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Detection Flagellin Gene of Enteric Isolates of *Salmonella enterica* serovar *Typhi* Using Conventional PCR Technique

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Abstract

The study was conducted to study the virulence gene (Flagellin gene) of *Salmonella typhi*. Stool samples were collected from patients with are ranging (15 – 60) years of the both sexes who suffering from diarrhea and visited AL-Yrmouk Teaching Hospital in Baghdad Governorate for the period between March to September 2017..The samples were cultured on suitable media then Api20 test was done in order to confirm the diagnosis, the polymerase chain reaction (PCR) was tested to diagnose the Flagellin gene responsible for the virulence of the bacteria. We conclude from the present study that flagellin gene in *S.typhi* is responsible for pathogenicity in this bacteria.

Keywords: Flagellin, gene, *Salmonella*, *Typhi*, Typhoid fever.

الخلاصة

أجريت هذه الدراسة لمعرفة جينات الضراوة لجراثيم سالمونيلا تايفي حيث تم جمع عينات الخروج من المرضى البالغين الذين تتراوح اعمارهم من (١٥ – ٦٠) سنة ومن كلا الجنسين الذين يعانون من الإسهال والمراجعين لمستشفى اليرموك التعليمي في محافظة بغداد للفترة من شهر اذار الى شهر ايلول ٢٠١٧. حيث تم زرع العينات على الأوساط الزرع المناسبة وبعدها تم عمل فحص Api20 وذلك لتأكيد التشخيص بعد ذلك تم اجراء فحص تفاعل البلمرة المتسلسل PCR لتشخيص جين الفلاجلين المسؤول عن ضراوة البكتيريا. نستنتج من هذه الدراسة وجود جين الضراوة الفلاجلين في جراثيم السالمونيلا تايفي والذي يعتبر مسؤول عن الامراضية في هذه البكتيريا.

Introduction

Typhoid fever consider the main problem of public health and spread in developing countries, the main causes belong to *Salmonella typhi* bacteria [1][2]. The cases of typhoid fever are large amount in the world and very difficult to detect because the clinical Signs are widely spread [3]. The World Health Organization Proven cases of typhoid fever were 17 million [4]. The pathogenicity of *S. typhi* depends on virulence factor which causes the disease in human, The villi or fimbria which present on the cell surface of *S. typhi* which playing an important roles in the process of adhesion and colonization to the human cells [5]. Flagella in *S. typhi* is responsible for motility and play critical role in host cell invasion and escape from immunity by intra cellular site.

Keeping the concept of the introduction the same, different documents have different styles

to introduce the written text. For example, the introduction of a Functional Specification consists of information that the whole document is yet to explain. If a User guide is written, the introduction is about the product. In a report, the introduction gives a summary about the report contents.

Materials and Methodologies

Samples collection

167 stool samples were obtained from diarrhea cases from patients who attended to AL-Yarmouk Teaching Hospital in Baghdad Governorate during the study period from March to September 2017. The samples were transferred in cold conditions to the laboratory for diagnostic tests.

Identification of *S.typhi*

Morphological and cultural characteristics of bacterial isolates

Microscopical characteristics of bacterial cells as well as cultural properties of the colonies grown on specific culture media were determined using standard methods as primary diagnosis of *S. typhi* [8].

Biochemical Tests and API 20 E System

The bacterial isolates were identified using different biochemical tests such, oxidase, catalase test, in addition to growth on Kligler Iron agar [9]. Bacterial identification of *S.typhi*

was conformed using API 20E strips (Biomerieux, France).

Extraction of the total DNA

The extraction of DNA was done by extraction kit.

Amplification of the total DNA

Specific primers and the amplification conditions were presented in Table 1.

Table 1: Primers used in this study.

Primers		Sequence	Amplicon size
<i>fliC18</i>	F	ACT GCT AAA ACC ACT ACT	363 bp
		TGG AGA CTT CGG TCG CGT	
	R	AG	

Table 2: Amplification conditions used in this study.

PCR cycle	repeat	Temperature	Time
Initial denaturation	1	94°C	5min
Denaturation	30	94°C	20sec.
Annealing		50-65°C	30sec
Extension		72°C	30sec
Final extension	1	72°C	7min

Results and Discussion

Distribution of *S. typhi* Positive Culture

A total of 45 (26.9%) *S. typhi* isolates were obtained from 167 patients with clinically suspected typhoid fever. The results were confirmed with positive cultures.

Distribution of Flagellin Gene

Detection of flagellin gene performed by PCR using primers according to [11], as shown in Figure 1.

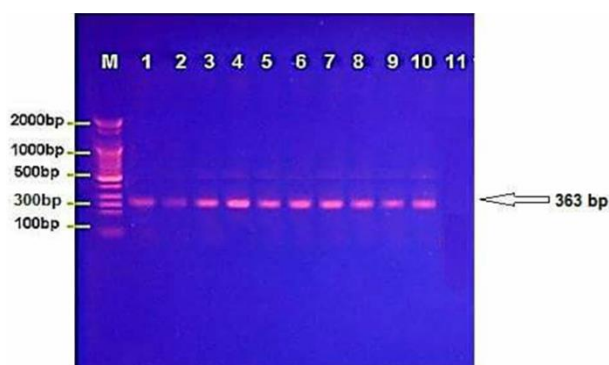


Figure 1: Agarose gel electrophoresis (1% agarose, 7v/cm² for 30 min) of flagellin gene of *S.typhi*. (363bp) amplicon size compared with M line (M100 bp DNA ladder).

Typhoid fever caused by *S.typhi* is a worldwide health problem in different countries [12]. This problem is particularly prominent in developing countries because of several correlated factors like unplanned urbanization with growing of peri-urban towns that lack safe water supply, lacking of sanitation services, increased local migration of great sums of workers, in addition to inefficiencies of vaccine preparations programs [13].

The results showed isolation and identification of *S.typhi* from diarrhea samples these results in agreement with [14] which recorded the morphological properties of the *S.typhi* exhibited Gram negative, small Virulence of *S. typhi* possessed an important factor the infection of human by typhoid fever, these factors are fimbria or pill present on the cell surface of *S.typhi* which possess effective role in the infection human [18]. The role of motility of *S.typhi* in pathogenicity was Penetration of the intestinal mucosa which consider the important step in the causes of infection because it allows bacteria to pass through the epithelial barrier, the Flagella are consist of a protein called flagellin which also act as an antigen [19].

The results detected the presence of flagellin gene which responsible for the pathogenicity of *S.typhi*, who reported that flagellin genes might be responsible for an amplified risk for emerging severe disease. The study of [7] found that flagellin gene present in three types which responsible for the gastrointestinal bleeding and make the *S.typhi* more invasive compared to others microorganism.

References

- [1] Crump JA, Mintz ED: Global trends in typhoid and paratyphoid Fever. *Clinical infectious diseases: an official publication of the Infectious Diseases Society of America* 2010,50(2):241-246.
- [2] Sankar S, Kuppanan S, Nandagopal B, Sridharan G: Diversity of Salmonella entericaserovarTyphi strains collected from india using variable number tandem repeat (VNTR)-PCR analysis. *Molecular diagnosis & therapy* 2013, 17(4):257-264.
- [3] Ley B, Mtove G, Thriemer K, Amos B, von Seidlein L, Hendriksen I, Mwambuli A, Shoo A, Malahiyo R, Ame SM *et al*: Evaluation of the Widal tube agglutination test for the diagnosis of typhoid fever among children admitted to a rural hdospital in Tanzania and a comparison with previous studies. *BMC infectious diseases* 2010, 10:180.
- [4] (WHO) WHO: Background document: The diagnosis, treatment and prevention of typhoid fever. In. Geneva: World Health Organization.; 2003.
- [5] Burrows LL: Weapons of mass retraction. *Molecular microbiology* 2005, 57(4):878-888.
- [6] Ibarra JA , Steele-Mortimer O , 2009 . *Salmonella* –the ultimate insider. *Salmonella* virulence factors that modulate intracellular survival . *Cell Microbiol* 11: 1579 – 1586.
- [7] Nataniel T., Yadi Y., M. Sabir, Masyhudi A., Moch. H. 2015.Distribution Flagellin Gene Variants of Salmonella Typhi in Patients with Typhoid Fever in West Kutai, East Kalimantan,Indonesia.American Journal of Biomedical and Life Sciences 3(5): 98-103.
- [8] Jawetz E., Melnick J.I. and Adelberg E.A. (2007) *Medical Microbiology*, 24th ed., Appleton and Lange, USA.
- [9] MaccFadin,J.K. (2000). Biochemical test for identification of medical bacteria. (3rd ed.). Lippincott Williams and Winkins . AwolterKlumer Company . Philadelphia Baltimor, New York.
- [10] Parry, C. M., Wijedoru, L., Arjyal, A. and Baker, S. (2011). The Utility of Diagnostic Tests for Enteric Fever in Endemic Locations. *Expert Review ofAnti Infection Therapy*, 11: 711 725.
- [11] Taylor M, Coovadia HM, Kvalsvig JD, Jinnabhai CC, Reddy P. Helminth control as an entry point for health –promoting schools in KwaZulu Natal. *S Afr Med J* 1999; 89:273 - 79.
- [12] Levine MM. Typhoid fever. In: Evans AS, Brachman PS, (eds). *Bacterial Infections of Humans*. Epidemiology and Control. Third edn. New York, Plenum Medical Book Company, 1998,pp.839-58.
- [13] Thong KL, Cheong YM, Puthucheary S, Koh CK, Pang T. Epidemiologic analysis of sporadic *Salmonella typhi*isolates and those from outbreaks by pulsed –field gel electrophoresis. *J ClinMicrobiol* 1994; 32:1135-41.
- [14] Sogard M, Norgaard M and Schonheyder HC (2007). First notification of positive blood cultures and the high accuracy of the gram stain report. *Journal of Clinical Microbiology* 45(4):1113-1117.
- [15] Buxton A and Fraser G (1977). *Animal Microbiology*. Blackwell Scientific Publications, Oxford. Vol.1:85- 110.
- [16] Han J, David JD, Aaron ML, Pravin K, Rajesh N, Rossina S and Steven LF (2011). Comparison of Salmonella entericaserovar Heidelberg isolates from human patients with those from animal and food sources. *Journal of Clinical Microbiology* 49:1130-1133.
- [17] Gallati C, Stephan R, Hächler H, Malorny B, Schroeter A, Nüesch-Inderbinen M, 2013. Characterization of Salmonella enterica Subsp. entericaSero var 4,[5],12:i:- Clones Isolated from Human and Other Sources in Switzerland Between 2007 and 2011. *Foodborne Pathog Dis*. 2013 Jun; 10(6):549-554.
- [18] Burrows L.L., 2005 : Weapons of mass retraction. *Molecular microbiology*, 57(4):878-888.
- [19] Josenhans C, Niehus E, Amersbach S, Horster A, Betz C, Drescher B, Hughes KT, Suerbaum S. 2002: Functional characterization of the antagonistic flagellar late regulators FliA and FlgM of Helicobacter pylori and their effects on the H. pylori transcriptome. *Molecular microbiology*, 43(2):307-322.