
Isolation of *Enterobacter sakazakii* from Some Spices

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ABSTRACT

Enterobacter sakazakii is considered an opportunistic pathogen that has been associated with severe lethal infections especially in neonates, elderly, and immunocompromised adults. *E. sakazakii* is a Gram negative, facultative anaerobes rod-shaped bacterium. It belongs to the family Enterobacteriaceae and genus *Enterobacter*.

Although we don't know the natural habitat of this bacteria we find that it exists in high rate in herbs and spices which indicates that plant may be this natural habitat.

Our study focused on spices, and contained 59 specimen (mixed, unmixed), which cultivated on general and selective medium, then we detected its presence based on α -glucosidase activity by using ESIA medium that contains 5-bromo-4-chloro-3-indolyl- α -D-glucopyranoside (X α Glc) as a substrate which the bacteria hydrolyses it then producing blue-green colony, then we confirmed our results by cultivating these colonies on Tryptic Soy Agar (TSA), after this we chose the typical yellow one to identify it by using biochemical tests and Polymerase chain reaction PCR. Our results showed that 50% of the examined specimen contained *E. sakazakii*.

Key words: *Enterobacter sakazakii*, Spices, Biochemical tests, Polymerase chain reaction (PCR).

Enterobacter sakazakii

(*Cronobacter* spp.

) *Enterobacter sakazakii*

1

3

Enterobacteriaceae

peritrichous

Voges-)

(Proskauer test

. [3, 2, 1]

°47-5.5

44-37

°22

°43-37

. [4]

20

°37

2.5≥)

. [5]

(

28

)

(

Meningitis

E. sakazakii

Bacteremia

Necrotizing enterocolitis

. [7, 6] %80-10

%94
.[10, 9, 8]

Farmer *et al.* 1980
DNA-DNA

[1] Riichi sakazakii

Iversen *et al.*
Cronobacter gen nov. 2008

.[11]

E. sakazakii
Enterobacteriaceae *Cronobacter*
C. dublinensis *C. muytjensii* *C. malonaticus* *C. turicensis* *C. sakazakii*

.[12]

:

:

(BPW) Buffered Pepton Water
modified Luryl Sulphate Treptose
(mLST/Van) broth/Vancomycin
(ESIA) *Enterobacter Sakazakii* Isolation Agar
(TSA) Tryptone Soy Agar

:

SDS 10% (10mM Tris-HCl pH 8.0, 1mM EDTA pH 8.0) TE
 CTAB NaCl NaCl 5M 20mg/ml K
 .%70 (25:24:1) (24:1)

:

200mM Tris-HCl pH8.8, 100mM KCl, 100mM) 10X PCR
 (dCTP, MgSO₄ ((NH₄)₂SO₄, 1mg/ml BSA, 1% Triton
 (1) DNA dATP, dTTP, dGTP)
 0.4M) 50X TAE 1X TAE (W/V) %1.5
 .(Tris-Base, 10mM EDTA pH 8.0, 57.1 ml Glacial Acetic Acid

:

(BD,USA) BBL™ DrySlide™ %3
 .MacConkey agar MR-VP KOH

:

.%20

:

:

:

:

E. sakazakii

4.5

0.5

18 °37

(BPW)

5

BPW

50 µl

24 °44

(mLST/Vancomycin)

/

°44 (ESIA)

10 µl

5-bromo-4-chloro-3-

24

E. sakazakii

indolyl- α ,D-glucopyranoside

[13] *E. sakazakii*

(TSA)

72-48 °25

1.5-1

E. sakazakii

(ESIA)

(BD,USA) BBL™DrySlide™

%3

ESIA

5-bromo-4-chloro-3-indolyl- α ,D-glucopyranoside

MacConkey

6.8

pH

neutral red

3

24 °37

pH)

Agar

MR-VP

(4

	(7	pH)	(7	pH)
0.5 (KOH) A		0.5	:	-
)			(α -naftol) B	
			(
			PCR	
		:		TSA
4000 rpm		5		.1
			+4°C	
30 μ l		TE	567 μ l	.2
.20 mg/ml proteinase K		3 μ l 10% SDS		
.1000 rpm		°37		
80 μ l		5M NaCl	100 μ l	.3
°65		10	.CTAB/NaCl	
			.1000 rpm	
(24:1) chloroform/isoamyl alcohol			780 μ l	.4
		14500 rpm	5	
phenol/chloroform/isoamyl alcohol				.5
14500 rpm		5	(25:24:1)	
	0.6			.6
	DNA		6-4	
	14500 rpm		5	
		.DNA		
14500	5	%70	1	.7
) Concentrator				
			rpm	
			(eppendorf	

TE 25 µl .8
 .100 ng/µl (Nano Drop)

DNA PCR
 TSA

.(1)

(1)

Primer	Sequence (5 to 3)		Size (bp)
SI-F	5 -CAG-GAG-TTG-AAG-AGG-TTT-AAC-T-3	22	251 bp
SI-R	5-GTG-CTG-CGA-GTT-TGA-GAG-ACT-C-3	22	
SG-F	5-GGG-TTG-TCT-GCG-AAA-GCG-AA-3	20	282 bp
SG-R	5-GTC-TTC-GTG-CTG-CGA-GTT-TG-3	20	
EsAg-F	5-TGA-AAG-CAA-TCG-ACA-AGA-AG-3	20	1680 bp
EsAg-R	5-ACT-CAT-TAC-CCC-TCC-TGA-TG-3	20	

(2) 25 µl

(3)

.PCR (2)

Materials	Final conc. µl	25 µl PCR
Genomic DNA	200-500 ng	2 µl DNA (100ng)
Primer 1 -10 µM	20 µM	2 µl
Primer 2-10 µM	20 µM	
dNTPs 20 mM	0.4 mM	0.5 µl
Buffer 10X	1X	2.5 µl
MgSO ₄ 50 mM	3 mM	1.5 µl
Taq 5U	2 U	0.2 µl
H ₂ O	----	16.3 µl

.PCR (3)

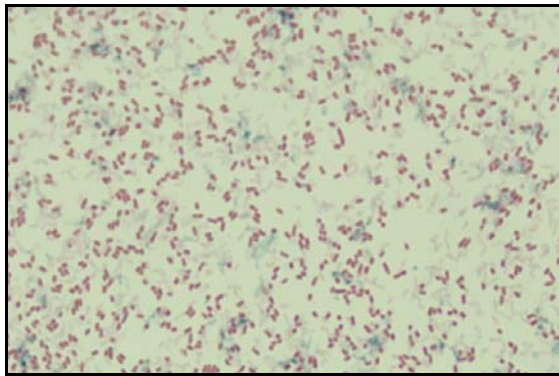
		Temperature	Time
	Initial denaturation	C°95	5 mins
35 Cycle	Denaturation	C°95	1 min
	Annealing	C°57	1 min
	Extension	C°72	1.5 min
	Final Extension	C°72	10 mins

(W/V) %1.5
 70V
 %20
 PCR
 1X TAE
 BPW
 °80-
 ESIA
E. sakazakii

)
 .PCR (

TSA
 (1)

.100X

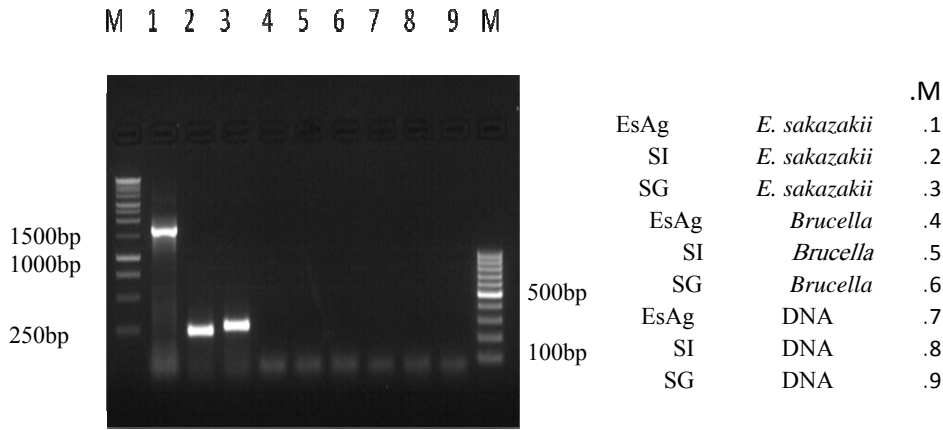


100X *E. sakazakii* (1)

E. sakazakii

. [3,2,1]

(4)
 .PCR



(W/V) %1.5

(2)

282 bp 251 bp 1680 bp
EsAg SG SI EsAg
SG SI [14] *gluA*
.[15] rDNA 23S 16S ITS

E. sakazakii

(5)

E. sakazakii

(5)

	<i>E. sakazakii</i>		
55%	18	33	
42%	11	26	
49%	29	59	

E. sakazakii

[16]

E. sakazakii

3 PCR

[19,18,17] DNA

(%49) 59 29

[16] Friedemann [20] Forsythe

[22]

[21]

E. sakazakii

REFERENCES

1. Farmer J. J., III, Asbury M. A., Hickman F. W., Brenner D. J. (1980). The Enterobacteriaceae Study Group (USA): *Enterobacter sakazakii*: a new species of "Enterobacteriaceae" isolated from clinical specimens. *Int. J. Syst. Bacteriol.* 30:569-584.
2. Iversen C., Waddington M., On S. L., Forsythe S. (2004). Identification and phylogeny of *Enterobacter sakazakii* relative to *Enterobacter* and *Citrobacter* species. *J. Clin. Microbiol.* 42:5368-5370.
3. Iversen C., Waddington M., Farmer J. J. III, Forsythe S. (2006). The biochemical differentiation of *Enterobacter sakazakii* genotypes. *BMC Microbiol.* 6:94.
4. Dauga C., Breeuwer P. (2008a). Taxonomy and physiology of *Enterobacter sakazakii*. Ed. By Farber J. M. and Forsyth S. J. ASM press. Washington, D. C. pp 15-16.
5. M. Leclerc. (2006). *Enterobacter sakazakii*. Agence Française de Sécurité Sanitaire des Aliments.
6. Corti G., Panunzi I., Losco M., Buzzi R. (2007). Post-surgical osteomyelitis caused by *Enterobacter sakazakii* in a healthy young man. *J. Chemotherapy*, 19:94-94.
7. Muytjens H. L., Zanen H. C., Sonderkamp H. J., Kollee L. A., Wachsmuth I. K., Farmer J. J. (1983). Analysis of eight cases of neonatal meningitis and sepsis due to *Enterobacter sakazakii*. *J. Clin. Microbiol.*, 18:115-120.
8. Gallagher P. G. (1990). *Enterobacter* bacteremia in pediatric patients. *Rev. Infect. Dis.*, 12:808-812.
9. Lehner A., Stephan R. (2004). Microbiological, epidemiological, and food safety aspects of *Enterobacter sakazakii*. *J. Food Prot.* 67:2850-2857
10. Gurtler J. B., Kornacki J. L., Beuchat L. (2005). *Enterobacter sakazakii*: A coliform of increased concern to infant health. *Int. J. Food Microbiol.*, 104:1-34.
11. Graves R., (Ed). (1992). The Dethronement of Cronos. In *The Greek Myths*. Combined Edition. London: Penguin Books;:39-44.
12. Iversen C., Mullane N., McCardell B., Tall B. D., Lehner A., Fanning S., Stephan R., Joosten H. (2008). *Cronobacter* gen. nov., a new genus to accommodate the biogroups of *Enterobacter sakazakii*, and proposal of *Cronobacter sakazakii* gen. nov., comb. nov., *C. malonaticus* sp. nov., *C. turicensis*, sp. nov., *C. muytjensii* sp. nov., *C. dublinensis* sp. nov., *Cronobacter genomospecies* 1, and of three subspecies. *C. dublinensis* sp. nov. subsp. *dublinensis* subsp. nov. *C. dublinensis* sp. nov. subsp. *lausannensis* subsp. nov., and *C. dublinensis* sp. nov. subsp. *lactaridi* subsp. nov. *Int. J. Sys. Evol. Microbiol.* 58:1442-1447.
13. ISO/TS 22964 (2006). Milk and milk products – Detection of *Enterobacter sakazakii*. International Organization for Standardization.

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14. Lehner A., Riedel K., Rattei T., Ruepp A., Frishman D., Breeuwer P., Diep B., Eberl L., Stephan R. (2006). Molecular characterization of the α -glucosidase activity in *Enterobacter sakazakii* reveals the presence of a putative gene cluster for palatinose metabolism. *Syst. Appl. Microbiol.*; 29:609–625.
 15. Liu Y., Gao Q., Zhang X., Hou Y., Yang J., Huang X. (2006). PCR and oligonucleotide array for detection of *Enterobacter sakazakii* in infant formula. *Mol. Cell Prob.* 20:11–17.
 16. Friedemann M. (2007). *Enterobacter sakazakii* in food and beverages (other than infant formula and milk powder). *Int. J. Food Microbiol.* 116:1-10.
 17. Iversen C., Lehner A., Mullane N., Marugg J., Fanning S., Stephan R., Joosten H. (2007). Identification of "*Cronobacter*" spp. (*Enterobacter sakazakii*). *J. Clin. Microbiol.*, 45:3814-3816.
 18. Drudy D., Rourke M.O., Murphy M., Mullane N.R., O'Maony R., Kelly L., Fisher M., Sanjaq S., Shannon P., Wall P., O'Mahony M., Whyte P., Fanning S. (2006). Characterization of a collection of *Enterobacter sakazakii* isolates from environmental and food sources. *Int. J. Food Microbiol.*, 110:127-134.
 19. Fanjat N., Leclercq A., Joosten H., Robichon D. (2007). Comparison of the Phenotyping Methods ID32 and VITEK 2 Compact GN with 16S rDNA Gene Sequencing for the Identification of *Enterobacter sakazakii*. *J. Clin. Microbiol.*, 45:2048-2050.
 20. Forsythe S. J. (2005). *Enterobacter sakazakii* and other bacteria in powdered infant milk formula. *J. Matern Child Nutr.*, 1:44-50.
 21. Nazarowec-White M., Farber J. M. (1997). Thermal resistance of *Enterobacter sakazakii* in reconstituted dried-infant formula. *Lett. Appl. Microbiol.*, 95:967-973.
 22. Breeuwer P., Lardeau A., Peterz M., Joosten H. M. (2003). Desiccation and heat tolerance of *Enterobacter sakazakii*. *J. Appl. Microbiol.*, 95:967-973.