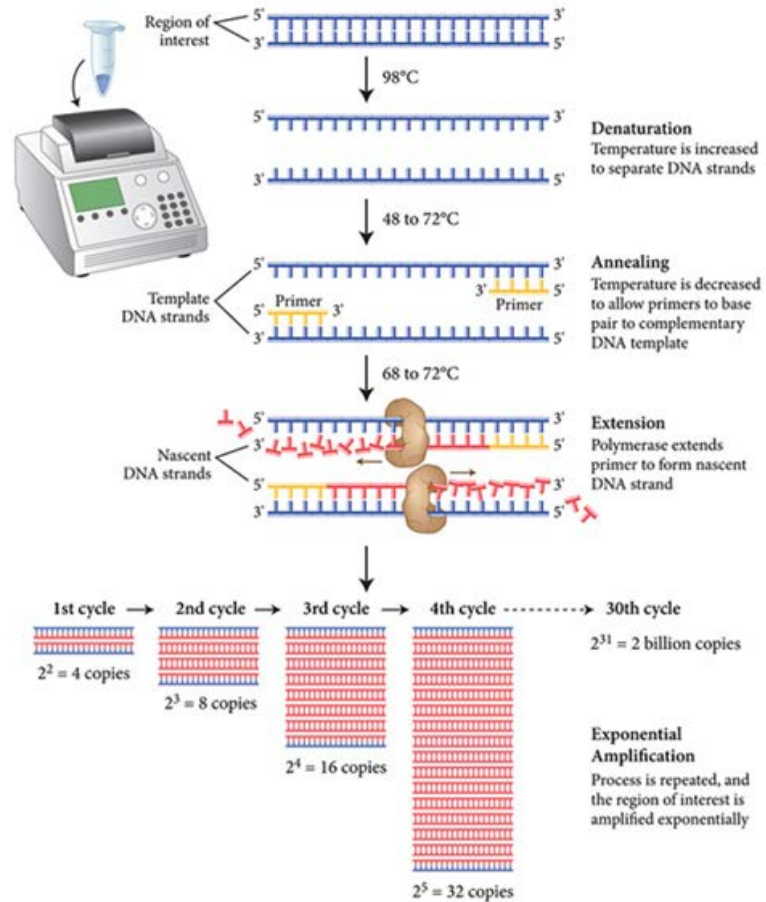
- Single-Step PCR
- Nested-PCR
- Multiplex PCR
- PCR-RFLP (Restriction fragment length polymorphism)
- Quantitative PCR (Real-time PCR or qPCR)
- Capillary Electrophoresis-Single Strand Conformation polymorphism (CE-SSCP)
- Random Amplified Polymorphic DNA (RAPD)-PCR

Requirements:

- thermal cycler (or more sophisticated instruments with detector)
- high quality nucleic acid preparation
- technical expertise of personnel
- cold chain to preserve heat-labile reagents

Drawbacks:

- time consuming
- high cost
- easy contamination
- susceptibility to false amplification



Non-PCR nucleic acid-based assays

- **Loop-mediated isothermal Amplification (LAMP)**
- **Nucleic Acid Sequence-Based Amplification (NASBA)**
- **Recombinase polymerase amplification (RPA)**
- **Strand displacement amplification (SDA)**
- **Rolling circle amplification (RCA)**
- **Helicase-dependent amplification (HDA)**

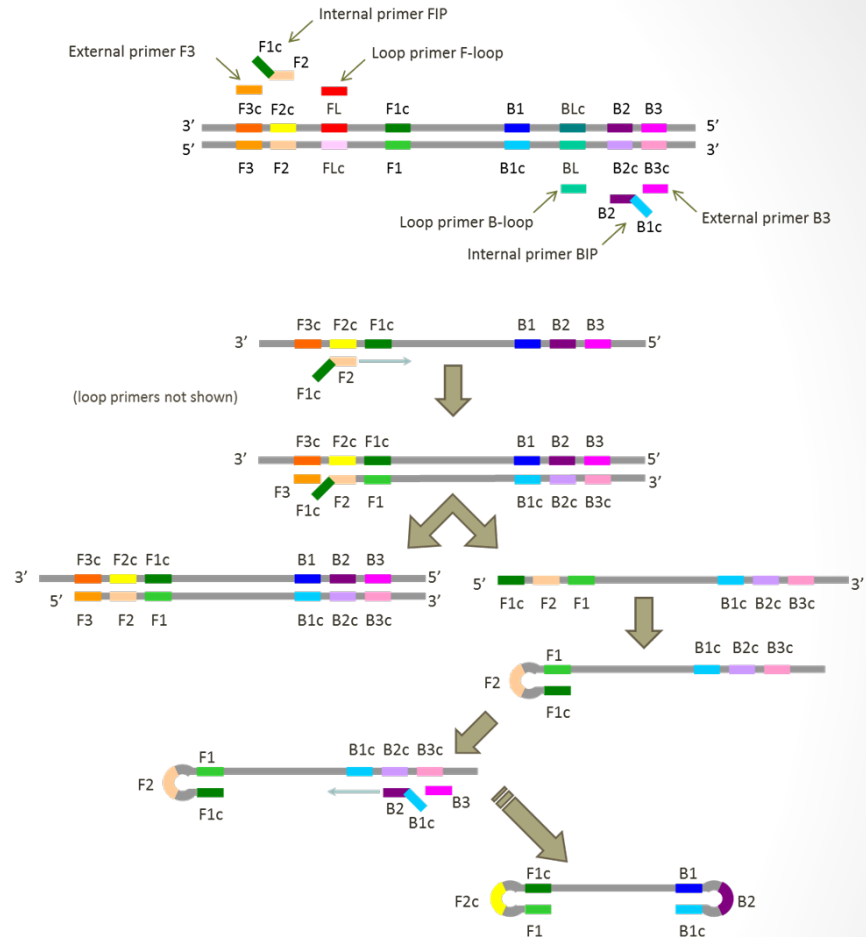
Isothermal amplification techniques have several advantageous qualities:

- short reaction times
- high sensitivity
- not readily affected by inhibitory substances that are often abundant in complex matrices (highly pure nucleic acids not required)
- simple heating device is sufficient to run the reactions (not need of expensive instruments)
- are well suited for incorporation into lab-on-a-chip devices
- amplification feasible in low-resource settings or at the point-of-need

| Isothermal technique | Target | Primers needed | Initial heating | Incubation temperature (°C) | Amplification time (min) | Limit of detection (copies) | Multiplexing | Lyophilised reagents | FDA approved tests |
|----------------------|---------|----------------|-----------------|-----------------------------|--------------------------|-----------------------------|--------------|----------------------|--------------------|
| NASBA | RNA | 2 | No | 41 | 60–180 | 1 | Yes | Yes | Yes |
| SDA | DNA | 4 | Yes | 30–55 | 60–120 | 10 | Yes | No | Yes |
| RCA | DNA/RNA | 1 | Yes | 30–65 | 60–240 | 10 | No | No | No |
| LAMP | DNA | 4–6 | Yes | 60–65 | 60 | ≈ 5 | Yes | No | Yes |
| HDA | DNA | 2 | No | 65 | 30–120 | 1 | Yes | No | Yes |
| RPA | DNA/RNA | 2 | No | 37–42 | 20–40 | 1 | Yes | Yes | No |

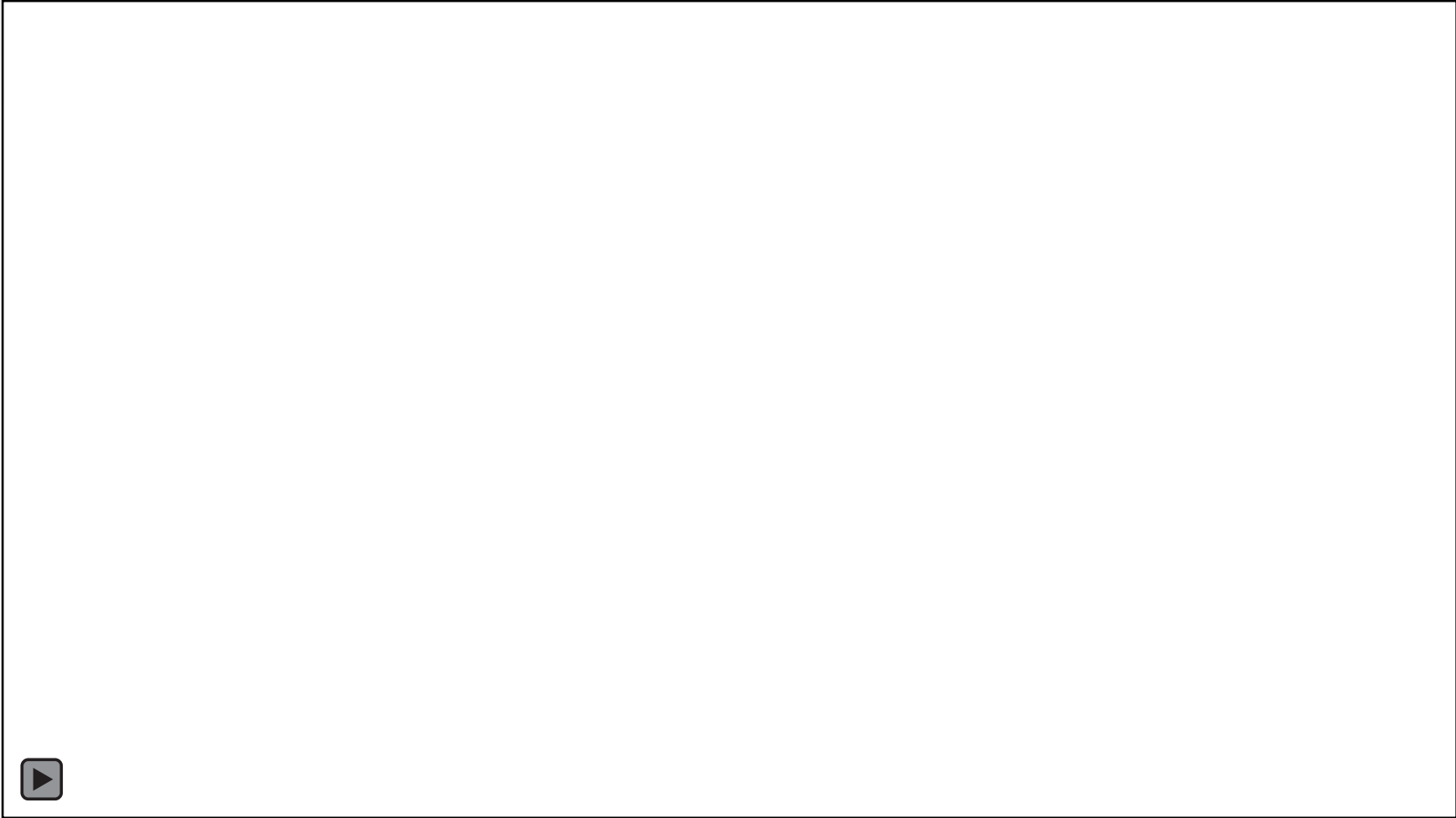
Loop-Mediated Isothermal Amplification (LAMP)

- To increase the sensitivity and specificity, amplification is performed under isothermal conditions employing 4-6 primers spanning 6-8 distinct sequences on the target DNA.
- It uses a robust polymerase (BST), originally from *Geobacillus stearothermophilus* to amplify target DNA (or RNA by inclusion of reverse transcriptase) proceeding to an autocycling strand displacement mechanism, at a **constant temperature (61-65°C)**
- Producing detectable product in approximately 30-60min (amplify from a single copy to 10^9).
- Production of magnesium pyrophosphate during the reaction increases turbidity and allows naked eyes detection of end product (with or without the addition of fluorescent dye)



Notomi et al. 2000. Nucleic Acids Research 28:e63.

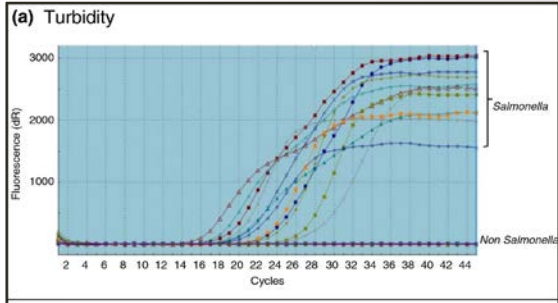
Loop-Mediated Isothermal Amplification (LAMP)



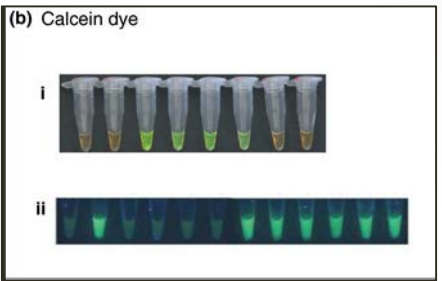
Advantages

- Bst polymerase: 5'→3' DNA polymerase activity and strand-displacement activity. Not affected by inhibition due to the secondary structure of DNA
- Amplification may be performed simply in a heating block or water bath.
- Very rapid amplification (30-60 minutes)
- High specificity
- Tolerance of polymerase inhibitors** can be applied to nucleic acid extracts of unpurified samples or even to samples without nucleic acid extraction
- Suitable for synthesis of DNA strands having high GC content.
- Amplification and detection may be carried out in one single tube.
- Can amplify RNA sequences through combination with reverse transcription.
- Versatile detection (turbidimer, fluorescent dye, gel electrophoresis, flow-dipstick).

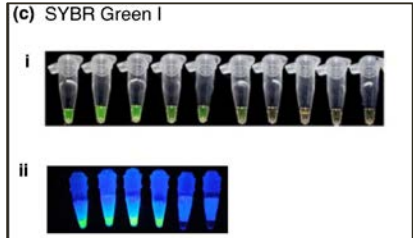
LAMP detection using Loopamp Realtime Turbidimeter LA-500. Increase in graph indicate positive LAMP amplification (indicated with *Salmonella*) whereas no amplification curve indicates negative LAMP amplification (indicated with non *Salmonella*) (image adapted from Mashooq *et al.*, 2016)



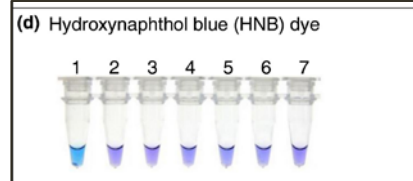
Color change using DNA intercalating dye, Calcein. Color changes from orange to green indicate positive LAMP reaction (i.) (image adapted from Zhou *et al.*, 2014). The reaction between free calcein and magnesium ions intensified the fluorescence of LAMP product (ii.) (image adapted from Chen *et al.*, 2014).



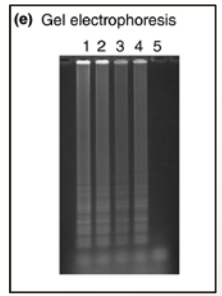
Naked eye observation using DNA fluorescence SYBR Green I dye. Notable color changes from orange to green, indicate positive reaction (i., image adapted from Chen *et al.*, 2017). The LAMP positive tubes show bright green fluorescence, negative tubes remain dark (ii., image adapted from Tao *et al.*, 2011).



LAMP end-point detection using HNB dye as visual indicator. Positive LAMP amplification changes the violet (tube labeled 2-7) to sky blue/teal (tube labeled 1), image adapted from Luo *et al.*, 2014.

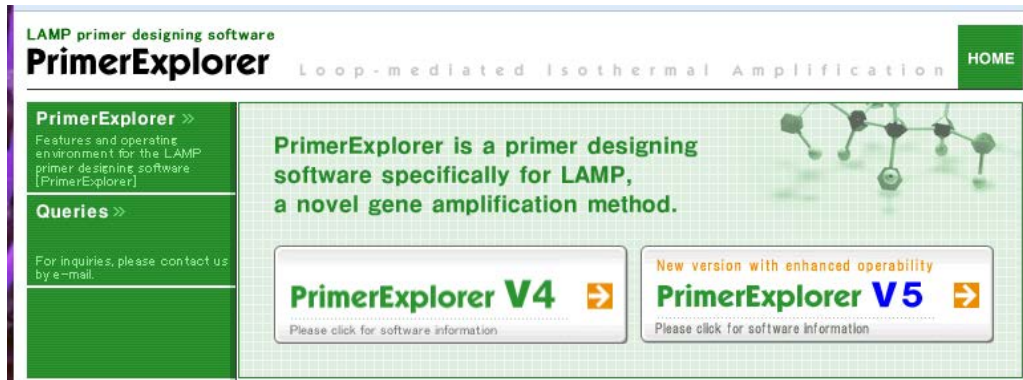


Gel electrophoresis with typical ladder like pattern (positive indicated by 1-4, negative indicated 5)



❑ Disadvantages

- Primer design quite complicated, a software is needed.



The screenshot shows the homepage of the PrimerExplorer software. At the top, it says "LAMP primer designing software" and "PrimerExplorer" in large green letters. Below that, it reads "Loop-mediated isothermal Amplification" and "HOME" in a green box. The main content area has a green background with a molecular structure image. It states: "PrimerExplorer is a primer designing software specifically for LAMP, a novel gene amplification method." There are two buttons: "PrimerExplorer V4" and "PrimerExplorer V5", both with arrows and the text "Please click for software information". A sidebar on the left contains links for "PrimerExplorer >>", "Queries >>", and contact information.

- Inadequate for the detection of unknown or unsequenced targets
- Challenges to using LAMP for multiplex assays in a single sample
- Challenges in quantitation of target DNA.
- High risk of carryover contamination which often leads to false-positive results in negative controls. Procedure for in-tube detection of DNA amplification are necessary.
- In case contamination, perform LAMP in a laminar flow hood and use separate pipettes and filtered tips.
- For procedures that involved opening of the reaction tubes, carry out LAMP in an isolated room

Table 1
Overview of LAMP assays for the detection of bacterial pathogens and zoonoses as well as mycotoxigenic fungi in food.

| Target species | DNA target | Assay type | Food sources | Detection limit | Reference |
|---|---|--------------------------------------|--|---|---|
| Gram negative bacterial pathogens | | | | | |
| <i>Salmonella</i> spp. | invA gene | LAMP ^a | Clareto (on-leaves), lettuce, parsley, spinach, tomato, jalapeno pepper | 2 cfu/25 g | Zhang et al., 2011 |
| spp. | MD ^b | LAMP | Raw milk | 100 fg DNA/reaction | Wang et al., 2008b |
| spp. | MD ^b | LAMP | Milk | 10 ² cfu/ml | Zhu et al., 2008 |
| spp. | MD ^b | LAMP | Raw milk | 10 ² cfu/ml | Wang et al., 2008a |
| spp. | MD ^b | LAMP kit ^c | Egg, milk, yogurt, omelet, hamburger, raw pork, beef, chicken, roast beef, lettuce, vegetable salad, watermelon, apple juice, spinach, sauté, cake, blood and human stool of healthy human | 10 ² cfu/ml | Ueda and Kuwabara, 2009 |
| spp. | EMA ^d LAMP | ND | 100 fg DNA/tube | | Lu et al., 2009 |
| spp. | EMA ^d LAMP | ND | 6.1 × 10 ⁻³ –6.1 × 10 ⁴ cfu/g | | Chen et al., 2011a |
| spp. | EMA ^d LAMP | ND | 1 cfu/ml ^e | | Ye et al., 2008, 2011 |
| spp. | LAMP kit | ND | 10 ² cfu/ml | | He et al., 2010 |
| enterica serovar Enteritidis | LAMP | Reverse-transcription LAMP | Liquid egg | 2.2 cfu/test tube | Hara-Kudo et al., 2005 |
| Enteritidis | Reverse-transcription LAMP | Reverse-transcription LAMP | Liquid whole eggs | 10 ² cfu/25 ml; 10 ² cfu/25 ml after enrichment | Teethavanan and D'Souza, 2012 |
| <i>Typhimurium</i> | LAMP | Reverse-transcription LAMP | Pork products | 10 ² cfu/25 g after enrichment | Teethavanan et al., 2010 |
| <i>Typhimurium</i> | LAMP | Real-time reverse transcription LAMP | Pork processing environment | 10 cfu/ml | Teethavanan et al., 2011 |
| <i>Typhimurium</i> | hlyJ gene | LAMP | Fresh pork, whole chickens, green vegetables | 16 cfu/reaction | Zhang et al., 2012 |
| O9 serogroup | insertion element B2000/S1351 gene | LAMP | Chicken | 10 ² cfu/ml | Okamura et al., 2008 |
| O9 serogroup serotype D serovar Enteritidis | phoP gene | LAMP | Meat, milk | 35 cfu/250 ml | Li et al., 2009 |
| | prf gene | LAMP | Chicken meat | 10 cfu/reaction | Rajan and Vaidyanathan, 2012 |
| | sdh1 region | LAMP | ND | 4 copies/μl | Yang et al., 2010 |
| Escherichia coli | | | | | |
| (630 cfu/reaction serogroup O157, viable cells serotype O157:H7 serotype O157:H7 serotype O157:H7 serotype O157:H7 serotype O157:H7 serotype O157:H7 H7 serogroups O157, O26 and O111 STEC serotypes) | Enterohaemorrhagic <i>E. coli</i> (EHEC) | agf8 gene | LAMP | Radish sprouts, broccoli sprouts, ready-to-eat salads, ground pork and beef | |
| | Yitoyama et al., 2010 | rdh gene | LAMP | 410 cfu/ml | Wang et al., 2009b |
| | rdh gene | EMA-LAMP | Raw milk | 440 cfu/ml | Wang and Hao, 2012 |
| | rdh gene | LAMP | Pork meat | 10 cfu/ml | Jiang et al., 2012 |
| | sdh1 gene | LAMP | ND | 100 fg/reaction | Zhao et al., 2010a |
| | sdh2 gene | LAMP | ND | 100 fg/reaction | Zhu et al., 2009a |
| | rdh gene | LAMP | ND | 26 cfu/reaction | |
| | flc gene | LAMP | Meat | 26 cfu/reaction | |
| | sdh2 gene | LAMP | Meat | 1.8 cfu/g; 1.8 cfu/g | Liu et al., 2011a |
| | rdh gene flc gene | Duplex ^f LAMP | Beef, radish sprouts, alfalfa sprouts | 0.7 cfu/reaction | Hara-Kudo et al., 2007; Hara-Kudo et al., 2008a |
| | sdh1 gene sdh2 gene | Duplex ^f LAMP | ND | 10 ² cfu/reaction | Koguchi et al., 2010 |
| | sdh1 gene | Real-time duplex LAMP | Ground beef, beef trim, lettuce, spinach | 10–20 cfu/25 g after enrichment | Wang et al., 2012a |
| way gene | Shiga toxin-producing <i>E. coli</i> (STEC) | wzx gene | LAMP | 10–20 cfu/25 g after enrichment | Wang et al., 2012a |
| Vibrio | | | | | |
| | parhaemolysin | gyrB gene | LAMP | Raw fish | Sheil et al., 2011 |
| | | rdh gene | LAMP | Oysters | Tian et al., 2009 |
| | | rdh gene | LAMP without loop primers | ND | Nemoto et al., 2011 |
| | | rdh gene | LAMP | 5.3 × 10 ² cfu/g Shrimp | Li et al., 2008 |
| | | LAMP-LFD ^g | 5.3 × 10 ² cfu/g Shrimp | | Yamazaki et al., 2008a,b,c |
| | | EMA-LAMP | ND | | 9 fg/reaction |
| | | rdh gene | LAMP | 100 fg DNA/reaction | ND |
| | | rdh gene | 100 fg DNA/reaction | | Zhao et al., 2010a,b |
| | | rdh gene | LAMP | 0.8 cfu/reaction | Yamazaki et al., 2010 |
| | | rdh1 gene | LAMP | 21.3 cfu/reaction | Li et al., 2011 |
| | | rdh2 gene | LAMP | 5.0 cfu/reaction | Nemoto et al., 2011 |
| | | rdh3 gene | LAMP | 3.7 cfu/reaction | |
| | | rdh4 gene | LAMP | 4.50 cfu/reaction | |
| | | rdh5 gene | LAMP | 1.1, 1.3 × 10 ² cfu/g | Oyler and Ge, 2010 |
| | | rdh6 gene | LAMP-LFD | 11 cfu/reaction | Surasit et al., 2011 |
| | | wg gene | Real-time LAMP | 2.5 × 10 ² cfu/g | Han et al., 2011 |
| | | vh gene | LAMP | 10 ² cfu/g | Han and Ge, 2008 |
| | | vh gene | LAMP | 1–10 cfu/reaction | Han and Ge, 2010 |
| Real-time LAMP | | | | | |
| <i>cholerae</i> | ctx gene | LAMP | Human stool | 1.4 cfu/reaction | Yamazaki et al., 2008a,b,c |
| | ompW gene | LAMP | Shrimp | 2.2 × 10 ² cfu/g | Srinak et al., 2010 |
| | | LAMP | Freshwater, human stool, milk | 1.6 × 10 ² cfu/ml | 1.6 × 10 ² cfu/g |
| | | LAMP | Shrimps, crabs, oysters, meat and human diarrhea complex | 3.2 cfu/g | Xu et al., 2010a,b |
| <i>cholerae</i> serogroup O139 oligofactor | whfK gene | LAMP | ND | 63 cfu/reaction | Zhu et al., 2009b |
| | ompK gene | LAMP | Fish, shrimps, clams | 10 ² cfu/ml | Xu et al., 2012 |
| | ompK gene | LAMP | Water | 3.8 cfu/ml | Ding et al., 2009 |
| <i>Yersinia enterocolitica</i> | gyrB gene | LAMP | Pork | 65 cfu/ml | Gao et al., 2009 |
| | phoP gene | LAMP | Pork meat, milk | 2.3 cfu/100 g | Li et al., 2010b |
| <i>Cronobacter sakazakii</i> (<i>Enterobacter sakazakii</i>) | 16S-23S rRNA gene | LAMP | Infant formula | 1.2 cfu/100 g | Liu et al., 2009 |
| Shigella spp. | sdh4 gene | LAMP | Infant formula | 1.1 cfu/g | Hu et al., 2009 |
| <i>Shewanella putrefaciens</i> (spp. putrefaciens) | ITS region within putrefaciens rRNA gene | Reverse transcription LAMP | Raw milk | 6–8 cfu/ml | Wang et al., 2009 |
| | | LAMP | Fish | 5.0 copies/reaction | Li et al., 2012 |
| <i>Campylobacter coli</i> | cj0414 gene | LAMP-kit | Human stool | 5.6 × 10 ² cfu/g | Yamazaki et al., 2008b |
| | gyrB gene | LAMP | 4.8 × 10 ² cfu/g | | Yamazaki et al., 2009 |
| <i>Brucella abortus</i> | BCSP31 gene | LAMP | Liver and spleen tissues of mice, milk | 7.9 cfu/reaction | 3.8 cfu/reaction |
| | | LAMP | 4.9 × 10 ² cfu/ml | | Ohbuki et al., 2008 |
| <i>Leptospira</i> spp. | lipL41 gene | Real-time LAMP | ND | 100 copies/reaction | Lin et al., 2009 |
| | 16S rRNA gene | LAMP | Blood samples | 10 copies/reaction | Sonhayanon et al., 2011 |
| | 16S rRNA gene | LAMP | Urine from field rats, farmed pigs, farmed buffaloes | 10 ² cfu/ml | Kotsuma et al., 2012 |
| <i>Burkholderia pseudomallei</i> | BPS1-1406 gene | LAMP | Sputum, urine, throat swabs, blood, pus | 38 copies/reaction | Chantratta et al., 2008 |
| Gram positive bacterial pathogens | | | | | |

Table 1 (continued)

| Target species | DNA target | Assay type | Food sources | Detection limit | Reference | |
|--|----------------------------------|----------------|---|---------------------------------|---|------------------------|
| <i>Sapthococcus aureus</i> | fnrA gene | LAMP | Raw milk | 8–9 cfu/ml | Wang et al., 2008a | |
| | fnrA gene | LAMP | ND | 8–9 cfu/ml | Li et al., 2010a | |
| | fnrA gene | LAMP | Infant formula, meat, milk | 10.3 cfu/reaction | Yang et al., 2011 | |
| | fnrA gene | LAMP | Meat samples, eggs, dairy products | 42 cfu/g | Xu et al., 2010a,b | |
| | fnrA gene | LAMP | Pork meat | 10 ² cfu/g | Suwampani et al., 2011 | |
| | fnrA gene | LAMP | ND | 2.3 cfu/reaction | Goto et al., 2007 | |
| | fnrA gene | LAMP | 30.2 cfu/reaction | | | |
| | fnrA gene | LAMP | 6.0 cfu/reaction | | | |
| | fnrA gene | LAMP | 38 cfu/reaction | | | |
| | fnrA gene | LAMP | Street food, fermented food, bakery products, milk, milk products | 100 fg/reaction | Nagarajappa et al., 2012 | |
| | fnrA gene | LAMP | 100 fg/reaction | | | |
| | fnrA gene | LAMP | 100 fg/reaction | | | |
| | fnrA gene | LAMP | 100 fg/reaction | | | |
| | fnrA gene | LAMP | 100 fg/reaction | | | |
| <i>Listeria monocytogenes</i> | hlyA gene | LAMP | Chicken | 2.0 cfu/reaction | Tang et al., 2011 | |
| | hlyA gene | LAMP | ND | 100 cfu/reaction | Wang et al., 2011 | |
| | hlyA gene | Real-time LAMP | Spiced chicken, pork, ground beef, milk powder | 6 cfu/reaction | Shan et al., 2012 | |
| | hlyA gene | Real-time LAMP | 100 cfu/ml | | Wan et al., 2012 | |
| | hlyA gene | LAMP | Raw milk | 186 cfu/ml | Wang et al., 2010 | |
| | hlyA gene | LAMP | ND | ND | Yan et al., 2009 | |
| <i>Bacillus cereus</i> | hlyE gene | FTA-LAMP | Milk | 57 cfu/ml | Qi et al., 2009 | |
| | hlyE gene | LAMP | Milk | 1 cfu/ml | Liu et al., 2011b | |
| | hlyE gene | LAMP | Milk | 11 cfu/ml | | |
| <i>Clostridium botulinum</i> | BoNT/A gene | Real-time LAMP | Fish, honey | 1 cfu/reaction | Sakuma et al., 2009 | |
| | BoNT/B gene | Real-time LAMP | 10 cfu/reaction | | | |
| | parfingens Aicydlobacillus | qpc gene | LAMP | Meat | 3 cfu/g | Kaneho et al., 2011 |
| | addoerestrin | 16S rRNA gene | LAMP | Apple juice | | |
| 22.5 cfu/ml | Chen et al., 2011a,b | | | | | |
| Mycotoxin producers and fungal spoilage organisms | | | | | | |
| <i>Fusarium graminearum</i> | gokV gene | LAMP | Wheat | 2 pg DNA/reaction | Niessen and Vogel, 2010; Abdel-Hamid et al., 2011 | |
| <i>Fusarium</i> | corallin calumorum | hlyD gene | LAMP | Barley | 0.74 pg/reaction | Denschlag et al., 2012 |
| | graminearum | Real-time LAMP | Barley | 0.5% infection level | Denschlag et al., 2012 | |
| <i>Aspergillus</i> | act1 gene | LAMP | Peanut, Brazil nut, coffee bean | 2.4 pg/reaction | Luo et al., 2012 | |
| | nomis amy1 gene | LAMP | Peanut, Brazil nut, coffee bean | 7.6 pg/reaction | | |
| | parafiticus carbonatus | amy1 gene | LAMP | Peanut, Brazil nut, coffee bean | 20 pg/reaction | |
| | OTA producing (strains) | Acple gene | LAMP | Pure culture mycelia | 50 pg/reaction | Storani et al., 2013 |
| | niger (OTA producing strains) | Anple gene | LAMP | Pure culture mycelia | 50 pg/reaction | |
| <i>Deikera anomala</i> | ITS region within 5.8S rRNA gene | LAMP | Distilled water, wine, beer | 10 cfu/ml | Hayashi et al., 2007 | |
| | bruvllesis | | | 10 cfu/ml | | |
| | caustriensis | | | 10 cfu/ml | | |
| | naardensis | | | 10 cfu/ml | | |
| | bruvllesis | | | 10 cfu/ml | | |

^a LAMP – loop-mediated isothermal amplification under standard conditions as described in Notomi et al. (2000).

^b ND – not defined.

^c Commercially available.

^d Ethidium monoazide.

^e Propidium monoazide.

^f Reverse-transcription LAMP – master mix contains reverse transcriptase to provide a cDNA target for amplification during LAMP.

^g Duplex LAMP – two primer sets combined in one LAMP assay.

^h LFD – lateral flow device.

ⁱ Real-time LAMP – amplification of DNA is monitored continuously during the LAMP reaction by measuring turbidity or fluorescence.

^j LAMP assay using FTA (Fast Technology for Analysis of nucleic acids) membrane to bind target DNA.

Use of LAMP to detect bacteria and fungi in food



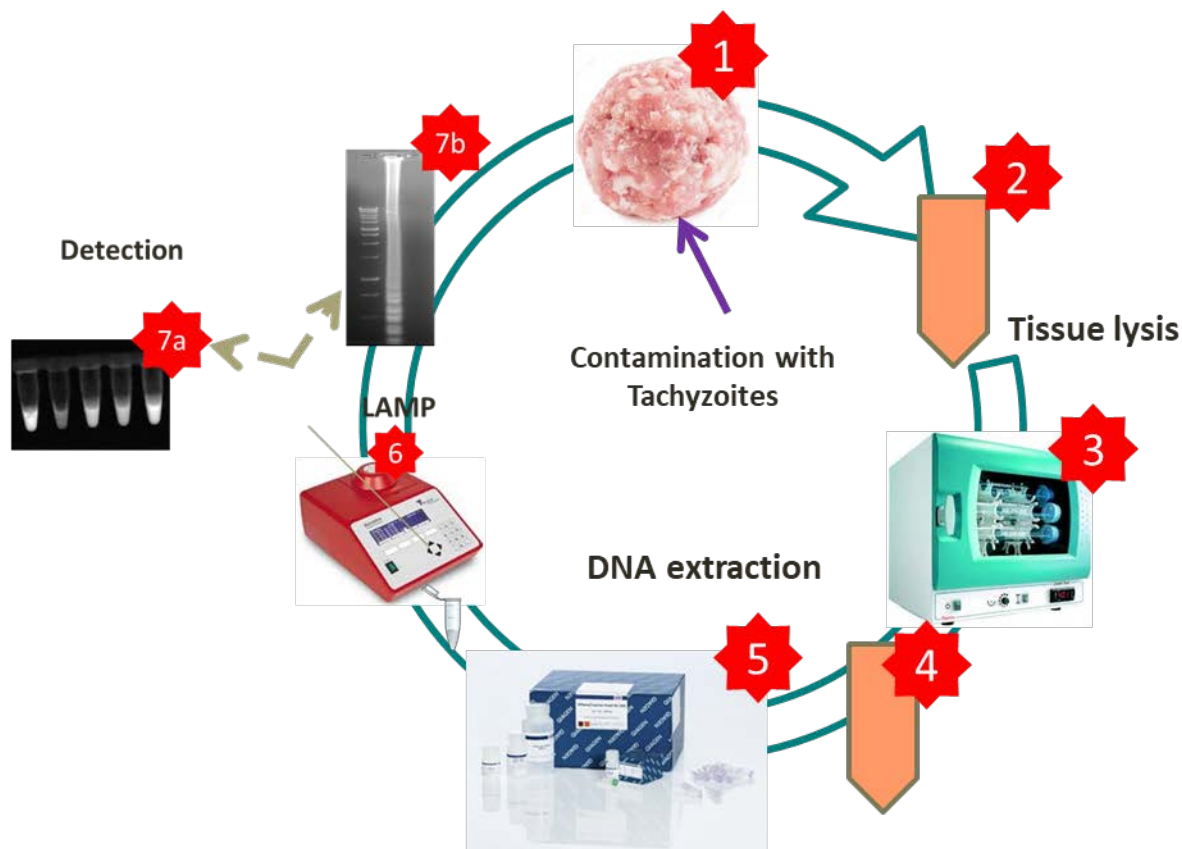
Toxoplasma detection by LAMP in food and environmental samples

| Target gene | sample type | parasite stage | sensitivity | comparison | ref. | year |
|------------------------------|--|----------------------------------|-------------------------|-----------------|---|------|
| 529 bp repetitive region | Leafy green vegetable | oocyst | 10 oocyst/50 g lettuce | LAMP=qPCR | Lalle et al., Food Microbiol. 2018 Apr;70:137-142 | 2018 |
| 529 bp repetitive region | formalin-fixed, paraffine embedded or frozen tissue from wild carnivores | tissue cysts | | | Suleman et al., J Vet Diagn Invest. 2016 Sep;28(5):536-42. | 2016 |
| SAG2 gene (single copy gene) | CNS homogenate and skeletal muscle of wild and farmed animal (25mg) | tissue cysts | 10 fg (~0.1 tachyzoite) | LAMP>PCR | Trisciuglio et al., J Vet Diagn Invest. 2015 Nov;27(6):754-7. | 2015 |
| ITS-1 (multicopy locus) | diaphragm samples obtained from pig | tissue cysts | 0.9 fg | LAMP>PCR | Zhuo et al., Veterinary Parasitology (2015) | 2015 |
| SAG1 | blood from experimentally infected pigs | tachyzoite (acute toxoplasmosis) | 100 fg (~1 tachyzoite) | LAMP<qPCR | Wang et al., Korean J Parasitol. 2013 Oct;51(5):573-7. | 2013 |
| MIC3 | soil | oocysts in the soil | 5 tachyzoites in soil | LAMP>PCR | Du et al., Veterinary Parasitology 184 (2012) 141–146 | 2012 |
| B1 (multicopy gene) | urine from experimentally infected mice | tachyzoite (acute toxoplasmosis) | 1 pg | LAMP>PCR | Hu et al., Folia Parasitol (Praha). 2012 Feb;59(1):21-6. | 2012 |
| 529 bp repetitive region | blood from experimentally infected mice | tachyzoite (acute toxoplasmosis) | 0.6 fg | LAMP529>LAMP_B1 | Kong et al. Parasites & Vectors 2012, 5:2 | 2012 |
| 529 bp repetitive region | blood samples from pigs and sheep | tachyzoite (acute toxoplasmosis) | 1 fg | LAMP>PCR | Lin et al., Veterinary Parasitology 185 (2012) 296–300 | 2011 |
| 529 bp repetitive region | lymph nodes from pigs | tissue cysts | 1 pg | LAMP>PCR | Zhang et al., Exp Parasitol. 2009 May;122(1):47-50. | 2009 |
| SAG1 | infected mouse organs | tissue cysts | 100 tachyzoites | LAMP>PCR | Krasteva et al., Vet Parasitol. 2009 Jun 10; 162(3-4):327-31. | 2009 |
| TgOWP and B1 | concentrated water | oocysts in water | 0.1 tachyzoite | LAMP=PCR | Sotiriadoua and Karanis. Diagnostic Microbiology and Infectious Disease 62 (2008) 357–365 | 2008 |

Aim

IDENTIFICATION OF DNA of *Toxoplasma gondii* IN FOOD MATRICES (MEAT AND MEAT PRODUCTS) BY LAMP

- ❖ Development of LAMP based procedure to detect *T. gondii* DNA (bradyzoites) in meat and meat products
- ❖ Evaluation of sensitivity and specificity of the LAMP assay



Materials

- ❖ *Toxoplasma gondii* tachyzoites of the RH stain
- ❖ gDNA from RH stain tachyzoites
- ❖ Pork, chicken, sheep and beef minced meat were used as matrices (5gr)
- ❖ LAMP assay targeting the 529 bp repetitive region (up to 300 copy per genome) was conducted using 6 primers (Zhang et al. 2009)

Toxoplasma gondii 529 repetitive region (GenBank: AF146527.1)

CTGCAGGGAGGAAGACGAAAGTTGTTTTTTTATTTTTTTTTCTTTTTGTTTTCTGATTTTTGTTTTTTTTGACTCGGGCCCAGCTGCGTCTGTCTGGGATGAGACCG
CGGAGCCGAAGTGCGTTTTCTTTTTTGGACTTTTTTTTTGTTTTTTCACAGGCAAGCTCGCCTGTGCTTGGAGCCACAGAAGGGACAGAAGTCGAAGGCGACTACAGA
CGCGATGCCTCCTCCAGCCGTCTTGGAGGAGAGATATCAGGACTGTAGATGAAGGCGAGGGTGAGGATGAGGGGGTGGCGTGGTTGGGAAGCGACGAGAGTCGGA
GAGGGAGAAGATGTTTTCCGCTTGGCTGCTTTTCTGGAGGGTGGAAAAAGAGACACCGGAATGCGATCCAGACGAGACGACGCTTTCTCGTGGTGATGGCGGAGA
GAATTGAAGAGTGGAGAAGAGGGCGAGGGAGACAGAGTCGGAGGCTTGGACGAAGGGAGGAGGGGTAGGAGAGGATCCAGATGCACTGTGTCTGCAG

ZF3 CCACAGAAGGGACAGAAGTC

ZB3 TCCGGTGTCTCTTTTTCCAC

ZFIP TCCTCACCTCGCCTTCATCTAGGACTACAGACGCGATGC

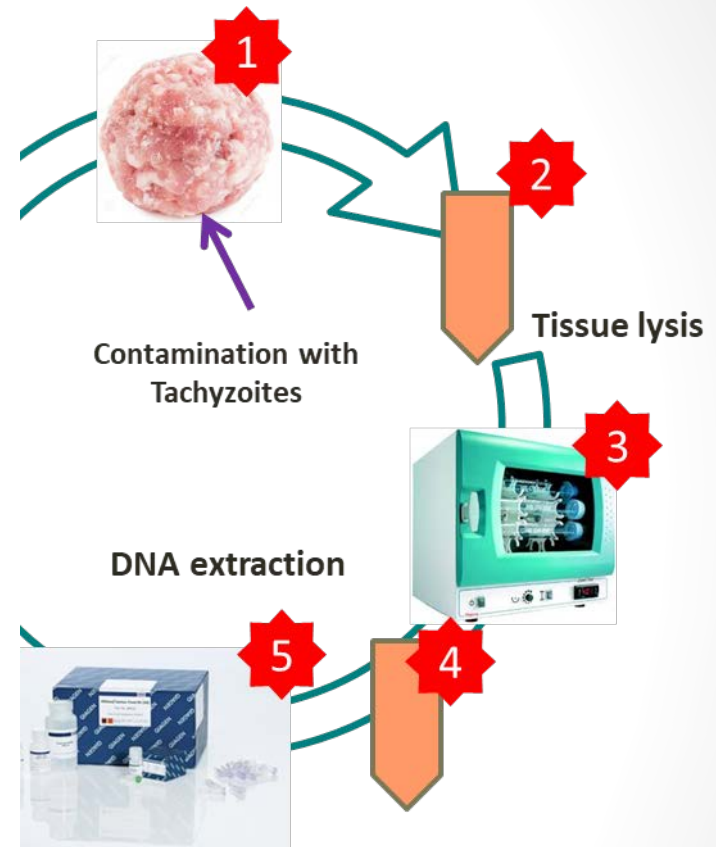
ZBIP TGGTTGGGAAGCGACGAGAGTCCAGGAAAAGCAGCCAAG

ZLF TCCAAGACGGCTGGAGGAG

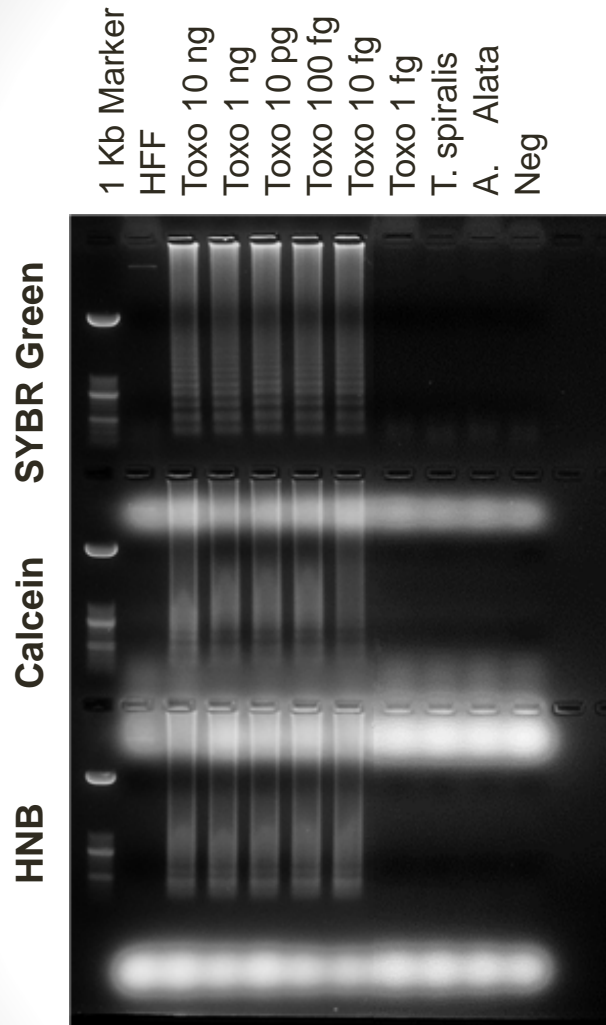
ZLB CGGAGAGGGAGAAGATGTTTTCC

Methods

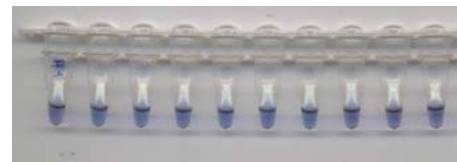
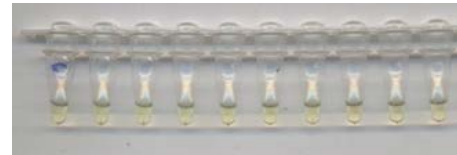
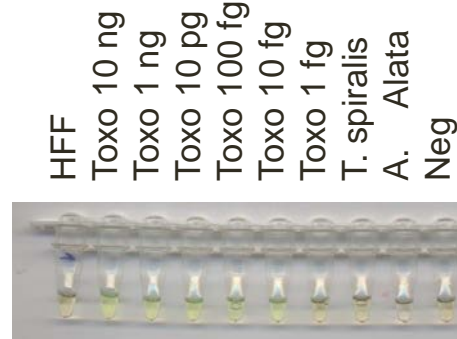
- ❖ DNA isolation from meatballs (5 gr) was performed combining homogenization with glass beads and tissue digestion with proteinase K at 55°C overnight (10 mL final volume) following Qiagen DNAeasy Tissue and blood kit DNA extraction (200 micL)
- ❖ LAMP sensitivity was determined using:
 - serial dilution of *T. gondii* RH DNA
 - DNA isolated from unspiked meatballs (pork, beef, lamb or chicken) and meatballs spiked with up to 10000 tachyzoites.



Detection of LAMP results with different visual methods

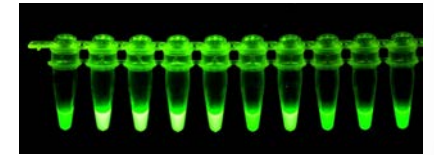
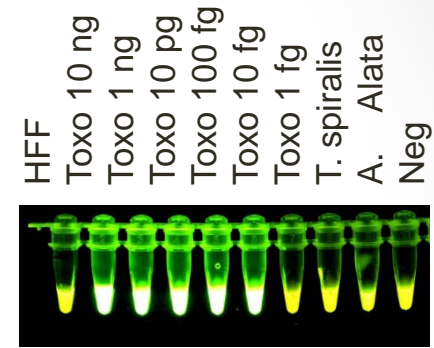


Specificity
No positive LAMP reaction was observed on gel electrophoresis using human, *T. spiralis* or *A. alata* gDNA



HNB (hydroxy naphthol blue)=metal ion indicator (Mg²⁺)

Sensitivity
Optimum condition for LAMP were 90 min at 63°C.
Detection limit 10 fg

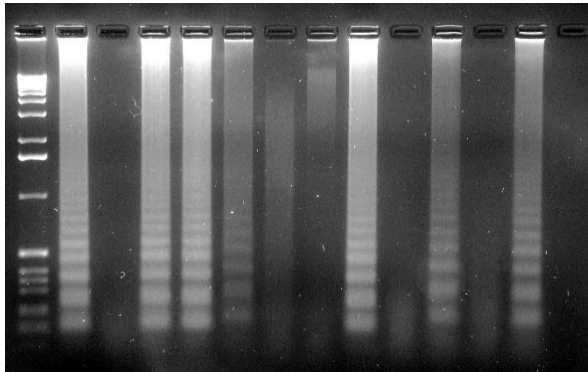


Spectrophotometric measurement at 650 nm

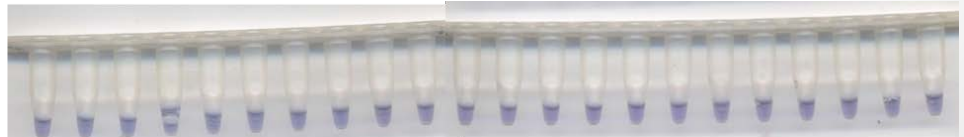
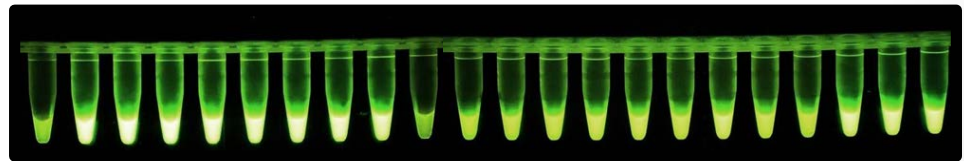
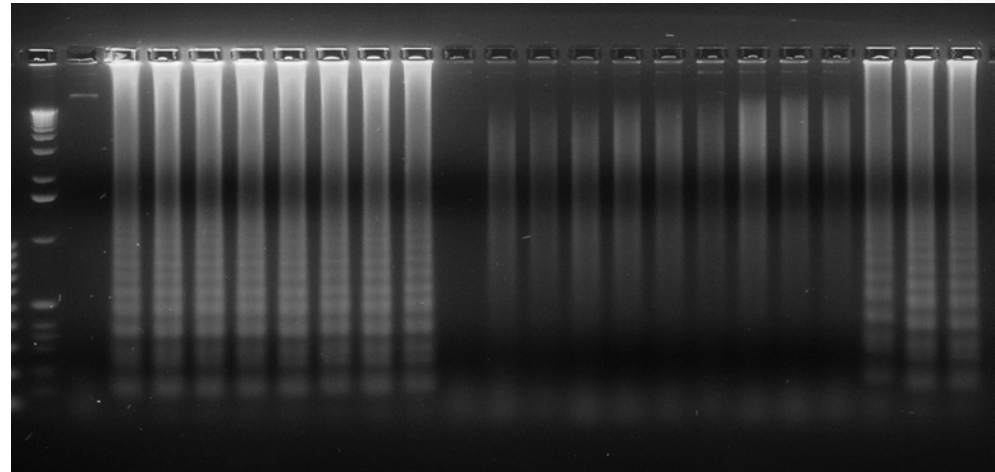
Detection performance
Both gel electrophoresis and SYBR green are suitable

Detection limit of tachyzoites contamination in meat by LAMP

1 Kb Marker
 Toxo 10 ng
 Neg
 Pork + 10000 tach
 Pork + 5000 tach
 Pork + 1000 tach
 Pork
 Lamb
 Lamb + 1000 tach
 Chicken
 Chicken + 1000 tach
 Beef
 Beef + 1000 tach



1 Kb Marker
 HFF
 Toxo 10 ng
 Toxo 1 ng
 Toxo 100 pg
 Toxo 10 pg
 Toxo 1 pg
 Toxo 100 fg
 Toxo 10 fg
 Toxo 10 fg
 Neg
 Pork
 Pork
 Pork
 Pork + 10 tach
 Pork + 10 tach
 Pork + 10 tach
 Pork + 100 tach
 Pork + 100 tach
 Pork + 100 tach
 Pork + 1000 tach
 Pork + 1000 tach
 Pork + 1000 tach



| | | | | | | | | | | | |
|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|
| 0,301 | 0,437 | 0,380 | 0,356 | 0,375 | 0,459 | 0,398 | 0,408 | 0,370 | 0,306 | | |
| 0,327 | 0,288 | 0,315 | 0,296 | 0,340 | 0,291 | 0,332 | 0,310 | 0,315 | 0,326 | 0,294 | 0,314 |

Specificity

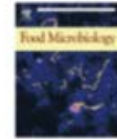
No positive LAMP reaction was observed on gel electrophoresis using as template DNA from unspiked matrices

Sensitivity

Detection limit 200 parasite/g of meat

Detection performance

SYBR green UV test was often prevented by high background fluorescence (high amount of DNA from matrix)



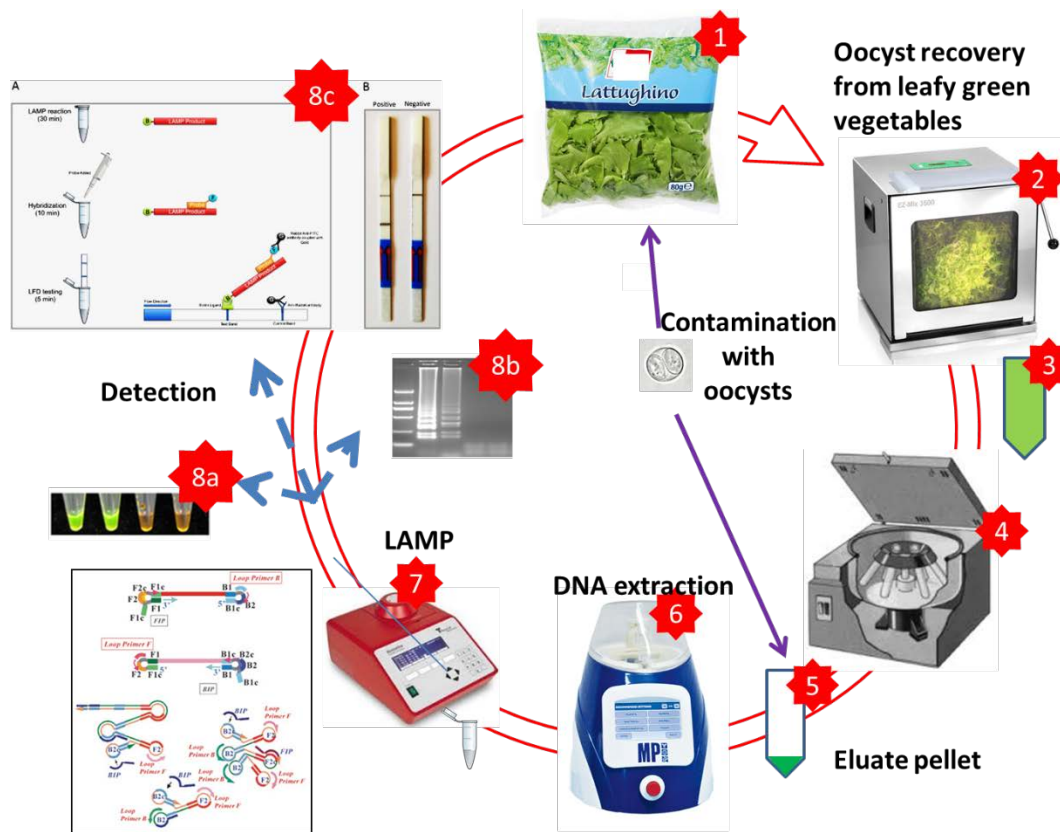
Aim

Loop-Mediated Isothermal Amplification-Lateral-Flow Dipstick (LAMP-LFD) to detect *Toxoplasma gondii* oocyst in ready-to-eat salad



Marco Lalle ^{a,*}, Alessia Possenti ^a, Jitender P. Dubey ^b, Edoardo Pozio ^a

- ❖ Development of LAMP-LFD based procedure to detect *T. gondii* oocysts in ready-to-eat leafy green vegetable.
- ❖ Evaluation of sensitivity and specificity of the LAMP-LFD assay



Materials

- ❖ *Toxoplasma gondii* oocysts of the VEG strain
- ❖ gDNA from tachyzoites of the RH strain
- ❖ Ready-to-eat baby lettuce was used as vegetable matrix
- ❖ LAMP assay targeting the 529 bp repetitive region (up to 300 copy per genome) was conducted using 6 primers (Zhang et al. 2009)
- ❖ For BIO-LAMP the a 5'-biotinylated BIP primer was used and a specific FITC-labeled DNA probe were designed to amplify and to hybridize with, respectively, the Bio-LAMP amplicon and being visualized on a chromatographic lateral flow dipstick (LFD) (Milenia® GenLine HybriDetect, GieBen, Germany).

Toxoplasma gondii 529 repetitive region (GenBank: AF146527.1)

```
CTGCAGGGAGGAAGACGAAAAGTTCTTTTTTATTTTTTTTTCTTTTTGTTTTCTGATTTTTGTTTTTTTTGACTCGGGCCCAGCTGCGTCTGTGGGATGAGACCG  
CGGAGCCGAAGTGCCTTTTTCTTTTTTACTTTTTTTTTGTTTTTTCACAGGCAAGCTCGCCTGTGCTTGGAGCCACAGAAGGGACAGAAGTCAAGGCGACTACAGA  
CGCGATGCGCTCCTCCAGCCGTCTTGGAGGAGAGATATCAGGACTGTAGATGAAGGCGAGGGTGAGGATGAGGGGGTGGCGTGGTTGGGAAGCGACGAGAGTCGGA  
GAGGGAGAAGATGTTTCCGGCTTGGCTGCTTTTCCTGGAAGGGTGGAAAAAGAGACACCGGAATGCGATCCAGACGAGACGACGCTTTTCTCGTGGTGATGGCGGAGA  
GAATTGAAGAGTGGAGAAGAGGGCGAGGGAGACAGAGTCGGAGGCTTGGACGAAGGGAGGAGGGGGTAGGAGAGGAATCCAGATGCACTGTGTCTGCAG
```

ZF3 CCACAGAAGGGACAGAAGTC

ZB3 TCCGGTGTCTCTTTTTCCAC

ZFIP TCCTCACCTCGCCTTCATCTAGGACTACAGACGCGATGC

ZBIP TGGTTGGGAAGCGACGAGAGTCCAGGAAAAGCAGCCAAG

ZLF TCCAAGACGGCTGGAGGAG

ZLB CGGAGAGGGAGAAGATGTTTCC

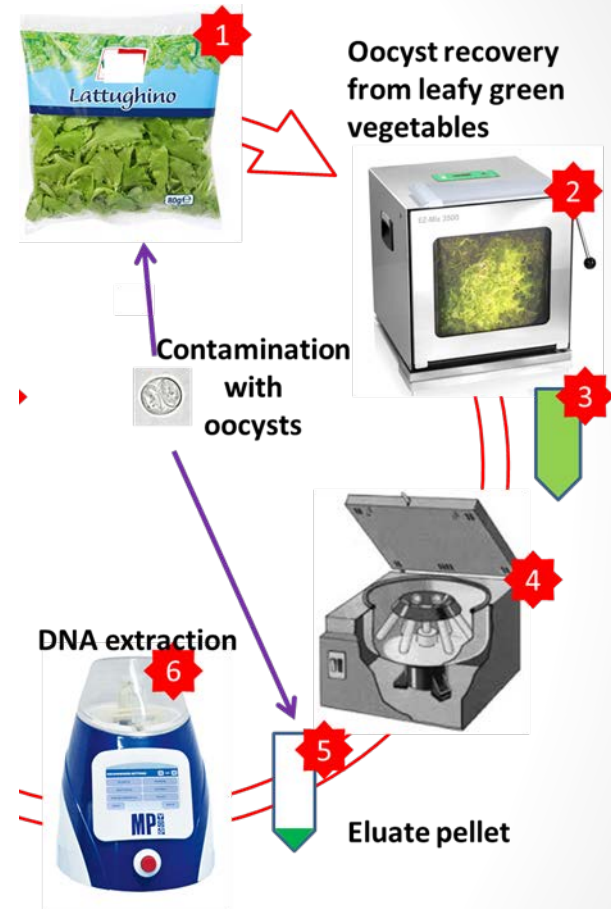
LAMP probe: FITC-5'-TGAGGGGGTGGCG-3'

bioZBIP: Biotin-5'-TGGTTGGGAAGCGACGAGAGTCGGAGAGGGAGAAGATGTTTCC-3'



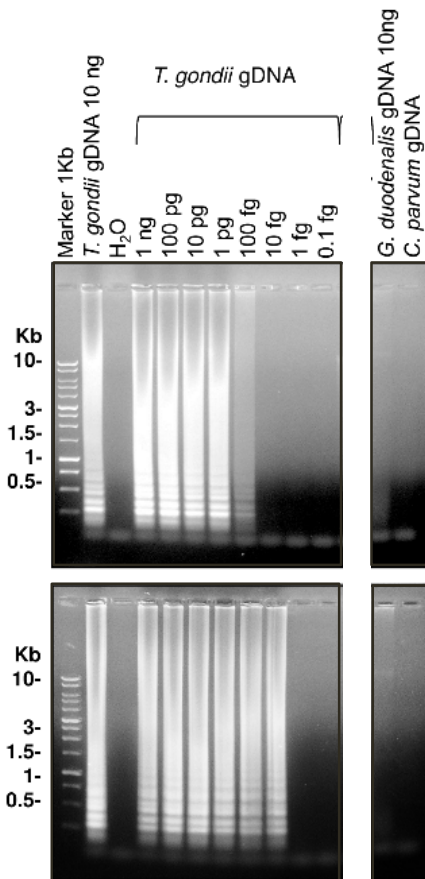
Methods

- ❖ A procedure for oocyst recovery was adapted from ISO 18744:2016 (*Detection and enumeration of Cryptosporidium and Giardia in fresh leafy green vegetables and berry fruits*)
- ❖ DNA isolation from vegetables with FastPrep 24 5G homogenizer and FastDNA™ SPIN Kit for Soil
- ❖ LAMP and Bio-LAMP-LFD sensitivity were determined using:
 - serial dilution of *T. gondii* RH DNA
 - DNA isolated from sediment of baby lettuce contaminated with *T. gondii* oocysts at different stages of sediment preparation: 5, 10, 50 oocysts/ml sediment or 10, 25, 50, 100 oocysts/50g of baby lettuce.



Optimization of LAMP

Addition after the reaction of the SYBR Green dye allow visualization of positive reaction directly in the vials. However, reaction negative vials containing DNA from control plant sediment fluoresce but a Ladder-like bands were evident in agarose gel electrophoresis only if the amplification product was present. **High amount of vegetable DNA background!**



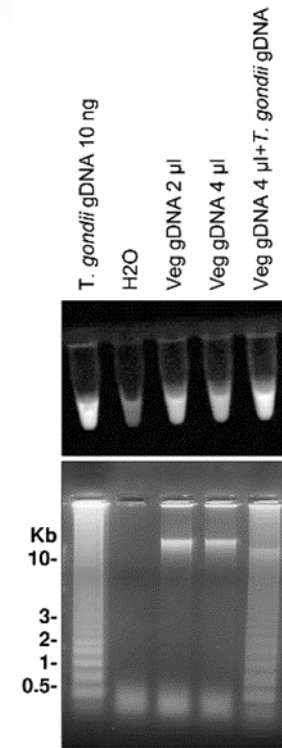
Specificity

No positive LAMP reaction was observed on gel electrophoresis using human, *C. parvum*, *G. duodenalis* or baby lettuce DNA!

Sensitivity

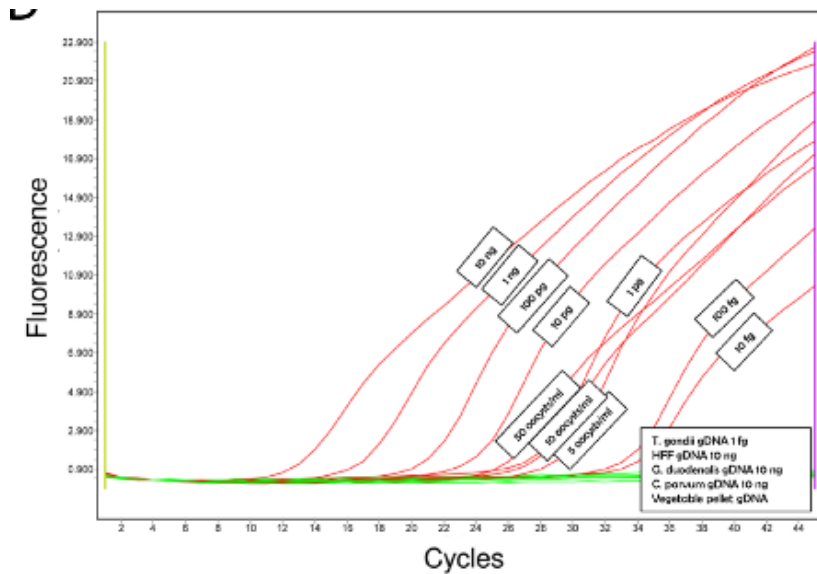
Optimum condition for LAMP were **2h at 63°C**, improving the detection limit! (B and C)

❖ From 100 fg to 10 fg of gDNA



LAMP vs qPCR

LAMP was as sensitive as qPCR, targeting the same locus, with a 6-FAM-labelled Taqman probe in LightCycler480 thermal-cycler (Opsteegh et al., 2010).



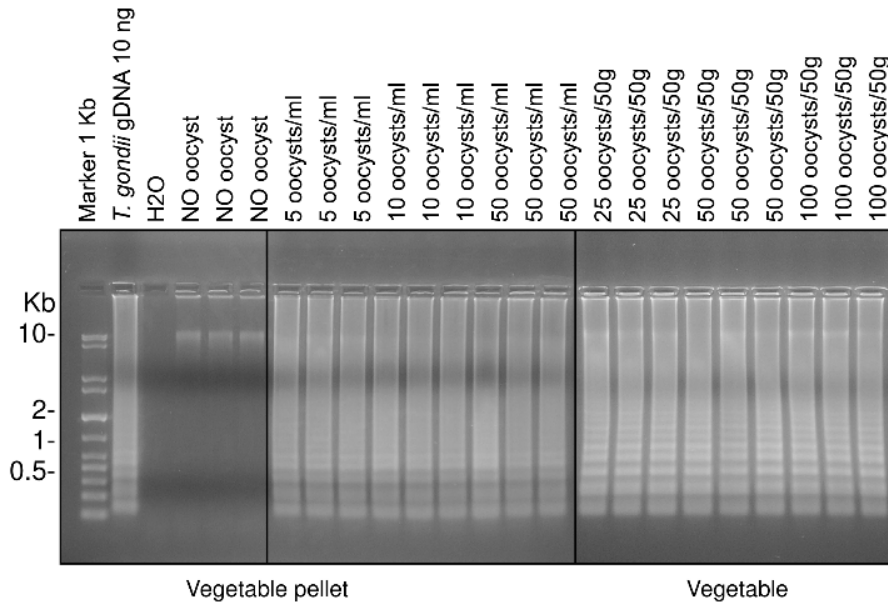
| Sample | <i>N. of T. gondii</i> haploid genomes | Mean Cp (\pm SD) |
|--|---|---------------------|
| HFF | NA | NA |
| <i>Giardia duodenalis</i> | NA | NA |
| <i>Cryptosporidium parvum</i> | NA | NA |
| <i>T. gondii</i> (RH strain) 10 ng | 140000 | 12.72 \pm 0.38 |
| <i>T. gondii</i> (RH strain) 1 ng | 14000 | 16.50 \pm 0.25 |
| <i>T. gondii</i> (RH strain) 0.1 ng | 1400 | 20.33 \pm 0.37 |
| <i>T. gondii</i> (RH strain) 10 pg | 140 | 23.70 \pm 0.38 |
| <i>T. gondii</i> (RH strain) 1 pg | 14 | 27.31 \pm 1.05 |
| <i>T. gondii</i> (RH strain) 0.1 pg | 1.4 | 31.07 \pm 1.33 |
| <i>T. gondii</i> (RH strain) 10 fg | 0.14 | 34.74 \pm 1.51 |
| <i>T. gondii</i> (RH strain) 1 fg | 0.014 | NA |
| Vegetable pellet suspension+50 oocysts | 16 | 26.19 \pm 1.12 |
| Vegetable pellet suspension+10 oocysts | 3.2 | 27.34 \pm 0.43 |
| Vegetable pellet suspension+5 oocysts | 1.6 | 27.65 \pm 1.47 |
| Vegetable pellet suspension | NA | NA |



Detection limit of oocysts contamination in vegetable by LAMP

LOD evaluation with baby lettuce artificially spiked with 25, 50 and 100 *T. gondii* oocysts per 50 grams of leaves (six samples for each condition).

Contaminated leaves were subjected to recovery procedure with glycine wash buffer and stomaching, following by DNA isolation.



| Sample type | LAMP positive ^a /Total |
|---------------------------|-----------------------------------|
| Leaves (50g) | |
| 25 oocysts | 5/6 (83%) |
| 50 oocysts | 6/6 (100%) |
| 100 oocysts | 6/6 (100%) |
| Pellet suspensions (1 mL) | |
| 5 oocysts | 4/6 (66%) |
| 10 oocysts | 5/6 (83%) |
| 50 oocysts | 6/6 (100%) |

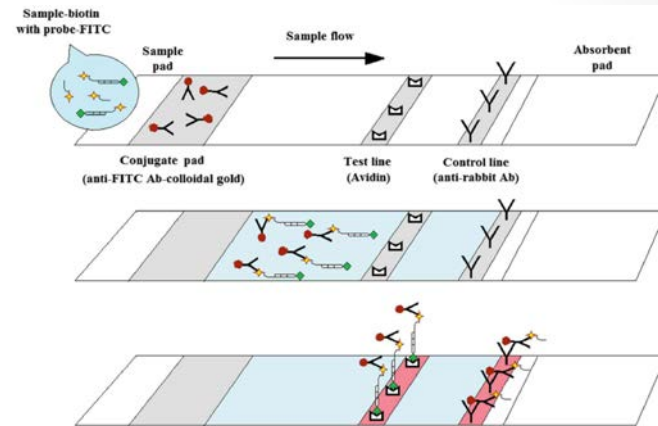
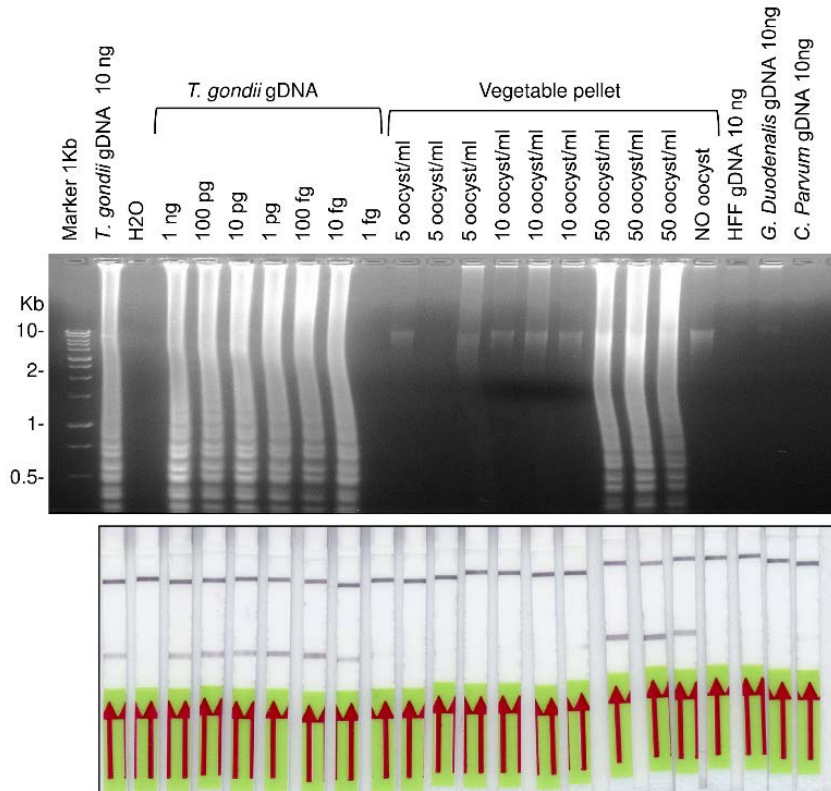
LAMP LOD 25 oocysts/50 gr (or 0.5 oocyst/gr) of contaminated leaves and 5 oocysts/mL of vegetable pellet suspension

In lettuce spiked with *T. gondii* oocysts, LOD was 10 to ≥ 100 oocysts per microliter by PCR targeting the B1 gene or the 529-bp repeat element (Marchioro et al., 2016)

A 33 oocysts/g LOD was reported using artificially contaminated basil applying immunomagnetic separation followed by qPCR targeting 529-bp repeat element (Hohweyer et al., 2016)

A LOD of 100 oocysts by B1 qPCR targeting was reported for an undefined amount of artificially contaminated radish (Lass et al., 2012).

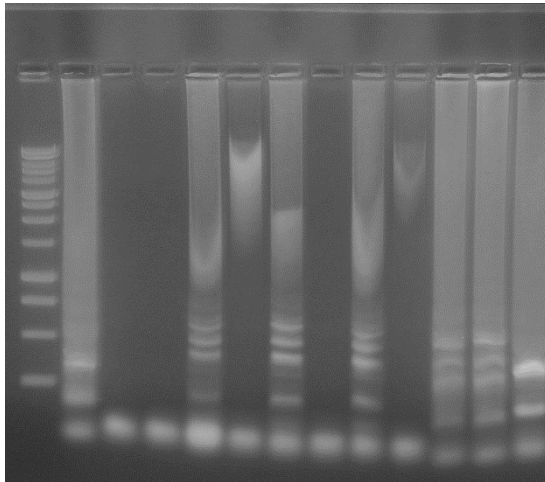
Faster visualization of LAMP by lateral flow deepstick



- ❖ BIO-LAMP and LAMP showed a comparable sensitivity on *T. gondii* gDNA
- ❖ BIO-LAMP was 10 times less efficient than LAMP (LOD 50 vs 5 oocysts) in detecting parasite gDNA in vegetable pellet suspension samples spiked with *T. gondii* oocysts

Detection of *T. gondii* in wild boar meat

1 Kb Marker
Toxo 10 ng
Neg
Neg
WB 307H1
WB 269H1
WB 146H1
WB 72H1
WB 48H1
WB 268H1
WB 269H1 spiked
WB 146H1 spiked
WB 72H1 spiked

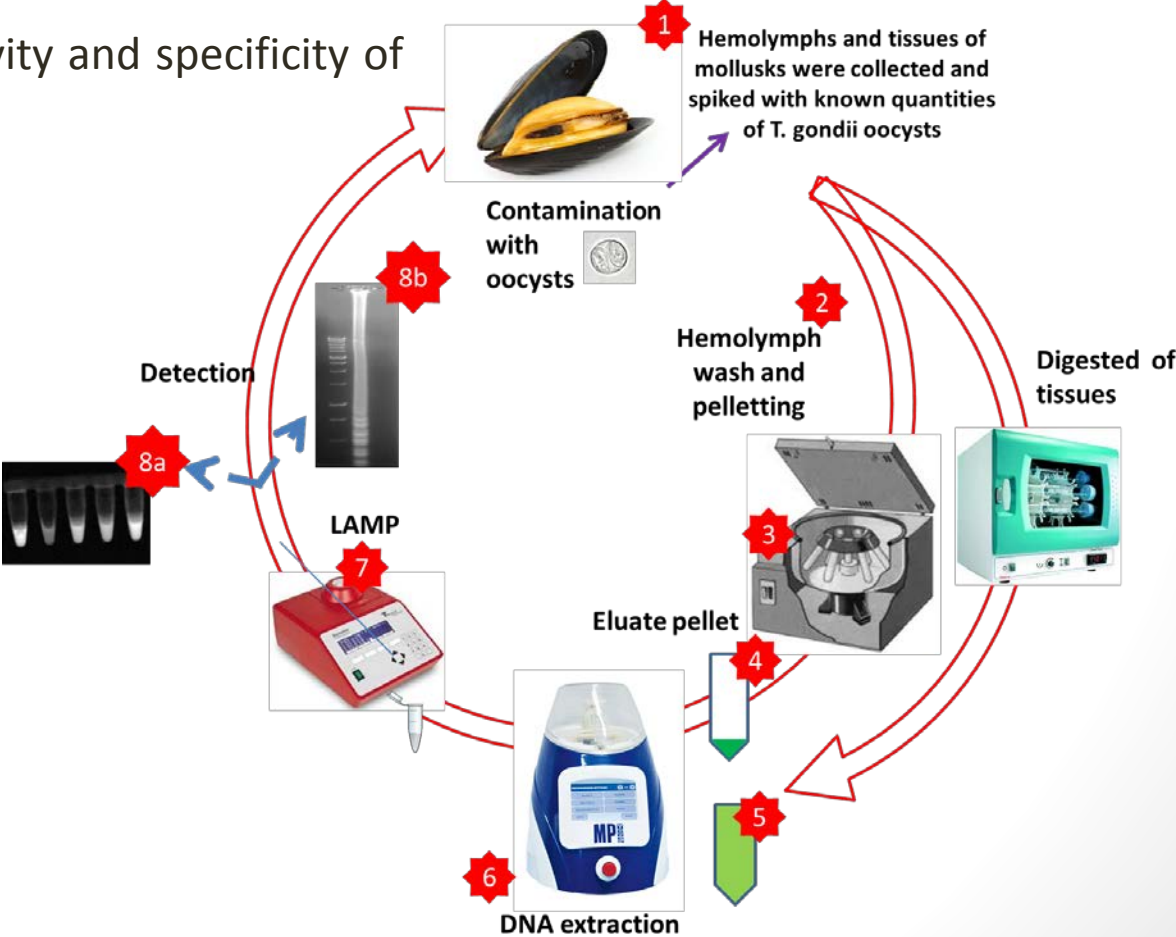


Main preliminary results

- Extending LAMP assay to 2 hours improved sensitivity
- DNA preparation do not contain inhibitors
- Samples will be further tested by qPCR for comparison with LAMP.

Aim

- ❖ Preliminary development of LAMP based procedure to detect *T. gondii* oocysts in mussels
- ❖ Evaluation of sensitivity and specificity of the LAMP assay



Materials

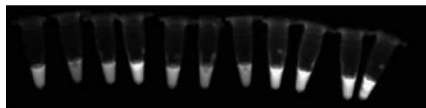
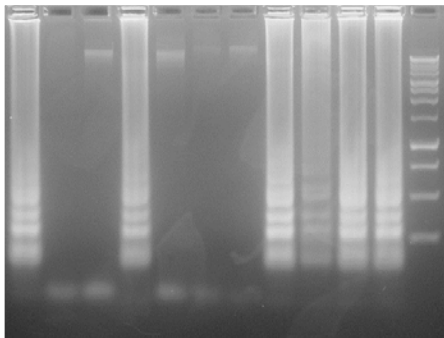
- ❖ *Toxoplasma gondii* oocysts of the VEG strain
- ❖ gDNA from tachyzoites of the RH strain
- ❖ Fresh blue mussels (hemolymph and tissue) as matrix
- ❖ LAMP assay targeting the 529 bp repetitive region (up to 300 copy per genome) was conducted using 6 primers (Zhang et al. 2009)

Methods

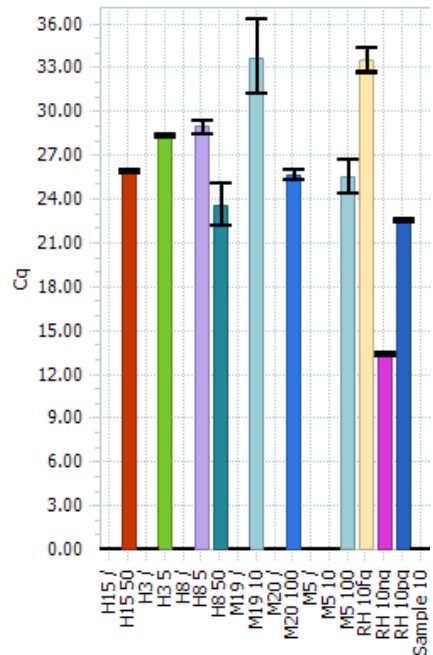
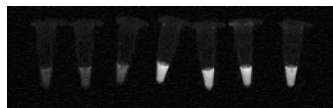
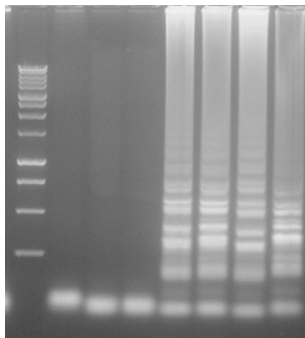
- ❖ DNA isolation from mussels tissue and emolymph with FastPrep 24 5G homogenizer and FastDNA™ SPIN Kit for Soil
- ❖ LAMP sensitivity was determined using:
 - DNA isolated from hemolymph and tissue contaminated or not with different amount of *T. gondii* oocysts 5 and 50 for 700 µl hemolymph, 10 and 100 oocysts/2g tissue.

Detection limit of oocysts contamination

- 1 : Pos control (20ng)
- 2 : Neg control
- 3 : Neg vegetable
- 4 : Pos vegetable
- 5 : Hemolymph 3 without spiking
- 6 : Hemolymph 8 without spiking
- 7 : Hemolymph 15 without spiking
- 8 : Hemolymph 3 with 5 oocysts
- 9 : Hemolymph 8 with 5 oocysts
- 10 : Hemolymph 15 with 50 oocysts
- 11 : Hemolymph 8 with 50 oocysts



- Mussel 5 without spikes
- Mussel 19 without spikes
- Mussel 20 without spikes
- Mussel 19 with 10 oocysts
- Mussel 5 with 10 oocysts
- Mussel 5 with 100 oocysts
- Mussel 20 with 100 oocysts



qPCR

LAMP LOD 10 oocysts/2 gr (or 5 oocyst/gr) of contaminated tissue and 5 oocysts/mL of washed hemolymph suspension

LAMP vs qPCR

| Sample ID | LAMP-PCR | | | | qPCR | | | | qPCR | | | | |
|-----------|---------------|----------|---------------|----------|---------------|--------|---------------|------|----------|--------|--------|-------|-----------|
| | LAMP-PCR (2h) | | LAMP-PCR (2h) | | LAMP-PCR (1h) | | LAMP-PCR (2h) | | 13/02/17 | | | | 14/02/217 |
| | 09/02/17 | 10/02/17 | 14/02/17 | 16/02/17 | Well 1 | Well 2 | Mean | SD | Well 1 | Well 2 | Well 3 | Mean | SD |
| H3 / | - | | - | | N/A | N/A | | | N/A | N/A | N/A | | |
| H3 5 | + | | + | | 27,81 | 27,37 | 27,59 | 0,22 | 28,36 | 28,08 | 28,35 | 28,26 | 0,13 |
| H8 / | - | | - | | N/A | N/A | | | N/A | N/A | N/A | | |
| H8 5 | + | | + | | 29,53 | 28,92 | 29,23 | 0,31 | 28,59 | 29,35 | 28,91 | 28,95 | 0,31 |
| H8 50 | + | | + | | 25,81 | 26,13 | 25,97 | 0,16 | 22,95 | 22,43 | 25,27 | 23,55 | 1,23 |
| H15 / | - | | - | | N/A | N/A | | | N/A | N/A | N/A | | |
| H15 50 | + | | + | | N/A | 26,27 | 26,27 | 0,00 | 25,84 | 25,95 | 25,94 | 25,91 | 0,05 |
| M5 / | | - | - | | N/A | N/A | | | N/A | N/A | N/A | | |
| M5 10 | | + | + | | N/A | N/A | | | N/A | N/A | N/A | | |
| M5 100 | | + | + | | 30,21 | NA | 30,21 | 0,00 | 26,87 | 25,16 | 24,38 | 25,47 | 1,04 |
| M19 / | | - | - | | N/A | N/A | | | N/A | N/A | N/A | | |
| M19 10 | | + | + | | N/A | N/A | | | 30,94 | 34,14 | 35,96 | 33,68 | 2,08 |
| M20 / | | - | - | | NA | 39,19 | 39,19 | 0,00 | N/A | N/A | N/A | | |
| M20 100 | | + | + | | 25,19 | 22,01 | 23,60 | 1,59 | 25,34 | 26,05 | 25,45 | 25,61 | 0,31 |

Preliminary results suggest that by applying the same extraction procedure and by targeting the same locus, LAMP is more sensitive and than qPCR especially to detect low amount of spiked oocyst in mussel tissue

Conclusions

- LAMP assay targeting the 529 bp repetitive region can be applied to detect *T. gondii* gDNA in meat, leafy green vegetables and mussels
- A well defined LOD was calculated in different conditions
- Despite qPCR is faster than LAMP (1h versus 3h), the comparable sensitivity of the two assays and the cheapest equipment required for LAMP makes the latest a valuable molecular test to be performed also in resource limited setting.
- Further optimization are necessary to apply BIO-LAMP to complex matrices as leafy green vegetable in order to provide a molecular detection tool more suitable to facilitate timely and effective food-borne disease outbreak investigations and response.

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Agricultural Research Service

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Stephanie la Carbona



Centre technique et d'expertise agroalimentaire

Kaya Stolberg
Nadja Bier



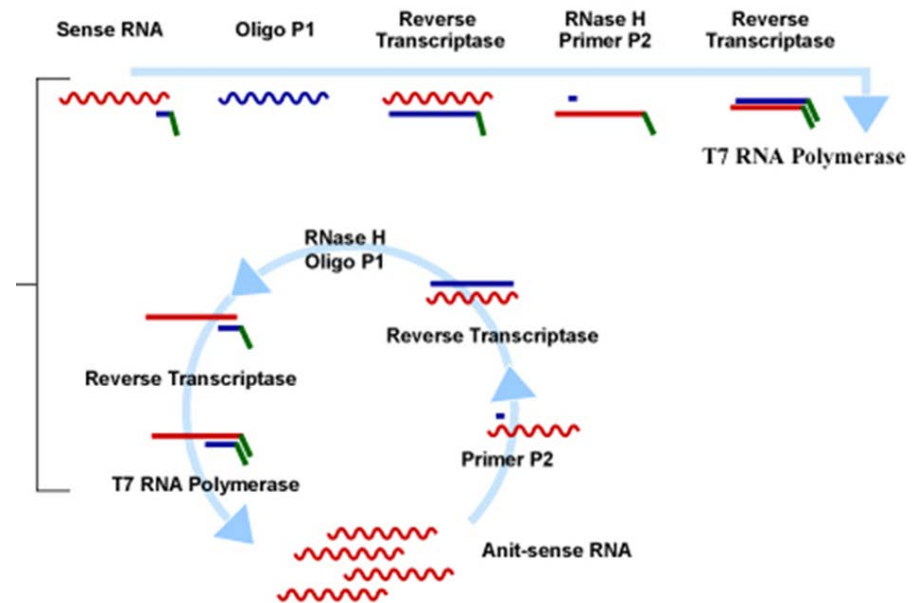
Federal Institute for Risk Assessment



Nucleic Acid Sequence-Based Amplification (NASBA)

- Specifically **designed to detect and amplify RNA** in the presence of DNA background without the use of a thermal cycler.
- 2 specific primers flanking the sequence to be amplified.
- 3 enzymes required: avian myeloblastosis virus reverse transcriptase (**AMV RT**), **RNase H**, and **T7 RNA polymerase**
- **Exponentially accumulate RNA/cDNA can be detected**

1. The primer P1 carries the T7 RNAPol binding sequence at its 5' end, it initiates the RT reaction
2. The RNase H then degrades the RNA strand in the RNA/cDNA hybrid molecules
3. Primer P2 binds cDNA and initiates the synthesis of the complementary strand.
4. T7 RNA Polymerase docks the double strand DNA on P1 5' end and transcribes RNA copies.
5. The process is repeated indeterminately starting from the newly transcribed RNA



❑ Advantages

- Lab equipment is less expensive
- not require a thermal cycler (a water bath or an isothermal block at 41°C is sufficient)
- High sensitivity ($>10^9$ copies of target in 90 min) compared to qPCR
- Suitable for lab-on-chip or portable apparatus
- Reduced sample prep compared with PCR due to less inhibition
- Can provide information of pathogen viability since the RNA is less stable than DNA.

❑ Disadvantages

- RNA integrity is the main cause of concern
- a single initial melting step is sometimes required.
- Temperature sensitive, use of thermolabile enzymes
- Amplified RNA target sequence 120–250 nucleotide

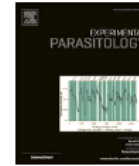


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Full length article

Real-time nucleic acid sequence-based amplification (NASBA) assay targeting *MIC1* for detection of *Cryptosporidium parvum* and *Cryptosporidium hominis* oocysts



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Both *C. parvum* and *C. hominis* are often associated with cryptosporidiosis in humans (humans are the main host for *C. hominis*, *C. parvum* is zoonotic and able to infect a variety of species.)

A real-time NASBA, targeting the *MIC1* transcript in *C. parvum* (CpMIC1) and *C. hominis* (ChMIC1).

LOD of *C. parvum* and *C. hominis* oocysts (down to 5 oocysts in 10 micl, and down to 1 oocyst using diluted RNA samples),

Distinguish between them. One of the primer sets targeted an exon only occurring in CpMIC1, distinguishing *C. parvum* from other *Cryptosporidium* spp.

NASBA assay detected *MIC1* mRNA also in inactivated oocysts. RNA does not seem to be a suitable marker for assessing oocyst viability.

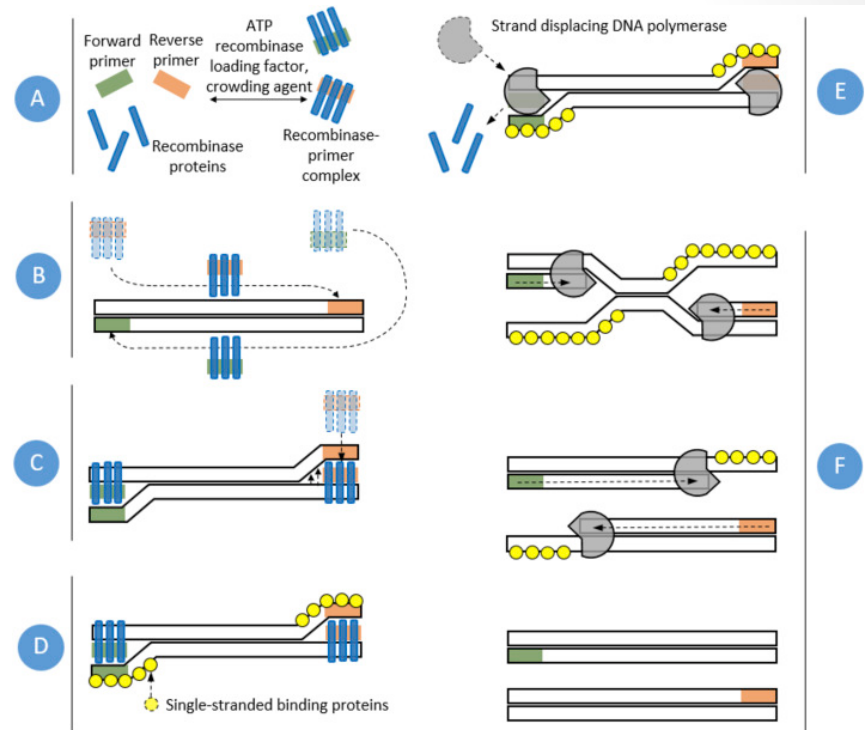


Recombinase polymerase amplification (RPA)

- RPA was first reported in 2006
- produced and under the intellectual license of TwistDx Limited (Cambridge, UK)

RPA utilizes:

- a bacterial recombinase, and a single-stranded DNA binding protein (SSB) to match primers to their target on the template DNA
- a *Staphylococcus aureus* Sau polymerase
- 2 primers of 30–35 mer (even shorter)
- ATP and a crowding agent (a high molecular polyethyleneglycol), forming a recombinase-primer complex.

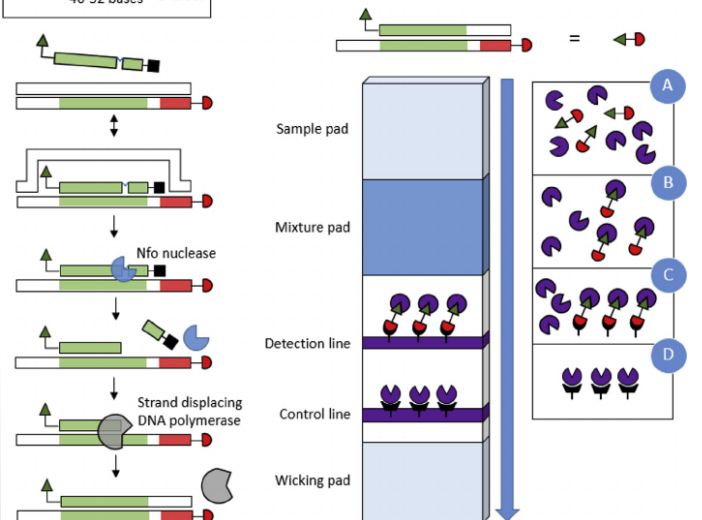
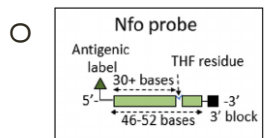
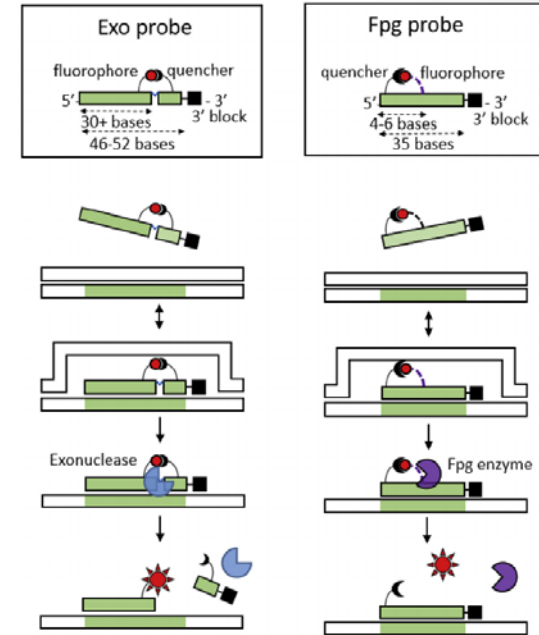


Lobato and O'Sullivan, Trends in Analytical Chemistry, 2018

- The recombinase enzyme forms a nucleoprotein complex with oligonucleotide primers and scans for homologous sequences in a DNA template.
- Recognition of a specific homologous sequence leads to the initiation of strand invasion and the opposing oligonucleotides are then extended by isothermal strand displacement amplification via Sau polymerase.

RPA can be monitored by end point detection (following amplification) or in real time (during amplification) and probes may be used depending on the detection strategy

- **The exo probe** (generally around 50 bases) contains an internal fluorophore and quencher. The quencher is cleaved by exonuclease III only upon binding to the template DNA. The probe contains a blocked 3' (not extendable)
- **The fpg probe** (generally 30–35 bases) contains a 5' quencher that is a few bases away from an abasic deoxyribose to which a fluorophore is linked. A nuclease (Fpg) specifically recognizes and cuts the linked fluorophore when the probe is annealed



The nfo probe here the quencher is cleaved by endonuclease IV (also called Nfo). This unblocks the probe 3' end of the and it can be extended by a polymerase to eventually join a segment amplified by a primer with 5' biotinylation. The biotinylated amplicon allows for easy application on a lateral flow

❑ Advantages

- the fastest isothermal amplification methods to date with minimal sample preparation, amplification of as low as 1–10 DNA target copies in less than 20 min
- Reaction is performed at 37–42°C with no requirement for a tight control of the temperature within this range
- Robustness in the presence of traditional PCR inhibitors (haemoglobin, ethanol and heparin)
- A reverse transcription step can be included to target RNA
- Multiplexing possible but highly dependent on target sequences, amplicon size and primer design
- The reagents necessary for RPA are sold in pellets, stable up to 6 months when stored at room temperature (22-28°C)

❑ Disadvantages

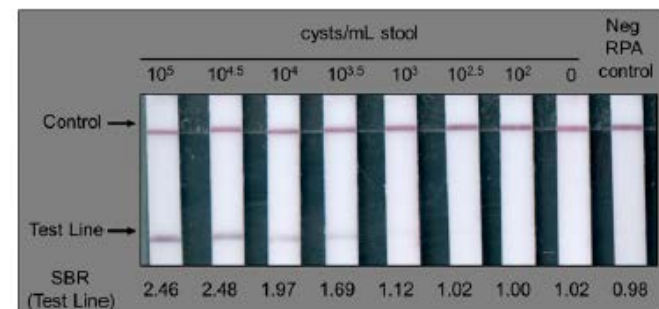
- Tolerance to mismatches can lead to cross-reactivity.
- Can be inhibited by high concentrations of genomic DNA,
- Kits are only sold by one company, (impact on pricing) and the user also has limited flexibility in the kit formulation.
- Requires purification/protein digestion following amplification (smearing or impaired flow in agarose gel electrophoresis and lateral flow)

Recombinase Polymerase Amplification-Based Assay to Diagnose *Giardia* in Stool Samples

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- RPA assay for *Giardia* utilizes DNA primers that target a unique 183 bp sequence on the beta giardin gene. Highly specific no cross amplification
- Detect as few as 10 *Giardia* cysts/ml spiked into PBS, similar to the analytical sensitivity of qPCR.
- The RPA assay detected both the A and B assemblages of *Giardia*
- Detected as few as 10^3 cysts per milliliter stool (50 cysts per reaction).
- RPA assay yielded 73% sensitivity and 95% specificity when tested using DNA extracted from clinical stool samples in comparison with gold standard (PCR microscopy)
- Lateral flow RPA has demonstrated a limit of detection of as few as 3000 *Giardia* cysts per gram of stool, slightly better than PCR-based assays



Conclusions

- There are a number of flexible, highly sensitive isothermal techniques that have been adapted to detect foodborne parasite at least in some matrices
- Most of the tests however are still confined to the labs where originally developed and still need to pass the test of wide and intense application to really evaluate their suitability in the field especially in the point-of-need (industry, slaughterhouse, farm)
- There are, however, limitations in the applicability of such techniques that will be difficult to overcome, as these are inherently linked to the nature of the samples under study.