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Summary of microbiological findings in healthy human

Materials and methods

- Fecal specimens were collected from healthy asymptomatic individuals, including farmers in Hanoi and Thai Binh, and marker sellers in Nha Trang.
- ✓ Sampling at Hanoi, Thai Binh and Nha Tran since 2013.
- ✓ Total collected sample numbers: more than 1500.
- CTX-M type ESBL-producing bacteria have been screened and characterized by following the protocol which had been established in this project.
- Collected fecal specimens were subjected to microbiological and molecular biological analysis, such as phylogenetic grouping, genotyping of antibiotic-resistant genes, antibiotic susceptibility test and PFGE.

Prevalence of the CTX-M type ESBL-producing *E. coli* in healthy asymptomatic individuals in Vietnam.

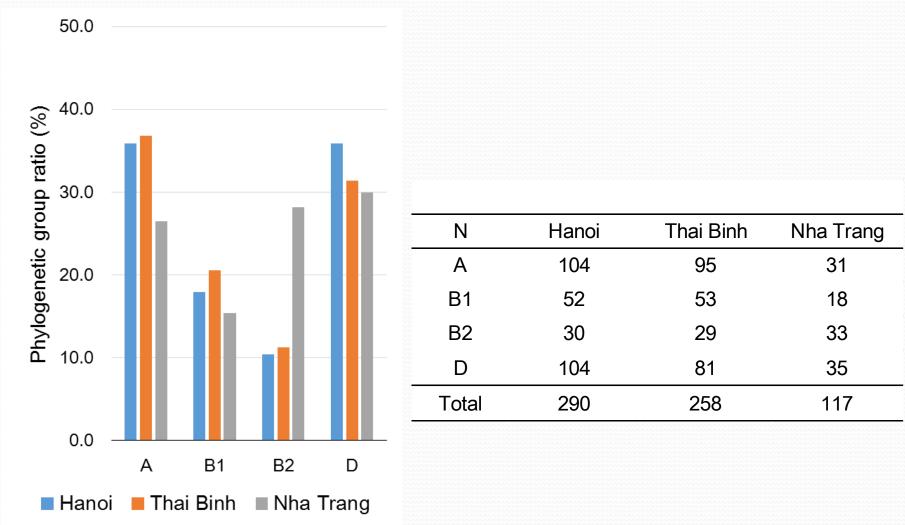
Sites	ESBL-producing bacteria in healthy people				
	Ν	%			
Hanoi ¹	1025	59.0%			
Thai Binh ¹	397	65.0%			
Nha Trang ¹	156	75.0%			
Total	1578	62.1%			
2 Thailand	417	65.9%			
Japan ³	218	6.0%			

1: Samples were collected from market seller in Nha Trang, from farmer in Hanoi and Thai Binh.

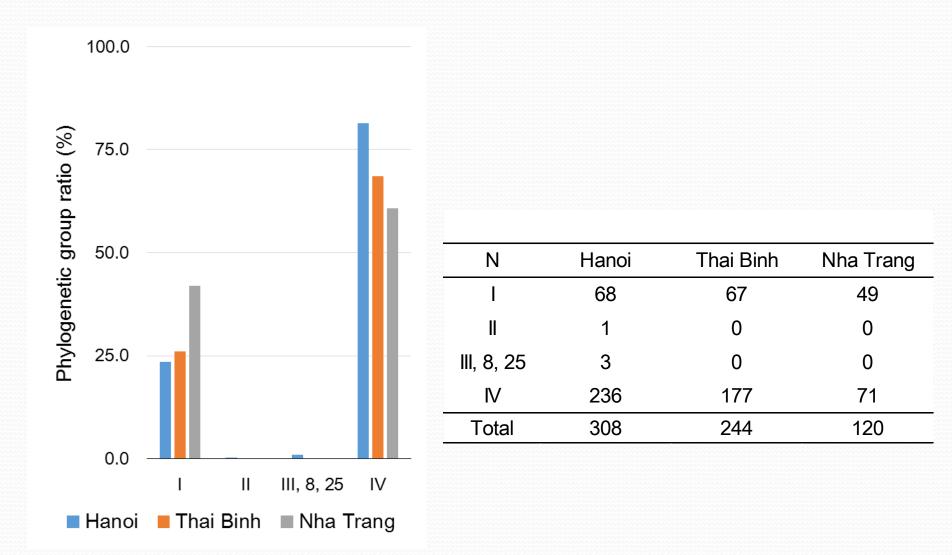
2: Luvsansharav UO, et al., J Antimicrob Chemother. 2012;67(7):1769–1774.

3: Luvsansharav UO, et al., J Infect Chemother. 2011;17(5):722-725.

Phylogenetic group classification of the isolated CTX-M type ESBL-producing *E. coli*



Genotyping of the detected *bla*_{CTX-M}s in healthy human.



Clinical isolates and isolates from UTI patients

- ✓ 86 clinical isolates were collected in Nha Trang, and 36 isolates (41.9%) were ESBL-producing *E. coli*. Among them, 35 isolates were CTX-M type ESBL-producing *E. coli*.
- ✓ Until now, 14 CTX-M-I group genes (11 bla_{CTX-M-15} and 3 bla_{CTX-M-15}) and 12 CTX-M-IV group genes (4 bla_{CTX-M-14} and 8 bla_{CTX-M-27}) were confirmed.
- ✓ 12 ESBL-producing *E. coli* isolates were obtained from 50 UTI patients (24.0%) in Thai Binh.
- ✓ 6 CTX-M-I group genes (5 bla_{CTX-M-15} and 1 bla_{CTX-M-55}) and 6 CTX-M-IV group genes (1 bla_{CTX-M-14} and 5 bla_{CTX-M-27}) were detected.

PFGE analysis (Phylogenetic group B2)

	:13TE074	13Aug31	Fish	Thong	62	TEM	ESEL	cbi-m-9s	FIB,Frep
	:15TBU027	DK767(1)	ຫ	т	E 2	TEM	ESBL	CTX-M-27	FIB
⊣∟	.14TB180	14Jun22	Strimp	Lua	62	none	ESBL.	ctv-m-9s	FIB,Frep
	31.093_Dec13	31.093	human	т	B 2	TEM	ESBL	CTX-M27	FIB,Frep
	.19.080_Dec13	19.060	human	т	E 2	TEM	ESBL	CTX-M27	FIB,FIA, Frep
	22.067_Dec13	22.067	human	т	62	TEM	ESBL.	CTX-M-27	FIB,FIA, Frep
	:15TEU008	DK597(1)	ຫ]	т	B2	TEM	ESBL	CTX-M-15	FIA, FIB, 11
	44.132_Dec13	44.132	human	т	E 2	т	ESBL	CTX-M-14	FIB/Frep
	:13TE019	13Jun15	Chicken	Thong	B2	none	ESBL	ctx-m-9s	FIB,HI2
	:15TEU020	DK999	UII)	т	62	TEM	ESBL.	CTX-M-15	FIB
l r	15TBU023	DK851(1)	UTI	т	B2	TEM	ESBL	CTX-M-27,CMY-8b	A/C, FIA, FIB, Frep
	15TBU025	DK888(1)	UTI	т	62	TEM	ESEL, A	CTX-M-27,CMY-8b	A/C, FIA, FIB, Frep
	15TBU032	DK702(1)	υπJ	т	B2	0	ESBL	CTX-M-27	A/C, FIB
	:13TE056	13Aug18	Strimp	Cau	E 2	TEM	ESBL	ctx-m-9s	FIB/I1,HI1,N
	70.212_Dec13	70.212	human	т	62	TEM	ESEL	CTX-M-14	FIB;FIA;like;Frep
	:16.046_Dec13	16.046	human	т	E 2	TEM	ESBL	CTX-M-14	FIB,FIA-like,Frep
	.16.048_Dec13	16.048	human	т	62	TEM	ESEL	CTX-M-14	FIB;FIA;like;Frep
	:14TE010	14Mar08	Chicken	Thong	E 2	TEM	ESEL	ctx-m-9s	FIBN,HI2
	:13TE007	13Jun06	Strimp	Thong	62	none	ESBL	cbi-m-9s	FIB,FIA, Frep
	:13TE011	13Jun07	Strimp	Thong	E 2	none	ESEL	none	FIB,FIA, Frep
	:13TE003	13Jun02	Park	Thong	62	none	ESBL	ctx-m-9s	FIB,FIA, Frep
	32.098_Dec13	32.098	human	т	62	NONE	ESEL	CTX-M27	FIBFIA
	20.064_Dec13	20.064	human	т	62	NONE	ESBL	GTX-M-27	FIBFIA
	06.020_Dec13	06.020	human	%	62	NONE	ESEL	CTX-M-27	FIB,FIA

Summary

- Prevalence of the CTX-M type ESBL-producing E. coli was high (62.1%) in Vietnamese healthy individuals.
- ✓ The CTX-M-type ESBL-producing E. coli was detected in clinical isolates (41.9%) and UTI patients urine samples (24.0%) with higher detection rate.
- ✓ PFGE analysis indicated similar ESBL-producing E. coli strain was detected in food item (fish) and UTI patient.

THANK YOU FOR YOUR KIND ATTENTION