

SALMONELLA SEROVARS ISOLATED FOR THE FIRST TIME IN POLAND, 1995–2007

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Abstract

Objectives: Almost each year, a few *Salmonella* serovars isolated for the first time are noted in Poland. Each such serovar should be regarded as potentially dangerous. Recognition of them and monitoring their occurrence and frequency of distribution in all sources is important to control salmonellosis, and is one of the roles of the National *Salmonella* Centre. **Materials and Methods:** Over 2000 *Salmonella* strains submitted to the National *Salmonella* Centre for reference identification were examined. All of them were isolated in Poland between 1995–2007 from human and non-human sources. They were serotyped and characterized biochemically according to standard techniques. The antigenic factors were identified by means of good-quality *Salmonella* specific rabbit antisera. The White-Kauffmann-Le Minor scheme was used to name the serovars. **Results:** One hundred and forty-five serovars were recognized. Fifty-two of them appeared for the first time in Poland. One serovar was found to be the new one. Its validation was done at the WHO Collaborating Centre for Reference and Research on *Salmonella* (Institut Pasteur, Paris, France). It will be included in the next (10th) edition of the White-Kauffmann-Le Minor scheme. **Conclusions:** *Salmonella* will continue to be the feature of humans, animals and the general environment. For practical purposes, we have to accept that *Salmonella* will not be eliminated and we must address our efforts to its controlling. Measuring trends in *Salmonella* serovars over time ought to be done to provide information about the efficacy of prevention and control measures. Results of this study are significant for these activities. They have also allowed to update the list of *Salmonella* serovars isolated in Poland during the last fifty years (1957–2007) which is presented as well.

Key words:

Salmonella, *Salmonella* control, *Salmonella* in Poland, *Salmonella* serovars

INTRODUCTION

Salmonellosis are currently the most important and most common infectious diseases. The genus *Salmonella* is responsible for a wide spectrum of human diseases and affects a wide range of warm- and cold-blooded animals. On a global scale, *Salmonella* is responsible for an estimated 3 billion human infections annually, and typhoid fever makes up 22 million of the cases with approximately 200 000 deaths per year [1]. Data concerning non-typhoidal *Salmonella* (NTS) serovars are notoriously

difficult to obtain, as most patients do not need to consult the health services. In the USA, there are approximately 200 000 infections annually [2]. In England and Wales, between 20 000 and 30 000 cases of human salmonellosis are reported per year, but it is suggested that there are at least three additional cases for every reported one. Within the European Union (EU), figures approach 350 000 reported cases of NTS infection annually with approximately 1000 deaths [3–4]. Fortunately, total case counts of salmonellosis in the EU have decreased since 2004,

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and this decreasing Community trend is statistically significant. The majority of EU member states reported a decrease in *Salmonella* notification rates (2004–2007), and only some of them experienced increase. Using returning Swedish tourists as a sentinel population and Swedish data of travel-associated salmonellosis acquired in Europe from 1997–2003, de Jong et al. [5] estimated the burden of salmonellosis in different European countries. The highest one was revealed for Bulgaria, followed by Turkey and Malta, and *Salmonella* Enteritidis was the dominating serovar. Overall, in the EU, *Salmonella* Enteritidis and *Salmonella* Typhimurium are the serovars most frequently associated with human illness. The epidemiology of *Salmonella* in Poland is parallel to general trends noted in Europe. Statistically significant and decreasing trend in *Salmonella* notification rates (1995–2007) was reported by Dera-Tomaszewska et al. [6]. The decrease in notification rates associated with *Salmonella* Enteritidis infections was also observed. The decreasing *Salmonella* Enteritidis trend was statistically significant as well. None of the remaining, statistically analysed serovars (i.e. *S.* Typhimurium, *S.* Hadar, *S.* Infantis, *S.* Virchow, *S.* Newport, *S.* Mbandaka and *S.* Agona) was found to be significantly increasing or decreasing in frequency over time. The distribution of their notification rates showed rather features of random walk.

Not all *Salmonella* serovars are equally pathogenic for man, but none of them can be considered as essentially harmless. Early, reliable identification is important for the provision of appropriate therapy and to control outbreaks. Monitoring the occurrence and frequency of distribution of all *Salmonella* serovars isolated from humans, animals, food, feeds and other sources is important to detect possible outbreaks, to identify possible sources and routes of transmission, and to take prevention and control measures. Travel and trade restrictions have never proved to be effective in controlling the international spread of *Salmonella* organisms but the sound public health practices

are the most effective approach. Serotyping is a frequently used component of a public health response to the global challenge of salmonellosis. Support for serotyping as the part of national *Salmonella* surveillance, and for rapid international communication of results will help to target future prevention strategies.

In Poland, the National *Salmonella* Centre provides, among others, the reference service for *Salmonella*, including identification of *Salmonella* serovars which appeared in our country for the first time. In this paper, 52 *Salmonella* serovars isolated for the first time in Poland between 1995–2007 are described. Antigenic formula of each such serovar was recognized by the National *Salmonella* Centre and confirmed in the WHO Collaborating Centre for Reference and Research on *Salmonella* (Institut Pasteur, Paris, France).

One of the roles of the *Salmonella* National Reference Centres has been to maintain the list of *Salmonella* serovars identified in their countries. One hundred and fifty-eight *Salmonella* serovars from human and non-human sources were reported in Poland up to the end of 1994. The results of this research work allowed to update this list. Updating and publication of the list is the responsibility of the National *Salmonella* Centre. During the last fifty years (1957–2007), two hundred and ten *Salmonella* serovars were recognized in Poland.

MATERIALS AND METHODS

Isolates

During the 13-year period 1995–2007, a total of 2038 isolates of *Salmonella* were submitted to the National *Salmonella* Centre by field public health, veterinary and other laboratories all over the country for reference identification. They mainly included cultures which were diagnostically difficult to be defined on the field laboratory level and those which were suspected to present serovars not recognized yet in Poland. Of all the examined strains,

972 were human isolates, 932 were isolated from non-human sources (animals, food, feed and feed ingredients, environment) and 134 strains were of unknown origin.

Serotyping and biochemical identification

All *Salmonella* isolates were serotyped and characterized biochemically according to standard techniques [7–10]. The principal biochemical tests by which *Salmonella* can be identified and subspecies of *Salmonella enterica* differentiated were done using validated in-house traditional media in tubes and commercially available tests. The antigenic factors of all examined strains were identified by means of good-quality in-house *Salmonella* specific rabbit antisera, antisera produced by Immunolab (Gdynia, Poland) and Statens Serum Institut (Copenhagen, Denmark). The White-Kauffmann-Le Minor scheme (formerly known as Kauffmann-White scheme) was used to name the serovars [11–12].

Surveillance

Together with *Salmonella* strains, the detailed information necessary from the epidemiological point of view was received by the National *Salmonella* Centre from field laboratories. Despite the inherent biases of such surveillance system, it still remains a good source of information. The bacteriological data are adjusted according to the results from the National *Salmonella* Centre which performed serological typing and biochemical characterization of all received *Salmonella* isolates using standard methods.

RESULTS AND DISCUSSION

Salmonellosis has been the subject of many studies throughout the world since Salmon and Smith, in 1885, isolated the first member of the genus *Salmonella* from an infected pig. A progress was made in the study of these infections after White and Kauffmann established the present method of antigenic analysis of the genus *Salmonella*

and the occurrence of numerous diverse serological variants were recognized. The antigenic diversity is a curiosity of this most numerous taxon within bacteria. The antigenic structures of *Salmonella* allowed to differentiate 2579 serovars within the genus [13], and new variants will continue to be described [14]. Differences of the antigenic structure is the main (although not only) criterion of strain differentiation within the genus *Salmonella*. Serotyping is the original means of *Salmonella* typing allowing for classification by the White-Kauffmann-Le Minor scheme. After initial isolation of *Salmonella* in the diagnostic laboratory, it is still the primary method by which *Salmonella* is described. Currently, the traditional *Salmonella* serotyping scheme according to White-Kauffmann-Le Minor is accepted world-wide as a 'gold standard' for the classification of *Salmonella* below subspecies level.

A total number of 2038 *Salmonella* strains were examined. They were submitted to the National *Salmonella* Centre for reference identification. All of them were isolated in Poland between 1995–2007 from human and non-human sources. One hundred and forty-five serovars were recognized within the examined strains, including 52 which appeared for the first time in our country (Table 1). One of them displayed an antigenic formula not known before. This serological variant was not recognized anywhere in the world yet. The identity of this newly discovered serovar was confirmed, also for the validation reason, by the researches performed by the WHO Collaborating Centre for Reference and Research on *Salmonella* (Institut Pasteur, Paris, France) and two other reference laboratories: Institut für Hygiene und Umwelt (Hamburg, Germany) and Centers for Disease Control and Prevention (Atlanta, USA). According to the current procedure, validation of new *Salmonella* is done at the WHO Collaborating Centre for Reference and Research on *Salmonella* in collaboration with laboratories in Hamburg and Atlanta mentioned above [15]. *Salmonella* serovar is homologated when these three laboratories agree on their validation. Serovar

Salmonella enterica subsp. *diarizonae* (48:k:z₅₇) isolated in Poland in 2007 was confirmed by the above laboratories to be the new one. This third *Salmonella* serovar discovered in Poland (after *Salmonella* Gdansk and *Salmonella* Lodz) will be included in the next (10th) edition of the White-Kauffmann-Le Minor scheme.

All *Salmonella* serovars isolated for the first time in Poland between 1995 and 2007 belonged to the species *Salmonella enterica*. Great majority of them were assigned

to subspecies *enterica* (37 serovars), 6 to subspecies *salmamae* (II), 7 to subspecies *diarizonae* (IIIb), one to subspecies *houtenae* (IV) and one to subspecies *indica* (VI). Thirteen of these first isolations were made only from human and 34 only from non-human sources. Five serovars, namely *Salmonella* Mississippi, *Salmonella* Hvittingfoss, *Salmonella* Isaszeg, *Salmonella* Lindern and *Salmonella* IV (11:z₄₇z₂₄:–) were isolated from both sources. The biochemical behaviour of all the strains was regular and

Table 1. *Salmonella* serovars isolated for the first time in Poland between 1995 and 2007, recognized by the National *Salmonella* Centre

Serovar name	Antigenic formula	Source	Year	Notes
<i>Salmonella enterica</i>				
subsp. <i>enterica</i>				
Agbeni	13,23:g,m:–	human stool (child)	1996	from adult in 2006
Alger	38:l,v:1,2	turtle faeces	1997	
Bardo	8:e,h:1,2	wild boar liver	1998	
Bergen	47:i:e,n,z ₁₅	human stool	2002	from pregnant woman
Berlin	17:d:1,5	henna	1998	
Bracknell	13,23:b:1,6	rape seed	2001	
Canstatt	1,3,19:m,t:–	meat-bone meal	1997	from Spanish pepper in 1998
Finkenwerder	6,14,25:d:1,5	water	2002	
Fischerkietz	1,6,14,25:y:e,n,x	turkey	2000	imported from England
Galiema	6,7:k:1,2	human stools	1998	from child and adult person
Grumpensis	13,23:d:1,7	human bile	1995	case of typhoid-like disease
Halle	28:c:1,7	turtle	2002	
Hull	16:b:1,2	human stool	2005	
Hvittingfoss	16:b:e,n,x	human stool	1998	from dried mushrooms in 1998 and sesame seed in 2002
Idican	13,23:i:1,5	animal meal	1998	also isolated in 2000 and 2001
Invernes	38:k:1,6	human stools	2004	from outbreak of food-poisoning
Isaszeg	48:z ₁₀ :e,n,x	lake water	2003	also from human stool
Istanbul	8:z ₁₀ :e,n,x	goose	1997	
Javiana	9,12:l,z ₂₈ :1,5	black pepper	1996	also isolated in 1999
Kiambu	4,12:z:1,5	feed	1996	
Kibusi	28:r:e,n,x	paprika powder	2000	imported from France
Lindern	6,14,24:d:e,n,x	human stool (child)	2003	from sesame seed in 2004
Liverpool	1,3,19:d:e,n,z ₁₅	sesame seed	1999	from trout's feed in 2000

Table 1. *Salmonella* serovars isolated for the first time in Poland between 1995 and 2007, recognized by the National *Salmonella* Centre – cont.

Serovar name	Antigenic formula	Source	Year	Notes
Llandoff	1,3,19:z ₂₉ ;–	animal meal	2000	from fish meal in 2003 – strain with H:Rz ₃₇ phase
Mississippi	13,23:b:1,5	human stool	1997	after return from Nigeria; and from cocoa pulp in 2001
Nyborg	3,10:e,h:1,7	dried basil	1996	imported from Austria
Oukam	9,46:z ₂₉ ;–	meat meal	1996	together with <i>S. Rissen</i>
Plymouth	9,46:d:z ₆	human stool	1996	after return from Nigeria
Salford	16:l,v:e,n,x	human stool (child)	1996	
Schleissheim	4,12,27:b:	river water	2003	from bathing beach area
Stockholm	3,10:y:z ₆	cocoa-beans pulp	2001	
Suelldorf	45:f,g;–	sesame seed	2001	
Telekebir	13,23:d:e,n,z ₁₅	feed mixture	1998	from broilers
Telhashomer	11:z ₁₀ :e,n,x	feed mixture	2000	
Tilene	1,40:e,h:1,2	human stool (child)	2007	
Uzaramo	1,6,14,25:z ₄ ,z ₂₃ :	human stool	2005	from healthy person
Yoruba	16:c:l,w	chocolate	2006	
subsp. <i>salamae</i>				
	9,12:b:e,n,x	paprika powder	1998	imported from Spain
	13,22;z ₂₉ :1,5	human stools	2002	2 children and adult person
	41:z:1,5	pasta	2001	imported from Italy
	42:b:e,n,x,z ₁₅	figs	2003	imported from Iran
	42:g,t;–	henna	1998	
	48:d:z ₆	turtle droppings	1998	
subsp. <i>diarizonae</i>				
	35:i:z ₃₅	viper organs	1998	
	38:r:z	human stool (child)	1995	
	48:k:z ₅₃	human stool (child)	2007	
	48:k:z57 ^a	river water	2007	from unguarded beach area
	58:z ₅₂ :z ₃₅	viper organs	1998	also from beaver faeces
	60:z ₅₂ :z ₅₃	viper organs	1998	
	61:k:1,5,7	human cerebrospinal fluid	2007	together with <i>Streptococcus pneumoniae</i> – case of purulent meningitis
subsp. <i>houtenae</i>				
	11:z ₄ ,z ₂₃ ;–	chameleon intestine	1997	from viper organs in 1998; from human stool in 2004 and 2005
subsp. <i>indica</i>				
	45:a:e,n,x	sesame seed	2002	

^a New serovar; its antigenic formula is not recorded in the current (9th) edition of the White-Kauffmann-Le Minor scheme [11]; it will be included in the next issue of the scheme.

was classified as typical for the genus *Salmonella*. Results of biochemical examination in dulcitol, lactose, L(+)-tartrate (= *D*-tartrate), mucate, malonate, salicin, gelatinase, galacturonate, β -glucuronidase and β -galactosidase (ONPG) activity tests, culture with potassium cyanide and lysis by phage O1 allowed to define serovar association with respective species or subspecies.

Unfortunately, human salmonellosis in Poland are also caused by serovars of subspecies other than *Salmonella enterica* subsp. *enterica*. Members of subspecies *salamae*, *diarizonae* and *houtenae* have been responsible for *Salmonella* infections in humans. Six of such serovars, including four of those which did not appear previously in Poland, were isolated during the reported period. Strains of subspecies *diarizonae*, like the majority of its representatives, showed the ability to ferment lactose. The participation of such serovars in causing infections of humans in our country may constitute more serious problem that it is suspected. The use of traditional media for *Salmonella* isolation as the only ones in official diagnostic procedures may result in omitting the bacteria's presence in the examined materials.

Appearance of new (in the meaning: occurring for the first time as well as new-discovered) *Salmonella* serological variants constitutes a major hazard, because people are at risk to be infected with microorganisms which are unknown for us, and which have not caused infections in our country yet. The fact that very young children, mainly infants, were involved in these infections, is especially worrisome. Such serovars as *Salmonella* Agbeni, *Salmonella* Galiema, *Salmonella* Lindern, *Salmonella* Salford, *Salmonella* Tilene, *Salmonella* II (13,22:z₂₉:1,5), *Salmonella* IIIb (38:r:z) and *Salmonella* IIIb (48:k:z₅₃) were etiologic agents of infections of children in the 4- to 13-month age range. Unfortunately, in great majority of these cases, the source of infection was unknown or assumed only, like it was in *Salmonella* Lindern-caused severe systemic infection (exposure to turtles).

Some of *Salmonella* serovars presented in this paper which were recognized as etiologic agents of animal salmonellosis, especially of imported animals, are rare in Poland. Such serovars as *Salmonella* Alger, *Salmonella* Bardo, *Salmonella* Fischerkietz, *Salmonella* Halle, *Salmonella* Istanbul, *Salmonella* II (48:d:z₆), *Salmonella* IIIb (35:i:z₆), *Salmonella* IIIb (58:z₅₂:z₃₅), *Salmonella* IIIb (60:z₅₂:z₅₃), *Salmonella* IV (11:z₄:z₂₃:–) just occurred for the first time. Although represented by single isolates, they constitute a hazard to human health, the more so that their ecology and pathogenesis of infection have not been completely explained. The behaviour of each such microorganism in one particular animal species is not necessarily predictive of its behaviour in another host species.

Animal feed is a recognized source of pathogenic microorganisms for animals [16]. It is still relatively common to find evidence of contamination of domestic and imported feed and animal-feed ingredients [17–26]. *Salmonella* can be isolated regularly from feedstuffs and feed ingredients, including both animal and vegetable proteins, such as soya, rape, palm kernel, rice bran and cottonseed. Feed appears as a major potential route by which new infections may be introduced into e.g., farm livestock, particularly poultry. Recent evidence presented in this paper indicates that a high proportion of both domestic and imported bone meal, meat meal, fish meal, and similar protein supplements used in animal and poultry feeds are contaminated with *Salmonella*. Specific feed-components may also be implicated in outbreaks of salmonellosis. *Salmonella* Mbandaka, *Salmonella* Infantis, *Salmonella* Havana, *Salmonella* Muenster and other serovars were defined in this study within strains isolated from them. The feedstuff and animal-feed ingredients were also the source of nine *Salmonella* serovars that previously had not been reported in Poland. Serovars *Salmonella* Bracknell, *Salmonella* Cannstatt, *Salmonella* Idican, *Salmonella* Kiambu, *Salmonella* Liverpool, *Salmonella* Lladoff, *Salmonella* Okam, *Salmonella* Telelkebir and *Salmonella* Telhashomer

appeared for the first time in our country between 1995 and 2007. Taking into consideration the scope of animal production and the feeding habits, it is easy to imagine how important in *Salmonella* dissemination the contaminated feeds are.

One hundred and fifty-eight *Salmonella* serovars isolated from human and non-human sources were reported in Poland up to the end of 1994. Fifty-two other serovars were identified during the next 13-year period as a result of this research work which allowed to update the list. During the last fifty years (1957–2007), two hundred and ten *Salmonella* serovars from 30 serological groups were recognized in Poland. They are presented in Table 2.

Group O:4 occupied the major part (16%) and together with groups O:7 (13%), O:8 (14%), O:9 (8%), O:3,10 (10%) and O:13 (6%) covered about 70% of these 210 serovars. Most of them have been isolated from both humans and animals, but some have so far been isolated only from human and others only from non-human sources [27–38]. Some of them were imported (from Asia, Africa, both Americas and from European countries) but the origin of most of the others has remained unknown. Many serovars produce only a very small number of infections year after year, but never entirely disappear. There are only comparatively few that have become well established in either humans or animals or both and

Table 2. *Salmonella* serovars isolated in Poland from human and non-human sources, 1957–2007^a

Group	Serovars	
	name	n
O:2	Paratyphi A	1
O:4	Kisangani, Bispebjerg, Paratyphi B, Abony (and <i>S. abortusovis</i> now combined with Abony), II (1,4,12,27:b:e,n,x), Schleissheim, Stanley, Schwarzengrund, Duisburg, Saintpaul, Reading, Chester, Sandiego, Derby, Agona, Essen, Hato, California, Kingston, Banana, Typhimurium, Agama, Ljubliana, Bredeney, Brandenburg, Kunduchi, Heidelberg, Coeln, Kiambu, Indiana, Stanleyville, Haifa, Abortusequi	33
O:7	Oslo, Coleypark, Brazzaville, Ohio (and its var. 14 ⁺), Paratyphi C, Choleraesuis, Isangi (and <i>S. mission</i> now combined with <i>S. Isangi</i>), Livingstone, Norwich, Braenderup, Rissen, Montevideo, Oranienburg, Galiema, Thompson, Singapore, Concord, Potsdam, Gdansk, Virchow, Infantis, Bareilly, Hartford, Oakland, Mbandaka, Jerusalem, Tennessee, Lille (and its var. 14 ⁺)	28
O:8	Tado, Virginia, Muenchen, Manhattan, Bardo, Newport, Kottbus, Tshiongwe, Emek, Yokoe, Takoradi, Bonariensis, Kentucky, Blockley, Haardt, Lichfield, Manchester, Breukelen, Hindmarsh, Bovismorbificans, Goldcoast, Altona, Inchpark, Chailey, Corvallis, Albany, Istanbul, Hadar, Glostrup, Molade	30
O:9	Miami, Saarbruecken, II (1,9,12:b:e,n,x), Durban, Typhi, Eastbourne, Berta, Enteritidis, Blegdam, Dublin, Kapemba, Javiana, Gallinarum (and biovar Pullorum)	16
O:9,46	Plymouth, India, Oukam, Fresno	4
O:3,10	Oxford, Butantan (and its var. 15 ⁺), Onireke, Vejle, Muenster (and its var. 15 ⁺), Anatum (and its var. 15 ⁺), Nyborg, Newlands, Meleagridis (and its var. 15 ⁺), Amsterdam (and its var. 15 ⁺), Westhampton (and its var. 15 ⁺), Falkensee, Zanzibar, Nchanga, London (and its var. 15 ⁺), Give (and its var. 15 ⁺), Uganda (and its var. 15 ⁺), Elizabethville, Weltevreden, Orion (and its var. 15 ⁺ and var. 15 ⁺ ,34 ⁺), Stockholm, Lexington	22
O:1,3,19	Liverpool, Senftenberg, Cannstatt, Taksony, Westerstede, Krefeld, Llandoff, Dessau	8
O:11	Adamstua, Stendal, Rubislaw, IV (11:z ₄ ,z ₂₃ :–), Telhashomer	5
O:13	Ibadan, Mississippi, Bracknell, Grumpensis, Telelkebir, Havana, Agbeni, Idican, Kedougou, Poona, Worthington, II (13,22:z ₂₉ :1,5), Cubana	13
O:6,14	Heves, Finkenwerder, Lindern, Fischerkietz, Uzaramo	5

Table 2. *Salmonella* serovars isolated in Poland from human and non-human sources, 1957–2007^a – cont.

Group	Serovars	
	name	n
O:16	Brazil, Hull, Hvittingfoss, Yoruba, Gaminara, Salford	6
O:17	Berlin	1
O:18	Cerro	1
O:21	Minnesota	1
O:28	Halle, Kibusi, Pomona	3
O:30	Urbana, Morehead	2
O:35	Adelaide, Anecho, IIIb (35:i:z ₃₅), Alachua	4
O:38	Thiaroye, Inverness, Alger, IIIb (38:r:z)	4
O:40	Johannenburg, Tilene	2
O:41	II (41:z:1,5), Waycross, Lodz	3
O:42	II (42:b:e,n,x,z ₁₅), II (42:g,t:-)	2
O:43	IV (43:z ₄ :z ₂₃ :-)	1
O:45	VI (45:a:e,n,x), Suelldorf	2
O:47	II (47:a:1,5), Bergen, Alexanderplatz	3
O:48	II (48:d:z ₀), IIIb (48:k:z ₅₃), IIIb (48:k:z57) ^b , IIIb (48:r:e,n,x,z ₁₅), IV (48:z ₄ :z ₂₃ :-), Isaszeg	6
O:50	IIIb (50:k:z ₅₃)	1
O:58	IIIb (58:z ₅₂ :z ₃₅)	1
O:60	IIIb (60:z ₅₂ :z ₅₃)	1
O:61	IIIb (61:k:1,5,7)	1

^a The order of groups and serovars within the groups – according to the listing in the current (9th) edition of the White-Kauffman-Le Minor scheme [11]; var. = variant.

^b New serovar; its antigenic formula is not recorded in the current edition of the White-Kauffmann-Le Minor scheme; it will be included in the next issue of the scheme.

continue being the principal contributors to *Salmonella* infections in our country every year. More and more numerous appearance of *Salmonella* serovars which have not been isolated in our country, may become a serious social and economic problem in the future. Each such *Salmonella* serovar means a potential epidemic threat and may cause an epidemic at any time.

All possible steps should be taken to monitor the occurrence of *Salmonella* organisms and the frequency of distribution of all serovars, not only the epidemic ones. Difficulties in the control are exacerbated by travelling and the global trade in food and food animals, which may assist in the movement of *Salmonella* serovars. The

ubiquity of *Salmonella* serovars and the frequency and rapidity with which the populations can change mean that it will never be possible to eradicate these bacteria. Measuring trends in serovars over time ought to be done to provide information about the efficacy of prevention and control measures. More complete knowledge of the relative prevalence of different *Salmonella* serovars, by time and place, is of high importance both from national and international standpoint. Global joint actions and involvement of all countries in this strategy will allow to reduce the risk of *Salmonella* infections. The results of the research work presented in this paper are significant for these activities.

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