

Contents lists available at ScienceDirect

Food Control

journal homepage: www.elsevier.com/locate/foodcont



Molecular epidemiology and antibiotic resistance phenotypes and genotypes of salmonellae from food supply chains in China



Guoxiang Chao ^{a, b, c, e}, Chao Wang ^c, Tianqi Wu ^a, Xiaorong Zhang ^{a, b}, Jianhao Chen ^a, Xiaoxing Qi ^d, Yongzhong Cao ^{a, b}, Yantao Wu ^{a, b, **}, Xinan Jiao ^{a, b, *}

- a Jiangsu Co-Innovation Center for Prevention and Control of Important Animal Infectious Diseases and Zoonoses, Yangzhou University, Yangzhou, China
- ^b Joint International Research Laboratory of Agriculture and Agri-Product Safety, The Ministry of Education of China, China
- ^c College of Medicine, Yangzhou University, China
- ^d China Institute of Veterinary Drug Control, Beijing, China
- ^e Yangzhou Center for Disease Control and Prevention, Yangzhou, China

ARTICLE INFO

Article history: Received 30 October 2016 Received in revised form 21 January 2017 Accepted 26 January 2017 Available online 30 January 2017

Keywords: Salmonella Multilocus sequence typing (MLST) Multidrug-resistant (MDR) clone β-lactamase genes

ABSTRACT

Salmonella is an important zoonotic agent and a vehicle for antibiotic resistance genes. Here, 294 isolates from humans and food-producing animals were subjected to serotyping, multilocus sequence typing, and assessment of phenotypic (15 antibiotics) and genotypic (32 resistance genes) antimicrobial resistance. Twenty-two serotypes and 35 sequence types (STs) were identified, the most common STs being ST11, including S. Enteritidis from chickens and humans; ST17, including S. Indiana from chickens; and ST40, including S. Derby from pigs and humans. Antimicrobial resistance phenotypes and genotypes exhibited ST- and serovar-specific features. ST11, clonal complex (CC) 19, ST40, and ST155 were moderately multidrug-resistant (MDR) clones, most of the isolates of which were resistant to between 3 and 6 antibiotics. Isolates of a super-MDR clone, ST17, demonstrated resistance against 9 to 14 antimicrobials, in particular, ampicillin, amoxicillin-clavulanic acid, cefotaxime, cefepime, cefoperazone, ceftriaxone, and ciprofloxacin. Consistent with this, ST17 (S. Indiana) was associated with a gene cluster comprising bla_{CTX-M} (and/or bla_{OXA-1}-like together with bla_{TEM-1}-like), sul1, aacC4, aac(6)-1b, floR, and dfrA17, while the moderately-MDR clones (ST11, S. Enteritidis) were more closely linked to the bla_{TEM-1}like gene. The similar genetic clones isolated from animals and humans indicate a common ancestor, and implicate animals as a major salmonellae source. Antibiotic abuse in animal production appears to be the origin of MDR and super-MDR isolates, the latter being closely associated with the β -lactamase genes bla_{CTX-M}, bla_{OXA-1}-like, and bla_{TEM-1}-like. Carried by chickens, ST17 (S. Indiana) is an emerging super-MDR clone whose associated resistance genes are expanding to other ST clones and serotypes being transmitted to animals and humans.

© 2017 Elsevier Ltd. All rights reserved.

1. Introduction

Salmonella is recognized as the causative agent of salmonellosis in both humans and animals. Many serovars of this bacterium, such as *S.* Enteritidis and *S.* Typhimurium, are common in food-producing animals, and a clear link between human illness and poultry consumption has been established (Foley & Lynne, 2008;

E-mail addresses: ytwu@yzu.edu.cn (Y. Wu), jiao@yzu.edu.cn (X. Jiao).

Guard-Bouldin, Morales, Frye, Gast, & Musgrove, 2007; Mead et al., 2010). Animals used in the production of food also play an important role in the transmission of antimicrobial-resistant *Salmonella* strains to humans (Beutlich et al., 2011). Several *S. enterica* serovars, including *S.* Typhimurium, *S.* Newport, *S.* Derby, and *S.* Agona, exhibit a super-multidrug-resistant (MDR) phenotype characterized by resistance to nine antimicrobial compounds, namely, ampicillin (AMP), amoxicillin/clavulanic acid (AMC), cefoxitin (FOX), cephalothin, ceftiofur, chloramphenicol (C), streptomycin, sulfamethoxazole, and tetracycline (TE). *Salmonella* genomic island 1 (SGI1) and its variants are associated with super-MDR isolates, in that this region contains a gene cluster conferring penta-resistance to AMP (*blaPSE*), C/florfenicol (*floR*), streptomycin/

^{*} Corresponding author. Yangzhou University, 88 South University Ave., Yangzhou, 225009, China.

 $^{\ ^{**}}$ Corresponding author. Yangzhou University, 88 South University Ave., Yangzhou, 225009, China.

Table 1 Salmonella isolates, serotypes, STs, and super-MDR isolates from different sources.

		Serotype (no.)	ST (no.)	No. of super-MDR isolates (ST
Chicken industry chains	(88) ^a			
Broiler farms (27)	Environmental soils (8)	S. enteritidis (2)	11 (2)	
		S. aberdeen (5)	426 (5)	
		S. agona (1)	40 (1)	
	Feeds (1)	S. montevideo (1)	17 (1)	
	Broiler (18)	S. enteritidis (3)	11 (2), 17 (1)	2 (ST11), 1(ST17)
		S. indiana (13)	17 (13)	13 (ST17)
		S. agona (1)	11 (1)	
		B group (1)	17 (1)	1 (ST17)
Slaughterhouses (49)	Plucking (20)	S. enteritidis (6)	11 (6)	1 (ST11)
		S. indiana (12)	11 (2), 17 (10)	9 (ST17)
		S. choleraesuis (1)	32 (1)	
		C group (1)	1541 (1)	
	Chilling (7)	S. enteritidis (5)	11 (5)	
		Non A-F group (1)	11 (1)	
		S. indiana (1)	17 (1)	1 (ST17)
	Tools (1)	S. kaapstad (1)	17 (1)	1 (ST17)
	Whole frozen carcasses (9)	S. enteritidis (1)	11 (1)	,
	,	S. indiana (5)	17 (5)	4 (ST17)
		C group (3)	1541 (3)	- ()
	Frozen split chicken (12)	S. enteritidis (7)	11 (7)	
		S. indiana (4)	17 (4)	2 (ST17)
		S. london (1)	463 (1)	·/
Distribution (2)	Frozen chicken on distribution (2)	S. enteritidis (1)	11 (1)	
	emercin on distribution (2)	S. lomita (1)	32 (1)	
Retail (10)	Frozen chicken at retail (10)	S. enteritidis (6)	11 (5), 17 (1)	
ictuii (10)	1102cm emeken at retair (10)	S. indiana (1)	17 (1)	1 (ST17)
		S. typhimurium (1)	11 (1)	1 (3117)
		S. dublin (1)	11 (1)	
		S. lomita (1)	32 (1)	
ll poultry (50) ^b		5. Iollita (1)	32 (1)	
ll poultry (50)	Diseased chicken, layer, and broiler (50)	Contoritidis (20)	11 (35), 17 (3), 19 (1),	3 (ST17)
iii poultry (50)	Diseased Chicken, layer, and broner (50)	, ,		• •
		S. indiana (2)	17 (2)	2 (ST17)
		S. gallinarum-pullorum (2)		2 (CT210)
		UT (Vi) (6)	319 (6)	3 (ST319)
Dia industry chains (21)	2	D group (1)	17 (1)	1 (ST17)
Pig industry chains (31)		C dorby (0)	17 (2) 40 (2) 64 (1) 462 (2) 2441 (1)	1 (\$\text{CT17}) 1 (\$\text{CT40}) 1 (\$\text{CT64})
Pig Farms (12) Slaughterhouses(19)	Living donor (12)	S. derby (9)	17 (2), 40 (3), 64 (1), 463 (2), 2441 (1)	1 (3117), 1 (3140), 1 (3164)
		S. ruzizi (1)	40 (1)	
	C(C)	S. new-rochelle (2)	40 (1), 2441 (1)	
	Carcass (6)	S. derby (5)	40 (2), 155 (1), 2441 (2)	
		S. ruzizi (1)	155 (1)	
	Iliac lymph nodes (13)	S. derby (10)	40 (9), 2441 (1)	
		S. ruzizi (1)	463 (1)	
		S. london (1)	463 (1)	
. 1/	. 1 . 1 . (5)4	S. new-rochelle (1)	40 (1)	
Food (raw meat) from h	3 8 ()			. (000.10)
Hotels (7)	Broiler-meat (3)	S. indiana (3)	17 (1),34 (1), 469 (1)	1 (ST17), 1 (ST34)
	Pork(4)	S. derby (3)	40 (3)	
		S.enteritidis (1)	11 (1)	
Animals in 1956, 1976, a	` ,			
Chicken (3)	Chicken (3)	S. enteritidis (3)	11 (2), 132 (1)	
Pigs (20)	Pigs (20)	S. enteritidis (1)	10 (1),	
		S. choleraesuis (9)	68 (7), 139 (1), 145 (1)	
		S. typhimurium (10)	19 (9), 2939 (1)	
Human from different ci			11 (6)	
	human carriers of food employees(44)	S. enteritidis (6)	• •	
		S. typhimurium (11)	19 (5), 34 (5), 40 (1)	1 (ST19), 2 (ST34)
		S. typhimurium (11) S. senftenberg (5)	• •	1 (ST19), 2 (ST34)
		S. typhimurium (11)	19 (5), 34 (5), 40 (1)	1 (ST19), 2 (ST34)
		S. typhimurium (11) S. senftenberg (5)	19 (5), 34 (5), 40 (1) 14 (5)	1 (ST19), 2 (ST34)
		S. typhimurium (11) S. senftenberg (5) S. dublin (2)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2)	1 (ST19), 2 (ST34)
		S. typhimurium (11) S. senftenberg (5) S. dublin (2) S. anatum (3)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2) 64 (3)	1 (ST19), 2 (ST34)
		S. typhimurium (11) S. senftenberg (5) S. dublin (2) S. anatum (3) S. derby (3)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2) 64 (3) 40 (3)	1 (ST19), 2 (ST34)
		S. typhimurium (11) S. senftenberg (5) S. dublin (2) S. anatum (3) S. derby (3) S. london (2)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2) 64 (3) 40 (3) 155 (2) 46 (1)	1 (ST19), 2 (ST34)
		S. typhimurium (11) S. senftenberg (5) S. dublin (2) S. anatum (3) S. derby (3) S. london (2) S. newport (1) S. aberdeen (1)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2) 64 (3) 40 (3) 155 (2) 46 (1) 426 (1)	
		S. typhimurium (11) S. senftenberg (5) S. dublin (2) S. anatum (3) S. derby (3) S. london (2) S. newport (1) S. aberdeen (1) S. rissen (1)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2) 64 (3) 40 (3) 155 (2) 46 (1) 426 (1) 469 (1)	1 (ST19), 2 (ST34) 1 (ST469)
		S. typhimurium (11) S. senftenberg (5) S. dublin (2) S. anatum (3) S. derby (3) S. london (2) S. newport (1) S. aberdeen (1) S. rissen (1) S. bovismorbificans (1)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2) 64 (3) 40 (3) 155 (2) 46 (1) 426 (1) 469 (1) 1499 (1)	
		S. typhimurium (11) S. senftenberg (5) S. dublin (2) S. anatum (3) S. derby (3) S. london (2) S. newport (1) S. aberdeen (1) S. rissen (1) S. bovismorbificans (1) B group (1)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2) 64 (3) 40 (3) 155 (2) 46 (1) 426 (1) 469 (1) 1499 (1) 36 (1)	1 (ST469)
		S. typhimurium (11) S. senftenberg (5) S. dublin (2) S. anatum (3) S. derby (3) S. london (2) S. newport (1) S. aberdeen (1) S. rissen (1) S. bovismorbificans (1) B group (1) C group (5)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2) 64 (3) 40 (3) 155 (2) 46 (1) 426 (1) 469 (1) 1499 (1) 36 (1) 26 (2), 33 (1), 292 (1), 314 (1)	
łuman (44)	human carriers of food employees(44)	S. typhimurium (11) S. senftenberg (5) S. dublin (2) S. anatum (3) S. derby (3) S. london (2) S. newport (1) S. aberdeen (1) S. rissen (1) S. bovismorbificans (1) B group (1) C group (5) E group (2)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2) 64 (3) 40 (3) 155 (2) 46 (1) 426 (1) 469 (1) 1499 (1) 36 (1) 26 (2), 33 (1), 292 (1), 314 (1) 516 (1), 1956 (1)	1 (ST469)
Human (44)	human carriers of food employees(44)	S. typhimurium (11) S. senftenberg (5) S. dublin (2) S. anatum (3) S. derby (3) S. london (2) S. newport (1) S. aberdeen (1) S. rissen (1) S. bovismorbificans (1) B group (1) C group (5) E group (2) S. enteritidis (19)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2) 64 (3) 40 (3) 155 (2) 46 (1) 426 (1) 469 (1) 1499 (1) 36 (1) 26 (2), 33 (1), 292 (1), 314 (1) 516 (1), 1956 (1) 11 (17), 14 (1), 40 (1)	1 (ST469)
Human from different ci Human (44) Foodborne pathents (51)	human carriers of food employees(44)	S. typhimurium (11) S. senftenberg (5) S. dublin (2) S. anatum (3) S. derby (3) S. london (2) S. newport (1) S. aberdeen (1) S. rissen (1) S. bovismorbificans (1) B group (1) C group (5) E group (2)	19 (5), 34 (5), 40 (1) 14 (5) 11 (2) 64 (3) 40 (3) 155 (2) 46 (1) 426 (1) 469 (1) 1499 (1) 36 (1) 26 (2), 33 (1), 292 (1), 314 (1) 516 (1), 1956 (1)	1 (ST469)

Download English Version:

https://daneshyari.com/en/article/5767548

Download Persian Version:

https://daneshyari.com/article/5767548

<u>Daneshyari.com</u>