

Development and Validation of SYBR Green-Based qPCR Technique of Detection and Quantification of *Salmonella enterica*

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Abstract

Salmonella enterica serovars is a leading cause of human gastroenteritis, and the incidence of salmonellosis is constantly increasing, causing millions of infections and many deaths annually. The detection of the pathogen in optimal terms is an essential factor for reducing the impact on the human body. In this work, SYBR Green I-based qPCR method of detection and quantification of *Salmonella enterica* subsp. *enterica*, two pairs of primers were designed using publically available Primer-BLAST software. Primer efficiency was calculated by establishing a standard curve. The specificity, sensitivity, accuracy, and precision of PCR results were tested. Both primer pairs showed an acceptable performance, proving the developed techniques were sensitive, reliable and precise. The validated qPCR technology has a good potential to replace the traditional culture method in microbial diagnosis.

Keywords

Dilution, DNA, Primer, Real-Time PCR, *Salmonella enterica*

1. Introduction

In recent decades, according to World Health Organization (WHO), the incidence of *Salmonella* infection (salmonellosis) has been increasing worldwide. Contamination of food with *Salmonella* spp. and, as a consequence, the development of salmonellosis is a serious threat that requires great attention to the control of the microbiological purity of food, the development of quick and rapid, accurate methods for its detection.

According to the Bergey's Manual of Systematic Bacteriology, *Salmonella* belongs to the Enterobacteriaceae family, genus—*Salmonella*, which includes 2 species: *Salmonella bongori* and *Salmonella enterica* [1]. *S. enterica* subdivided into six subspecies with about 2500 serovars (serotypes) [2]. These serovars can cause illnesses in human ranging from gastroenteritis to typhoid fever which kills around 500,000 people worldwide per year [3].

The source of salmonellosis, most often, are eggs (up to 90% of cases of salmonellosis are associated with the consumption of raw or insufficiently cooked eggs), meat and dairy products, and, to a lesser extent, fish and fish products, as well as plant products. *Salmonella* is relatively stable in the external environment and is well preserved in room dust (almost up to three months), in open water (11 to 120 days), in meat and sausages (2 - 6 months), in frozen meat (more than six months), in milk at room temperature for up to 10 days and in the refrigerator for up to 20 days; in butter (52 - 128 days); in eggs (over a year), on eggshells (24 days). At 70°C, salmonella die within 5 - 10 minutes, in the thickness of meat they withstand boiling for several hours. In meat stored in a refrigerator at a low above zero temperature, salmonella not only survive, but are also able to reproduce [4].

Recently, salmonella serotypes have become widespread, characterized by resistance to many modern common antibiotics and disinfectants, as well as increased heat resistance [5].

It has been extensively reported that classical microbiological methods, even though reliable, are lengthy and tedious. In this sense molecular applications, relies on the detection of specific *Salmonella* genes by PCR techniques, such as Real-Time PCR [6] [7], which have greatly allowed the reduction in the time needed to achieve the final result and to help control the spread of disease [8].

For designing a reliable PCR assay, it is necessary to pick a good target gene for PCR amplification. Several target genes of *Salmonella* have been reported, such as *invA* [6] [7] [9], *sdhA*, *fliC*, *fliB*, *sefB*, *hilA* [10] [11] [12] [13] [14].

The *invA* gene lies at centisome 63 region of the *Salmonella* chromosome and is widely used as a target in PCR assays for *Salmonella* detection [10].

The *invA* gene usually codes for a protein of the inner bacterial membrane, which is responsible for the invasion of the intestinal cells of the host [4] and is promising for primer design since it contains conservative and well characterized sequences. In order to validate the PCR assays of *Salmonella* molecular detection it is necessary to test specificity, sensitivity, accuracy, and precision of results [15].

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