

# Validation of the Detection Kit for Pathogenic Bacteria *Salmonella typhi* Causes Food Poisoning with Real Time Polymerase Chain Reaction

Irvan Maulana<sup>1,2</sup>, Muktiningsih Nurjayadi<sup>1,2,\*</sup>, Ulfi Rahma Efrianti<sup>1,2</sup>, Noer Azizah<sup>1,2</sup>, Nabilla Alya Pramudiyasih<sup>1,2</sup>, Ratna Nur Kusumawati<sup>1,2</sup>, Maharanianska Azzahra<sup>1,2</sup>, Jefferson Lynford Declan<sup>1,2</sup>, Gladys Indira Putri<sup>1,2</sup>, Dandi Akbar Juliansyah<sup>1,2</sup>, Ismaya Krisdawati<sup>1,2</sup>, Fera Kurniadewi<sup>1,2</sup>, Irma Ratna Kartika<sup>1,2</sup>, Dalia Sukmawati<sup>3,2</sup>, Lidwina Nastassya<sup>4</sup>, Vira Saamia<sup>5</sup>, Dwi Anna Oktaviani Saputro<sup>5</sup>, I Made Wiranatha<sup>5</sup>, Hesham Ali El-Enshasy<sup>6,7,8</sup>

<sup>1</sup>Department of Chemistry, Faculty of Mathematics and Natural Science, Universitas Negeri Jakarta, Gedung KH. Hasjim Asj'ari, 6th Floor, Jl. Rawamangun Muka, Jakarta Timur, 13220, Indonesia

<sup>2</sup>Research Center for Detection of Pathogenic Bacteria, Lembaga Penelitian dan Pengabdian Kepada Masyarakat, Universitas Negeri Jakarta, Jl. Rawamangun Muka, Jakarta Timur 13220, Indonesia

<sup>3</sup>Department of Biology, Faculty of Mathematics and Natural Science, Universitas Negeri Jakarta, Gedung KH. Hasjim Asj'ari, 6th Floor, Jl. Rawamangun Muka Jakarta Timur, 13220, Indonesia.

<sup>4</sup>Center for Agricultural Quarantine Standards Test, Jl. Pemuda Rawamangun, Jakarta Timur, 13220, Indonesia

<sup>5</sup>Center Forensic Laboratory of the Criminal Investigation, Police of the Republic of Indonesia, Cipambuan, Babakan Madang, Bogor, West Java 16810

<sup>6</sup>Institute of Bioproduct Development (IBD), Universiti Teknologi Malaysia (UTM), Skudai, Johor Bahru, Johor, Malaysia.

<sup>7</sup>School of Chemical and Energy Engineering, Universiti Teknologi Malaysia (UTM), Skudai, Johor Bahru, Johor, Malaysia.

<sup>8</sup>City of Scientific Research and Technology Applications, New Burg Al Arab, Alexandria, Egypt.

**Abstract.** *Salmonella typhi* is a gram-negative bacteria that causes food poisoning. *Salmonella typhi* bacteria can cause typhus, which can produce lipopolysaccharide complex endotoxin, which plays an important role in the pathogenesis of typhoid fever. Cases of food poisoning are still common and are one of the causes of death and death in Indonesia. In 2019, there were 77 cases of food poisoning in Indonesia. A fast, accurate, and specific detection method is needed to detect poisoning. In previous studies, the optimum conditions and formulas for detecting these bacteria have been obtained using Real Time PCR. The results of previous studies are used as the basis for the development of a prototype detection kit. In this study, validation was carried out, which aimed to confirm the results of previous studies so that a reproducible and accurate product for the detection of *Salmonella typhi* bacteria could be obtained. The results of this study showed that fim-C primers for *Salmonella typhi* were amplified at 95 base pairs (bp) with an annealing temperature of 60°C and a standard DNA concentration of 50 ng/μL. The results of the Real Time PCR confirmation test of *Salmonella typhi* bacteria at a pre-denaturation of 3 minutes with a concentration of 10 pmol obtained the Ct value according to the standard with previous studies. The Ct value obtained was 13.96 for *S. typhi* bacteria. Based on the results obtained, it can be concluded that the condition validation stage for pure cultures was successfully carried out by producing consistent and reproducible data.

\* Corresponding author: [muktiningsih@unj.ac.id](mailto:muktiningsih@unj.ac.id)

## 1 Introduction

Based on the report from the General Center for Food and Drug Administration, cases of food poisoning in Indonesia in 2019 reached 77 cases, with a total of 7,244 people exposed [1]. Food poisoning cases can be caused by microorganisms, one of which is the *Salmonella typhi* bacteria. *Salmonella typhi* bacteria is a pathogenic bacteria that is classified as gram-negative bacteria that can cause typhus in humans and is transmitted through food or drink [2][3]. *Salmonella typhi* bacteria can spread through contaminated foods or drinks. After entering the body, these bacteria will spread through the blood to the lymph glands, intestines, liver, and spleen, where they will multiply [4] In general, cases of food poisoning caused by *Salmonella* bacteria are more common in children under the age of 5, which causes 93.8 million children under the age of 5 to suffer from salmonellosis worldwide [5].

In Indonesia, many cases of food poisoning occur due to food made in households, services, factory processing, restaurants, households, and unlisted household industries [1]. In addition to bacteria, viruses and parasites, food poisoning can be caused by improper food storage, improper cooking and food contamination [6]. *Salmonella*, *E. coli*, *Campylobacter*, *Shigella*, *Listeria*, *Vibrio*, *Yersenia*, and *Staphylococcus* are common bacteria that can cause food poisoning [7].

Researchers have developed a variety of techniques to identify cases of food poisoning, but these incidents are still responsible for many deaths and painful illnesses in Indonesia. They are a very significant problem that can lead to unusual cases of food poisoning [3]. To find the source of food poisoning and lower the risk of death, this calls for swift, precise, and focused detection procedures. Real-time polymerase chain reaction techniques were used in the short-mixing methods based on nucleic acid research, that is, approaches that use template hybridization processes with primary or short nucleotide strands.

This study is part of the UNJ *Salmonella* research team. Previous research has successfully detected *Salmonella typhi* bacteria using a real-time polymerase chain reaction method. The principle of real-time PCR is that the amount of DNA can be measured through fluorescent signals, which can enhance the fluorescence signal and are directly proportional to the amplification of the DNA amount [8][9].

The results obtained in previous studies were successful in designing specific primers for *Salmonella typhi* and *Salmonella typhimurium* bacteria [10][11]. In this research, formula validation will be carried out to produce a prototype detection kit to detect *Salmonella typhi* bacteria, which cause food poisoning. Validation aims to confirm and improve the results of previous studies so that a reproducible product is obtained.

## 2 Material and Methods

### 2.1 Bacterial Culture

Culture of *Salmonella typhi* bacteria on Blood Agar media using a specific medium, namely *Salmonella Shigella* Agar. Bacterial culture was carried out by streaking a zig-zag (streak plate) with an ose needle and instilling an ose of bacteria on glycerol stock into Blood Agar media. then incubated for 1 day at 37 °C (overnight culture); after incubation, a colony of *S. typhi* bacteria was formed.

After that, the single colonies that formed on the agar medium were taken with a sterile loop and dipped into 10 mL of liquid medium, the liquid medium used is lactose broth. then incubated at 37 °C for 18 hours (overnight culture) with a shaker speed of 150 rpm until it produced turbidity.

### 2.2 Isolation of *Salmonella typhi* bacterial DNA

Isolation of DNA from pure cultures of bacteria was carried out using the commercial. The process of isolating gram-negative bacteria follows the protocol contained in the handbook kit. The DNase membran combine micro spin technology with the binding characteristics of a silica gel based membrane. The buffer conditions used in the DNeasy extraction technique are created to enable the adsorption of DNA specific to the silica-gel membrane and to provide the best possible removal of carbohydrates, polyphenols, and other plant metabolites [12]. After that, the isolated results were characterized by agarose gel electrophoresis and quantified using a nanodrop spectrophotometer.

### 2.3 Validation of the annealing temperature of the *fimC* gene of *Salmonella typhi* bacteria by PCR gradient

Primer annealing temperature validation was carried out using a Gradient PCR instrument. This process was carried out at an annealing temperature of 60 °C according to previous studies. At this stage it starts from making a recipe for the gradient PCR amplification process with a total volume of 25 µL of one reaction mixture containing a pair of primers (forward and reverse) of the *fim-C* gene, a template of bacterial DNA isolate *S. typhi* Colorless Master Mix and Nuclease Free water (NFW). The amplification results were characterized by 2% agarose gel electrophoresis and the DNA bands were observed with a UV transilluminator.

## 2.4 Validation of the confirmation test for the *fimC* gene of *Salmonella typhi* bacteria by real-time PCR

Confirmation tests were carried out using purecultured DNA isolates of *S. typhi* bacteria as a positive control and NTC (No Template Control) as a negative control. A mixture of positive and negative control reactions containing 20 µL each was made, which contained the master mix, primers (forward and reverse), bacterial DNA isolates, and Nuclease Free Water (NFW). However, negative controls were made without DNA isolates.

The real-time PCR program used included initial denaturation at 95°C for 3 minutes, denaturation at 95°C for 10 seconds, annealing at 60°C for 30 seconds, extension at 72°C for 30 seconds, and repetition for 40 PCR cycles. However, this study varied the predenaturation or initial denaturation time from 3, 4, and 5 minutes and varied the primary concentrations from 5, 10, and 15pmol.

## 3 Result and Discussion

### 3.1 Bacterial Culture

Bacterial culture on agar media is the first step in this research. *Salmonella typhi* bacteria were cultured on Salmonella Shigella Agar media. Culture on SSA media resulted in the formation of black colonies. This happens because SSA media contain bile salts, brilliant green, and sodium citrate to inhibit the growth of gram-positive bacteria. In addition, *Salmonella* bacteria cannot ferment lactose, but can reduce sodium thiosulfate in SSA media to produce sulfite and H<sub>2</sub>S gas [13]. The resultscan be seen in figure 1.



**Fig. 1.** Culturing bacteria on Salmonella-Shigella agar media

Lactose broth is the medium used after successful culture on agar and then culture on liquid media. Lactose is one carbohydrate that can differentiate to produce coliform. Gram-negative basil and optional aerobic and anaerobic spores are

members of the coliform group. They digest lactose with gas production in 48 hours at a temperature of 35 °C [14]. The result that is formed in liquid media is turbidity on the media, which indicates that the bacteria have succeeded in growing [15]. The purpose of culturing in liquid media is to enrich the growth of bacteria so that it can easily isolate bacterial DNA. The results can be seen in figure 2.



**Fig. 2.** Culturing bacteria on Lactose Broth

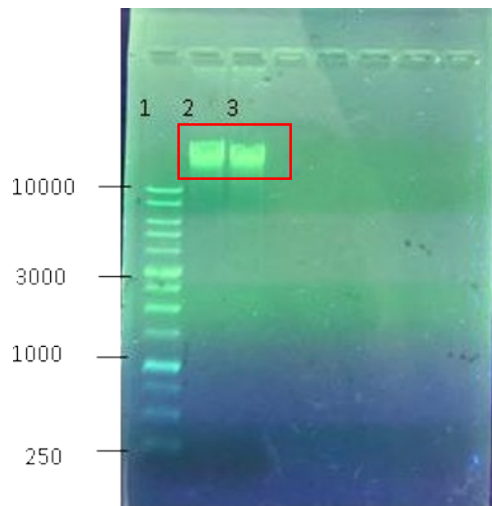
### 3.2 DNA Isolation

DNA isolation aims to obtain pure DNA isolates without contamination of protein, RNA, or other components. The results of DNA isolation were subjected to quantitative analysis using a nanodrop spectrophotometer, to determine the concentration and purity of the isolated DNA. The concentration and purity of DNA isolates can be seen in Table 1.

**Table 1.** Results of nanodrop spectrophotometer measurements

Sample	Concentration (ng/µL)	Purity (A260/280)
<i>Salmonella typhi</i>	49	1.837

Based on the table above, the purity of the resulting DNA isolates was 1.837. The purity of DNA isolates is said to be good, when the purity value is between 1.8-2.0 if the purity is below 1.8, then the isolate is contaminated with protein, if the purity is above 2.0, then the isolate is contaminated with RNA then the concentration of DNA used for real-time PCR testing has a minimum concentration of 0.1–500 ng [16]. The isolation results were then characterized by electrophoresis on 2% agarose, which was used to determine whether or not there was bacterial DNA present [17]. The results of agarose electrophoresis can be seen in Figure 3.

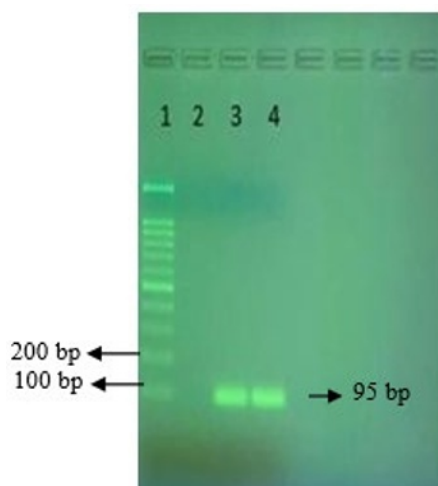


**Fig. 3.** Agarose Electrophoresis: (1) DNA ladder 1 kb; (2) Bacterial DNA isolates of *S. typhi*; (3) Bacterial DNA isolates of *S. typhi*.

Based on the results of the above agarose gel electrophoresis a band was formed above the 1 kb ladder. This is due to the larger genome size of *Salmonella typhi*  $5 \times 10^6$  [18]. Based on these results, it can be stated that bacterial DNA isolation has been successful.

### 3.3 Validation of the annealing temperature of the fim C gene of salmonella typhi Bacteria by PCR gradient

In previous research, annealing temperature optimization has been carried out using gradient PCR with a temperature range of 56–61 °C. The selection of this temperature is based on the melting temperature when the primer is synthesized, and usually the temperature range for this optimization is 5°C below the  $T_m$  of the primer [8][19][20]. The optimum temperature for *Salmonella typhi* bacteria in this temperature range is 60 °C [10]. The results of validation annealing temperature can be seen in Figure 4.



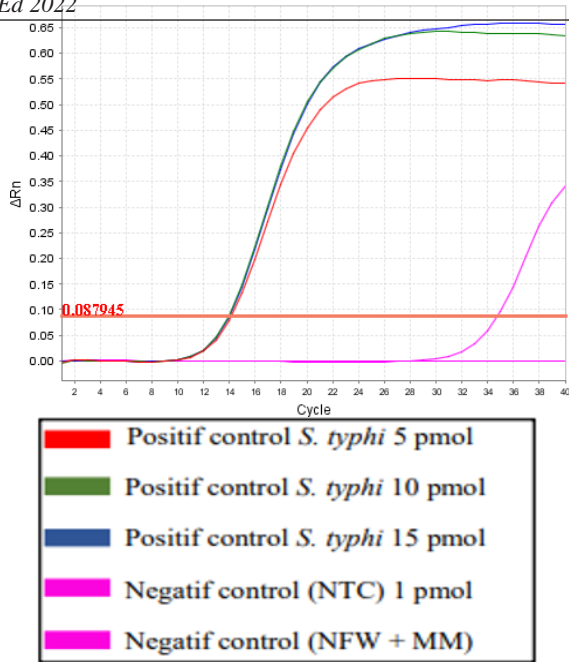
**Fig. 4.** Result of validation annealing temperature (1) DNA Ladder 100 bp; (2) NTC; (3)(4) Fragment fim-C gene of bacterial *S. typhi*;

Based on the results obtained, it can be seen that there is a band on the fim-C gene fragment of *Salmonella typhi* bacteria with an amplicon size of 95 bp at 60 °C. From these results it can be concluded that the annealing temperature validation is in accordance with the results of previous studies which showed that there was a band with an amplicon size of 95 bp [10].

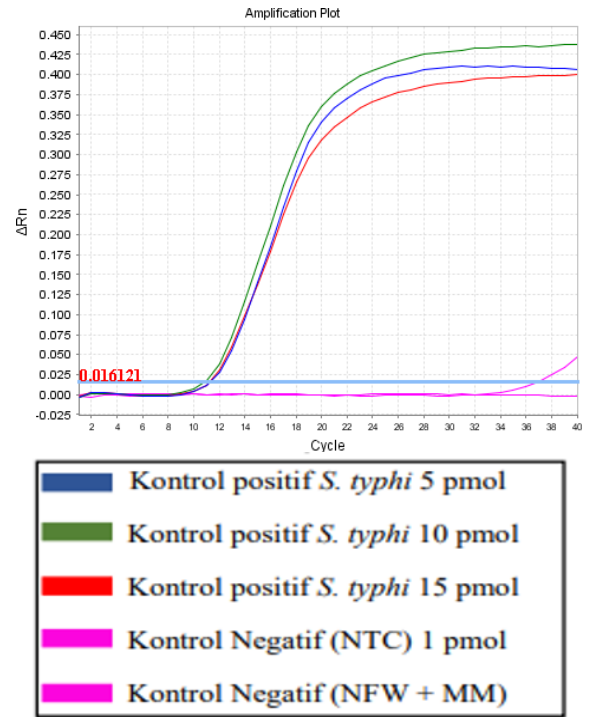
Validation of the annealing temperature is necessary because it is a step that influences the success of the PCR amplification process. In addition, this step is also used for attachment of primers to the template DNA, so this attachment process requires optimum timing, the temperature is too high, it will cause amplification failure because the primer does not stick to the template DNA; if it is too low, it will make the primer stick to the other side of the genome, resulting in DNA that is formed having low specificity [19].

### 3.4 Validation of the confirmation test for the fimc gene of Salmonella typhi bacteria by Real Time PCR

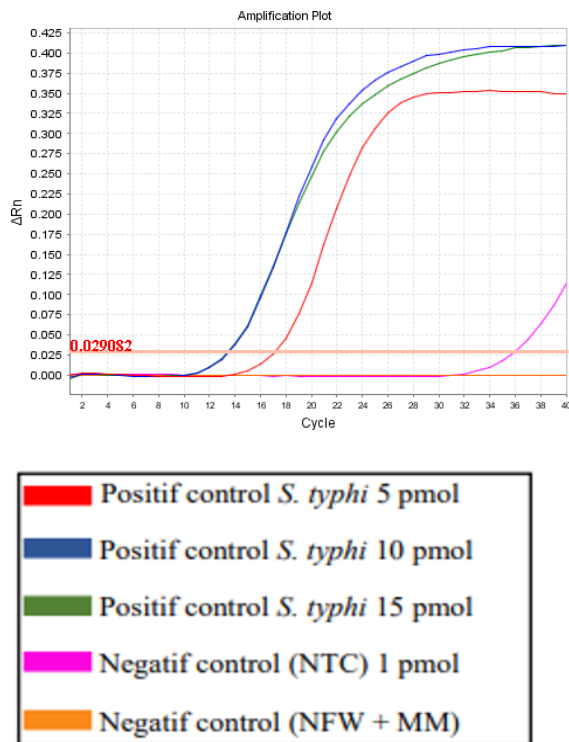
The confirmation test aims to determine whether or not the primer is capable of detecting and amplifying the target bacteria. Confirmation tests have been carried out in previous studies with a pre-denaturation time of 3 minutes and a primary concentration of 10 pmol. In this research, optimization of pre-denaturation time and primary concentration was carried out, optimizing the variations of pre-denaturation time used, namely 3 minutes, 4 minutes, and 5 minutes, while also optimizing the variations of primary concentrations used, namely 5 pmol, 10 pmol, and 15 pmol. The validation results of the confirmation test with variations in pre-denaturation time and variations in primary concentration are shown in Figures 5-7.



**Fig. 5.** Result of confirmation test Pradenaturation 3 minutes



**Fig. 7.** Result of Confirmation test Pradenaturation 5 minutes.



**Fig. 6.** Result of Confirmation test pradenaturation 4 minutes

Based on the results obtained by varying the primer concentration and predenaturation time, the amplification process in real-time PCR produces a stable curve and the optimum conditions for each. but to make a detection kit product, of course, it is necessary to streamline the time and materials to be used, so the results of 3 minutes of predenaturation with a concentration of 10 pmol will be used as a reference and compared with the results in previous studies. The Ct value obtained in the previous study was 14.35, with a negative control Ct value (NTC) of 32.63 [10]. The Ct value obtained from the results of 3 minutes of predenaturation validation and 10 pmol of concentration was 13.96, with a negative control Ct value (NTC) of 34.76. Shifts in CT values can indeed occur; ideally, the dynamic range for real-time PCR is 7-8 fold for plasmid DNA, and a range of 3-4 for genomic DNA [8]. However, the Ct value was also found in the negative control (NTC), while the negative control NFW and master mix did not amplify. but the curve can be ignored if the negative control CT is more than 10 positive control cycles [9]. From the ct value data obtained, it can be said that the validation using real-time PCR was declared successful and reproducible.

## 4. Conclusion

Based on the results obtained by validating the annealing temperature with gradient PCR, the same results were obtained in the previous study, namely, that there was an amplification band at 95 bp at 60 °C. A validation confirmation test showed that the primer was able to recognize the target DNA, resulting in amplification, which was indicated by the presence of Ct values and shifts in Ct values obtained in previous studies with validation results not too far away, so this validation can be said to be successful and reproducible.

## 5. Acknowledgements

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