

**Original Article**

## **Occurrence of multidrug-resistant *Salmonella enterica* serovar Enteritidis isolates from poultry in Iran**

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### **ABSTRACT**

*Salmonella enterica* is recognized as one of the major food-borne pathogens with more than 2,500 serotypes worldwide. The present study addresses antimicrobial resistance of *Salmonella enterica* serovar Enteritidis isolates in Iran. A collection of 151 *Salmonella spp.* isolates collected from poultry were serotyped to identify *Salmonella* Enteritidis. Sixty-one *Salmonella* Enteritidis were subsequently tested against 30 antimicrobials. A high frequency of antimicrobial resistance was observed against nitrofurantoin (n=55, 90.2%) followed by nalidixic acid (n=41, 67.2%), and cephalexin (n=23, 37.7%). Multi-drug resistance were observed in 35 (57.4%) out of 61 isolates. Twenty-six antimicrobial resistance patterns were observed among the 61 *Salmonella* Enteritidis. All isolates were susceptible to ofloxacin, imipenem, enrofloxacin, chloramphenicol, gentamicin, and 3<sup>rd</sup> and 4<sup>th</sup> generation cephalosporins. In conclusion, our results revealed that implementing new policies toward overuse of antimicrobial drugs in Iranian poultry industry are of great importance.

**Keywords:** antimicrobial resistance, resistance profile, multi-drug resistance, *Salmonella enterica*

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### **INTRODUCTION**

*Salmonella enterica* is a major foodborne pathogen with more than 2,500 serotypes (Grimont and Weill, 2007). Although typhoidal *Salmonella* serovars are host restricted (Uzzau et al., 2000), non-typhoidal *Salmonella* (NTS) are zoonotic agents, that are found in a wide range of animal reservoir (Hald et al., 2007) such as poultry, pigs, and cattle that are considered as the common vehicle of human salmonellosis (Braden, 2006). Globally, it is estimated that 93.8 million cases of gastroenteritis due to *Salmonella* species with

155,000 deaths occurs each year (Majowicz et al., 2010). Salmonellosis is mostly self-limiting and treatment with antimicrobials are not recommended, but this is not the case for very young, the elderly and immunocompromised patients (Martin et al., 2004; Varma et al., 2005). It has been proved that salmonellosis driven by drug-resistant strains has led to 30-50% longer duration of illness period, three times higher risk of hospitalization, and three times higher rate of death compared to pan-susceptible *Salmonella* (Molbak, 2004). *Salmonella* Enteritidis is among the most frequently reported serotypes in Iran (Rad et al.,

2008; Emaddi Chashni et al., 2009; Abdollahi et al., 2011). In a resistance trend study between 1996 and 2006 on clinical *Salmonella* Enteritidis isolates an increased level of resistance was observed for nalidixic acid (9.0% to 43.0%) and ceftazidime (3% to 23%) (Ashtiani et al., 2009). In a cross-sectional study including 1,950 fecal specimens from children with diarrhea, 26 *Salmonella* isolates were collected out of which 14 (54.0%) were identified as *Salmonella* Enteritidis. All the isolates were resistant to nitrofurantoin and ten (71.4%) to nalidixic acid. The resistance against streptomycin, co-trimoxazole, and tetracycline as the most common drugs prescribed for salmonellosis were 7.1%, 21.4%, and 28.6%, respectively (Eshraghi et al., 2010). In 2008, Ranjbar et al., detected 60 serogroup D isolates in a collection of 136 *Salmonella* spp.. Further research on resistance against ampicillin, chloramphenicol, co-trimoxazole, and tetracycline showed 6.7%, 3.3%, 8.3%, and 11.7% resistance rates, respectively. This collection demonstrated the highest level of resistance against doxycycline up to 51.7% (Ranjbar et al., 2008). In other continents, a study from New Zealand on antibiotic susceptibility of 1,560 human and 1,505 non-human *Salmonella* isolates between 2002-2009 revealed that almost all isolates were susceptible to ciprofloxacin and gentamicin (Broughton et al., 2010). Of non-human isolates in the United States of America between 1999 and 2003, there was increased sulfisoxazole, decreased tetracycline and fluctuating streptomycin resistance (Kiessling et al., 2007). In Morocco, 150 *Salmonella* Enteritidis isolates obtained between 2000 to 2008 were subjected to seven antibiotics, resulted in 42% of resistance to at least one class of antimicrobial agent with the largest numbers of resistance to nalidixic acid (36%) (Ohmani et al., 2010). Nowadays, progressive increase in resistance rates of microorganisms against antimicrobial agents has complicated control programs of infectious diseases (Hendriksen, 2003a; Okeke et al., 2005). Among the main reasons in emergence of antimicrobial resistance, the role of misuse and overuse of antimicrobial drugs cannot be neglected. In the

condition that, countries face growing trend of non-responsive microorganisms against prescribed treatments, surveillance programs on drug resistance phenomenon will help by implement proper antimicrobial consumption strategies to prevent disastrous consequences of antimicrobial resistance phenomena (Anonymous, 2012a). This study was conducted to determine the level of antimicrobial resistance found in *Salmonella* Enteritidis among Iranian poultry population.

## MATERIALS AND METHODS

**Bacterial Isolates.** During 2011-2012, a total of 585 fecal samples were collected and transferred to the Microbiology Dep. Laboratory of Razi Vaccine and Serum Research Institute Karaj, Iran. All the samples were collected from three different poultry abattoirs in Alborz (No. 35), Marakazi (No. 240), and Fars Provinces (No.310). Isolation and identification of samples were carried out according to standard procedures (Davies et al., 2001; Hendriksen, 2003b). Gram negative short-motile rods with characteristic red slope/yellow butt reaction on TSI (Merck, Darmstadt, Germany) with the production of H<sub>2</sub>S were interpreted presumptively as *Salmonella*. To enable comparative analysis of the results a *Salmonella* Enteritidis type strain (ATCC 13076) was added to our collection.

**Serotyping.** All the biochemically confirmed isolates were serotyped using slide agglutination with standard antisera (Mast, Bootle, England) for somatic and flagellar antigen identification according to the Kauffman-White classification scheme (Grimont and Weill, 2007).

**Antimicrobial Susceptibility Testing.** The Kirby-Bauer disc-diffusion test, was used according to CLSI standards (Anonymous, 2012b). The following antibiotic discs were used: amikacin, amoxicillin, ampicillin, ampicillin-sulbactam, cefazolin, cefepime, cefixime, cefotaxime, ceftizoxim, ceftazidime, ceftriaxone, cefuroximesodium, cephalixin, cephalothin, chloramphenicol, ciprofloxacin, co-trimoxazole, enrofloxacin, florfenicol, furazolidone, gentamicin,

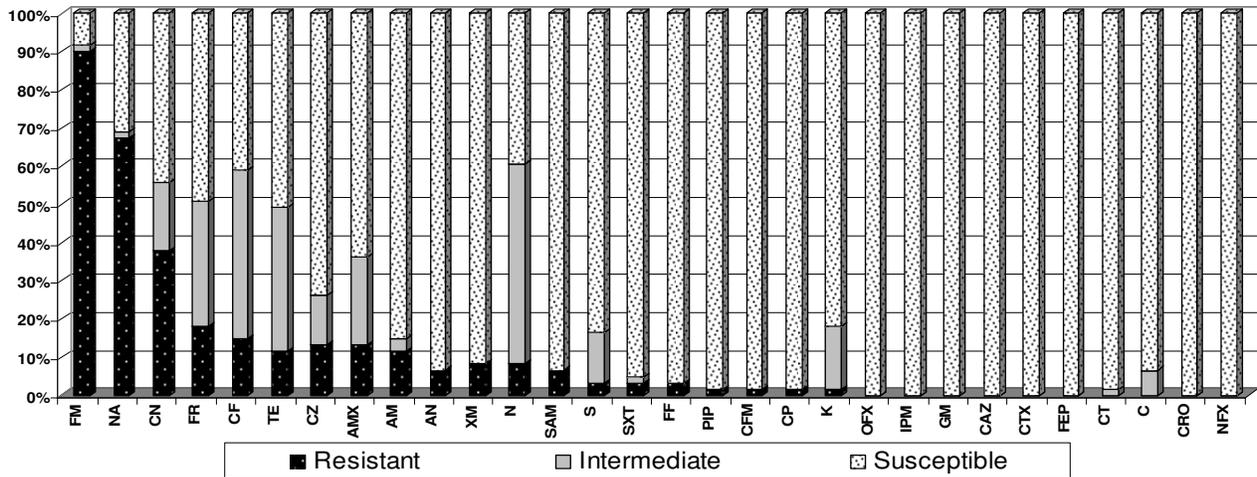
imipenem, kanamycin, nalidixic acid, neomycin, nitrofurantoin, ofloxacin, piperacillin, streptomycin, and tetracycline (Padtan Teb Co. Tehran, Iran). The ATCC reference strain *Escherichia coli* ATCC 25922, was used for quality control purposes.

## RESULTS

One-hundred-fifty-one (25.8%) *Salmonella* isolates out of 585 were analyzed of which, 61 (40.4%) were positive for *Salmonella* Enteritidis. All the isolates originated from three poultry abattoirs comprised of different urban regions; Arak (n = 29, 47.5%), Shiraz (n= 15, 24.6%), Qazvin (n = 10, 16.4%), Takestan (n = 6, 9.8%) and Abhar (n = 1, 1.6%). Antimicrobial susceptibility testing of 30 antimicrobial agents against the entire collection of isolates indicated that most resistance were observed towards nitrofurantoin (90.1%), nalidixic acid (67.2%), and cephalexin (37.7%) followed by furazolidone (18%), cephalothin (14.7%) and tetracycline (11.5%), respectively (Figure 1). On the other hand, the top eleven most susceptible drugs reported against the all isolates includes floxacin, imipenem, gentamicin, ceftazidime, cefotaxime, cefepime, ceftizoxime, chloramphenicol, ceftriaxone, and enrofloxacin (Table 1). Multi-drug resistance against three or more ( $\geq 3$ ) antibiotics was a common phenomenon which were observed with 34 (55.7%) of 61 isolates. Twenty-five antibiotic resistance patterns were detected (Table 2). Overall, this rate of resistance against at least 2 ( $2\leq$ ), 3 ( $3\leq$ ), and 4 ( $4\leq$ ) drugs was 90.1%, 55.7%, and 22%, respectively. The most frequent resistance profile no.7 (Table 2) CN, FM, NA (cephalexin, nitrofurantoin, nalidixic acid) were obtained from 4 (6.5%) isolates. The largest pan-resistant pattern was a panel of 12 drugs consisting NA, SXT, TE, CN, AMX, FM, CZ, FR, AM, XM, CFM, SAM as only one isolate from Arak exhibited this pattern. It is noteworthy, that five (8.1%) of the 61 isolates of *Salmonella* Enteritidis were resistant to seven or more antibiotics, among which two were isolated from Arak and three from Qazvin.

## DISCUSSION

Nowadays, food-borne diseases caused by non-typhoid *Salmonellae* represent an important public health problem. Intestinal salmonellosis will usually resolves spontaneously in five to seven days without the need to exert antibiotic therapy. However, this is not the case for those in which infection spreads beyond the intestinal tract. Hence appropriate and right choice of antimicrobial drugs are of great concern and in rare cases lifesaving (Hohmann, 2001). In this study, *Salmonella* Enteritidis was the predominant serotype isolated, comprising 40.4% of all isolates from five different cities of Arak, Qazvin, Takestan, Shiraz and Abhar. This prevalence rate is in agreement with the recent epidemiologic investigations in Iran (Eshraghi et al., 2010; Zahraei Salehi et al., 2005). Our findings demonstrated that, all isolates were resistant to at least one antimicrobial drug. Multi-drug resistance (MDR) was a remarkable observation in this study. The (CN, FM, NA) was the most prevalent pattern (6.5%) (Table 2) which corroborates our findings that these mentioned antimicrobial agents possess the three most resistance rate (Table 1) recorded against *Salmonella* Enteritidis isolates. Nalidixic acid is the prototype quinolone. It has been available in many countries since the mid 1960, but its use which once has been restricted for a while (Crump et al., 2003), currently is widely prescribed against invasive and systematic infections caused by *Salmonella* spp. (Malorny et al., 1999; Giraud et al., 2006). Some of the researchers believe that use of fluoroquinolone in Veterinary Medicine has an indicative role in emergence and dissemination of nalidixic acid resistance in *Salmonella* in food animals, with high probability of transmission to humans (Angulo et al., 2004). Regarding the importance of this antibiotic from human health point of view and its potential risk of damage to growing cartilage (Bennish and Salam, 1992) we gained the 2<sup>nd</sup> rate of resistance among our poultry study group. Likewise In another study conducted on a human group, this drug has been reported as the 2<sup>nd</sup> most resistant antibiotic (74.1%) after doxycycline



**Figure 1.** Antimicrobial Resistance against 30 drugs

Amikacin (AN), amoxicillin (AMX), ampicillin (AM), ampicillin-Sulbactam (SAM), cefazolin (CZ), cefepime (FEP), cefixime (CFM), cefotaxime (CTX), ceftizoxim (CT), ceftazidime (CAZ), ceftriaxone (CRO), cefuroxime sodium (XM), cephalixin (CN), cephalothin (CF), chloramphenicol (C), ciprofloxacin (CP), co-trimoxazole (SXT), enrofloxacin (NFX), florfenicol (FF), furazolidone (FR), gentamicin (GM), imipenem (IPM), kanamycin (K), nalidixic acid (NA), neomycin (N), nitrofurantoin (FM), ofloxacin (OFX), piperacilin (PIP), streptomycin (S), and tetracycline (TE)

(77.5%) (Firoozeh et al., 2011), which was in agreement with Morshed et al., 2010 investigation on a small group of nine human cases, with (77.8%) resistance rate of nalidixic acid followed by (66.7%) of flumequine. Furthermore In a trend study spanning 2001 till 2005, Ashtiani et al., 2009 showed a gradual increase in resistance to nalidixic acid which correlate with our results. Our study identified the well-known antimicrobial drug of phenicol group, chloramphenicol, demonstrated no sign of resistance which seems, to be due to strict regulations implemented to ban its use on food animals.

Among those antibiotic with no signs of resistance phenomena, ceftazidime and cefixime, can be mentioned which were not in support of Morshed et al. assumption of respectively 6.9% and 24.1% resistance rate in their poultry subgroup of their study. Our findings were in agreement to Mirmahdavi and co-workers to some extent, who worked on antimicrobial resistance of 413 *Salmonella* isolates collected from human clinical cases against five antibiotics, belonged to 3<sup>rd</sup> generation cephalosporin including ceftizoxime, ceftriaxone, ceftazidime, cefotaxime, and cefoperazone.

All, except cefoperazone with 21% resistance rate were pan-susceptible. In another investigation carried out by Shapouri et al. on 240 chickens and eggs, the maximum susceptibility rate (100%) was observed against Gentamicin, Kanamycin and Nalidixic acid, while our findings were in consistent with the first two antimicrobial drugs but in quite contrast with Nalidixic acid which has been recorded as the second most resistant drug (Shapouri et al., 2009). The results obtained in this study showed a similar pattern of antibiotic resistance in the city of Arak and the city of Qazvin for antibiotics CF, FM, NA, CN; however, there are no patterns of antibiotic resistance in other cities. Although Takestan and Abhar are closer in distance to Qazvin, this similarity in pattern between the city of Qazvin & Arak indicate the poultry transportation between the two cities. Furthermore, antibiotic resistance, antibiotic greater than 7, can be seen in these two cities, which can be a reason for the uncontrolled use of antibiotics in these two cities. In conclusion, our results regarding high resistance rate of nitrofurans group of antibiotics in poultry population of our investigation, along with the potential risk of

**Table1.** Antimicrobial resistance in *Salmonella* Enteritidis isolated from Poultry

Antimicrobials	Agent		Resistant No. (Percentages)	
	Class agents	Acronym Disc Content (µg.)	No.	(%)
<b>Aminoglycosides</b>				
Arukacin	AN	30	4	(6.5)
Gentamicin	GM	10	0	(0)
Kanamycin	K	30	1	(1.6)
Neomycin	N	30	5	(8.2)
Streptomycin	S	10	2	(3.3)
<b>Carbapenem</b>				
Imipenem	IPM	10	0	(0)
<b>Cephalosporins</b>				
Cefazolin	CZ	30	8	(13.0)
Cephalothin	CF	30	9	(14.7)
Cefuroxime sodium	XM	30	4	(6.5)
Cephalexin	CN	30	23	(37.7)
Cefixime	CFM	30	1	(1.6)
Cefotaxime	CTX	30	0	(0)
Ceftiozime	CT	30	0	(0)
Ceftazidime	CAZ	30	0	(0)
Ceftriaxone	CRO	30	0	(0)
Cefepime	FEP	30	0	(0)
<b>Nitrofuran</b>				
Furazolidone	FR	100	11	(18.0)
Nitrofurantoin	FM	300	55	(90.2)
<b>Penicillins</b>				
Amoxicillin	AMX	25	8	(13.0)
Ampicillin	AM	10	7	(11.5)
Ampicillin / Sulbactam	SAM	10+5	4	(6.5)
Piperacilin	PIP	100	1	(1.6)
<b>Phenicol</b>				
Chloramphenicol	C	30	0	(0)
Florfenicol	FF	30	2	(3.3)
<b>Quinolones &amp; Fluoroquinolones</b>				
Enrofloxacin	NFX	5	0	(0)
Ofloxacin	OFX	5	0	(0)
Ciprofloxacin	CP	5	1	(1.6)
Nalidixic acid	NA	30	41	(67.2)
<b>Sulfonamide</b>				
Co-trimoxazole	SXT	25	2	(3.3)
<b>Tetracycline</b>				
Tetracycline	TE	30	7	(11.5)
Total =			61	(100)

**Table2.** Multi-drug resistance (MDR) patterns among 61 *Salmonella* Enteritidis isolates

Antibiotype	Antibiotics	No.	%	Source	City
Resistotype 1	AN,FM,NA	3	4.9	Poultry	Arak
Resistotype 2	CF,CN,NA	1	1.6	Poultry	Ghazvin
Resistotype 3	CF,FM,NA	3	4.9	Poultry	Ghazvin & Arak
Resistotype 4	CF,FM,PIP	1	1.6	Poultry	Shiraz
Resistotype 6	CN,FM,FR	1	1.5	Poultry	Shiraz
Resistotype 7	CN,FM,N	1	1.6	Poultry	Shiraz
Resistotype 8	CN,FM,NA	4	6.5	Poultry	Ghazvin & Arak
Resistotype 9	CZ,NA,TE	1	1.6	Poultry	Arak
Resistotype 10	FM,FR,NA	3	4.9	Poultry	Arak
Resistotype 11	FM,NA,TE	1	1.6	Poultry	Arak
Resistotype 12	FM,TE,XM	1	1.6	Poultry	Abhar
Resistotype 13	AN,CN,FM,NA	1	1.6	Poultry	Takestan
Resistotype 14	CN,CZ,FM,FR	1	1.6	Poultry	Shiraz
Resistotype 15	CN,FM,FR,N	1	1.6	Poultry	Shiraz
Resistotype 16	CN,FM,FR,TE	1	1.6	Poultry	Arak
Resistotype 17	CN,FM,N,NA	1	1.6	Poultry	Takestan
Resistotype 18	AM,AMX,CN,CZ,FM	1	1.6	Poultry	Arak
Resistotype 19	AM,AMX,CN,NA,TE	1	1.6	Poultry	Arak
Resistotype 20	AM,AMX,CZ,FM,NA	1	1.6	Poultry	Arak
Resistotype 21	CN,FM,FR,K,S	1	1.6	Poultry	Shiraz
Resistotype 22	AM,AMX,CF,CN,CZ,NA,SAM	1	1.6	Poultry	Ghazvin
Resistotype 23	AMX,FM,FR,NA,S,SXT,TE	1	1.6	Poultry	Arak
Resistotype 24	AM,AMX,CF,CN,CZ,FF,FM,N,NA,SAM	1	1.6	Poultry	Ghazvin
Resistotype 25	AM,AMX,CF,CN,CZ,FF,FM,N,NA,SAM,XM	1	1.6	Poultry	Ghazvin
Resistotype 26	AM,AMX,CFM,CN,CZ,FM,FR,NA,SAM,SXT,TE,XM	1	1.6	Poultry	Arak
Total =		26	34		

resistance against nalidixic acid, demands new efforts to reduce the prevalence of resistant *Salmonella* in food animals. In view of high rate of sensitivity observed among quinolone class except nalidixic acid, carbapenem group and majority of 3<sup>rd</sup> generation of cephalosporins. Strategies including adoption of guidelines for the prudent use of antibacterial agents in animals used for food next to enforcing new food-safety regulations are of great importance. Important investigations that need to be addressed in future include the most frequent serotypes circulating in the region and public health effect of emerging antimicrobial resistance related to those strains.

### Ethics

I hereby declare all ethical standards have been respected in preparation of the submitted article.

### Conflict of Interest

The authors declare that they have no conflict of interest.

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### References

- Abdollahi, A., Najafipour, S., Kouhpayeh, S.A., Meshkibaf, M., 2011. Salmonella Enterica: Serotyping, Drug Resistance & Extended Spectrum of B-Lactamase (ESBLs). *J. Fasa Univ. Med.Sci.* 1, 38-44.
- Angulo, F.J., Nargund, V.N., Chiller, T.C., 2004. Evidence of an association between use of anti-microbial agents in food animals and anti-microbial resistance among bacteria isolated from humans and the human health consequences of such resistance. *J Vet Med B Infect Dis Vet Public Health* 51, 374-379.
- Anonymous, 2012a. Antimicrobial Resistance. In: Organization, W.H. (Ed.), (World Health Organization). World Health Organization, World Health Organization.
- Anonymous, 2012b. Clinical&Laboratory-Standard-Institute, Twenty-Second Informational Supplement performance standards for antimicrobial susceptibility testing, Wayne, PA, USA, p. 188.
- Ashtiani, M.T., Monajemzadeh, M., Kashi, L., 2009. Trends in antimicrobial resistance of fecal Shigella and Salmonella isolates in Tehran, Iran. *Indian J Pathol Microbiol* 52, 52-55.
- Bennish, M.L., Salam, M.A., 1992. Rethinking options for the treatment of shigellosis. *J Antimicrob Chemother* 30, 243-247.
- Braden, C.R., 2006. Salmonella enterica serotype Enteritidis and eggs: a national epidemic in the United States. *Clin Infect Dis* 43, 512-517.
- Broughton, E.I., Heffernan, H.M., Coles, C.L., 2010. Salmonella enterica serotypes and antibiotic susceptibility in New Zealand, 2002-2007. *Epidemiol Infect* 138, 322-329.
- Crump, J.A., Barrett, T.J., Nelson, J.T., Angulo, F.J., 2003. Reevaluating fluoroquinolone breakpoints for Salmonella enterica serotype Typhi and for non-Typhi salmonellae. *Clin Infect Dis* 37, 75-81.
- Davies, P., Turkson, P., Funk, J., Nichols, M., Ladely, S., Fedorka-Cray, P., 2001. Comparison of methods for isolating Salmonella bacteria from faeces of naturally infected pigs. *Journal of applied microbiology* 89, 169-177.
- Eshraghi, S., Dalall, M.M.S., Fardsanei, F., Salehi, T.Z., Ranjbar, R., Nikmanesh, B., Aminharati, F., Abdosamadi, Z., Akbari, A., 2010. Salmonella Enteritidis and antibiotic resistance patterns: a study on 1950 children with diarrhea. *Tehran University Med J* 67, 876-881.
- Firoozeh, F., Shahcheraghi, F., Zahraei Salehi, T., Karimi, V., Aslani, M., 2011. Antimicrobial resistance profile and presence of class I integrons among Salmonella enterica serovars isolated from human clinical specimens in Tehran, Iran. *Iran J Microbiol* 3, 112-117.
- Giraud, E., Baucheron, S., Cloeckaert, A., 2006. Resistance to fluoroquinolones in Salmonella: emerging mechanisms and resistance prevention strategies. *Microbes Infect* 8, 1937-1944.
- Grimont, P.A., Weill, F.X., 2007. Antigenic formulas of the Salmonella serovars WHO Collaboration Center for

- Resistance and Research on Salmonella, Institute Pasteur, Paris
- Hald, T., Lo Fo Wong, D.M., Aarestrup, F.M., 2007. The attribution of human infections with antimicrobial resistant Salmonella bacteria in Denmark to sources of animal origin. *Foodborne Pathog Dis* 4, 313-326.
- Hendriksen, R.S., 2003a. Global Salm-Surv A global Salmonella surveillance and laboratory support project of the World Health Organization- Identification of Salmonella. WHO.
- Hendriksen, R.S., 2003b. Global Salmonella Surveillance, Laboratory protocols: identification of Salmonella. World Health Organization, pp. 1-21.
- Hohmann, E.L., 2001. Nontyphoidal salmonellosis. *Clin Infect Dis* 32, 263-269.
- Kiessling, C.R., Jackson, M., Watts, K.A., Loftis, M.H., Kiessling, W.M., Buen, M.B., Laster, E.W., Sofos, J.N., 2007. Antimicrobial susceptibility of Salmonella isolated from various products, from 1999 to 2003. *J Food Prot* 70, 1334-1338.
- Majowicz, S.E., Musto, J., Scallan, E., Angulo, F.J., Kirk, M., O'Brien, S.J., Jones, T.F., Fazil, A., Hoekstra, R.M., International Collaboration on Enteric Disease 'Burden of Illness, S., 2010. The global burden of nontyphoidal Salmonella gastroenteritis. *Clin Infect Dis* 50, 882-889.
- Malorny, B., Schroeter, A., Helmuth, R., 1999. Incidence of quinolone resistance over the period 1986 to 1998 in veterinary Salmonella isolates from Germany. *Antimicrob Agents Chemother* 43, 2278-2282.
- Martin, L.J., Fyfe, M., Dore, K., Buxton, J.A., Pollari, F., Henry, B., Middleton, D., Ahmed, R., Jamieson, F., Ciebin, B., McEwen, S.A., Wilson, J.B., 2004. Increased burden of illness associated with antimicrobial-resistant Salmonella enterica serotype typhimurium infections. *J Infect Dis* 189, 377-384.
- Molbak, K., 2004. Spread of resistant bacteria and resistance genes from animals to humans--the public health consequences. *J Vet Med B Infect Dis Vet Public Health* 51, 364-369.
- Morshed, R., Peighambari, S.M. 2010. Drug resistance, plasmid profile and random amplified polymorphic DNA analysis of Iranian isolates of Salmonella enteritidis. *New Microbiol* 33, 47-56.
- Ohmani, F., Khedid, K., Britel, S., Qasmaoui, A., Charof, R., Filali-Maltouf, A., El Aouad, R., 2010. Antimicrobial resistance in Salmonella enterica serovar Enteritidis in Morocco. *J Infect Dev Ctries* 4, 804-809.
- Okeke, I.N., Laxminarayan, R., Bhutta, Z.A., Duse, A.G., Jenkins, P., O'Brien, T.F., Pablos-Mendez, A., Klugman, K.P., 2005. Antimicrobial resistance in developing countries. Part I: recent trends and current status. *Lancet Infect Dis* 5, 481-493.
- Rad, M., Kalidari, G., SH., K., 2008. Identification of salmonella spp. In a native poultry breeding and improvement center. *Pajouhesh & Sazandegi* 81, 87-93.
- Ranjbar, R., Naghoni, A., Panahi, Y., Izadi, M., 2008. Antimicrobial susceptibility pattern of salmonella strains isolated from clinical cases against 10 less ordinary prescribed antibiotics in salmonella infection treatment. *Iranian Journal of Infectious Diseases and Tropical Medicine* 46, 41-45.
- Shapouri, R., Rahnema, M., Eghbalzadeh, S., 2009. Prevalence of *Salmonella* serotypes in poultry meat and egg and determine their antibiotic sensitivity in Zanjan city. *The Quarterly Journal of Biological Sciences* 2, 63-71.
- Uzzau, S., Brown, D.J., Wallis, T., Rubino, S., Leori, G., Bernard, S., Casadesus, J., Platt, D.J., Olsen, J.E., 2000. Host adapted serotypes of Salmonella enterica. *Epidemiol Infect* 125, 229-255.
- Varma, J.K., Molbak, K., Barrett, T.J., Beebe, J.L., Jones, T.F., Rabatsky-Ehr, T., Smith, K.E., Vugia, D.J., Chang, H.G., Angulo, F.J., 2005. Antimicrobial-resistant nontyphoidal Salmonella is associated with excess bloodstream infections and hospitalizations. *J Infect Dis* 191, 554-561.
- Zahraei Salehi, T., Mahzounieh, M., Saeedzadeh, A., 2005. The isolation of antibiotic-resistant Salmonella from intestine and liver of poultry in Shiraz province of Iran. pp. 320-322.