

# Livestock salmonellosis in the Irkutsk region

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**Abstract.** The analysis of the incidence of livestock (cattle and pigs) of salmonellosis in the territory of the Irkutsk region during 2004–2016 was carried out. It was found that the incidence rate of cattle was  $5.4 \pm 0.62$  per 100 thousands livestock, pigs –  $4.0 \pm 0.04$  <sup>0/0000</sup>. The dynamics of reducing of the incidence of pigs was identified. Negative rates of increase in the incidence of animals were revealed. The epizootic index was 1.0. Salmonella of nine species (*Salmonella dublin*, *S. enteritidis*, *S. choleraesuis*, *S. typhimurium*, *S. london*, *S. lindi*, *S. wernigerode* and *S. othmarschen*) in cattle and five species (*S. choleraesuis*, *S. enteritidis*, *S. typhimurium*, *S. rochdale* and *S. arizonae*) in pigs were identified as the causative agent of the infection. *S. dublin* was dominated at the etiology of salmonellosis in cattle (75.1 %) ( $p \leq 0.05$ ), *S. choleraesuis* – in pigs (89.2 ± 2.9 %) ( $p \leq 0.01$ ). Salmonellosis in cattle was recorded in 11 districts of the Irkutsk region, in pigs – in nine districts.

## 1 Introduction

Salmonellosis, widespread in many countries of the world, is a serious threat to human health [1]. As a result of Salmonella infection, about 94 million cases of gastroenteritis in people are recorded annually in the world, and in hundreds of thousands of cases the disease is fatal [2]. In the United States alone, about 1.4 million cases of salmonellosis among the population are observed each year [3], while economic damage is \$ 2.5 billion [4]. According to the report of the European Food Safety Authority (EFSA) for 2014, salmonellosis is one of the most common zoonoses and foodborne toxic infections [5, 6]. About 8 % of all cases of human salmonellosis refer to direct contact with animals; 13 % are related to environmental sources, 14 % to travel; about 55 % – with food products and about 9 % – with direct transmission from person to person [1, 7]. The relevance of salmonellosis is determined by the fact that at present about 20 % of the world's population is employed in the livestock sector. Salmonella is widespread in herds of farm animals, mainly among those animals that are used by humans for food [6]. Cattle (cattle) play a primary role as a source of foodborne infection [7]. In Russia, salmonellosis of cattle (especially calves), pigs and sheep is registered in many regions [8, 9, 10], making up 15–45 % in different territories in the structure of the incidence of zoonoses in farm animals.

Currently, there is a wide list of salmonellosis pathogens (about 2400 serovars, of which 150 are diagnosed in Europe) [11]. The leading role in the etiological structure of animal salmonellosis belongs to *Salmonella enteritidis* (35.9 %), *S. typhimurium* (13.2 %), *S. dublin* (11.2 %), *S. choleraesuis* (10.1 %)

[6]. At the same time, *S. typhimurium* and *S. enteritidis* are, in epidemiological terms, the most relevant for humans. Serovars *S. panama*, *S. infantis* are also significant for humans. *S. Newport*, *S. agona*, *S. london* and others [6]. A number of *Salmonella* serovars enter Russia with imported food and raw materials [12].

*Salmonella* serovars vary greatly in terms of host range. Most of them are adapted to a certain circle of owners. The adapted salmonella serovars in cattle include *S. dublin* [13], in pigs – *S. choleraesuis* [11]. In lambs, the most common cause of abortion, stillbirth, and disease are *Salmonella S. abortusovis* [14; 15]. Pathogens such as *S. typhimurium* and *S. enteritidis* can cause diseases in a wide range of hosts [16, 17]. *Salmonella S. typhimurium* is often associated with the disease in many animal species [17].

Along with the dominant role of *S. dublin*, pathogens such as *S. typhimurium*, *S. enteritidis*, *S. choleraesuis*, *S. typhimurium*, *S. abortusovis*, *S. london*, *S. anfo*, *S. lindi* and others in the etiology of cattle salmonellosis [18]. As for sheep, other *Salmonella* serovars, including *S. typhimurium*, *S. dublin*, *S. anatum* and others, have also been described as causative agents of abortion [14, 16, 19]. According to O. Alvseike et al. (2002), *S. diarizonae* is another common sheep-adapted serovar (its prevalence in Norwegian sheep is about 12 %).

It should be emphasized that in the last 20–30 years there has been a change in the spectrum of salmonella pathogens in various animal groups: against the background of a decrease in the proportion of dominant salmonella serovars, an increase in the significance of those pathogens that previously did not play a significant role in the etiology of this infection [11]. Due to the fact that the clinical form of salmonellosis, the severity is largely determined by the serovar of the pathogen, the

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study of the etiological spectrum of the pathogens of salmonellosis in animals in the long term is of great practical importance [7].

The distribution of *Salmonella* serovars among farm animals varies greatly in time and differs between geographical regions, age groups, clinical manifestations, and production systems [7]. Moreover, salmonellosis is a major socio-economic problem for both developing and developed countries. The economic damage from salmonellosis is caused by both high mortality of young animals and loss of productivity, as well as the costs associated with treatment and infection control, which makes salmonellosis one of the most economically important diseases of both cattle and small ruminants [7]. The above indicates that measures aimed at combating this infection should be carried out on a global scale [2, 16].

## 2 Materials and methods

The study and analysis of the incidence of cattle and pigs salmonellosis in the Irkutsk region was carried out using materials from reporting forms of the Federal State Budgetary Institution “Irkutsk Interregional Veterinary Laboratory” (IMVL) according to the results of studies conducted in IMVL, as well as in Tulunskaya, Ust-Udinsk, Usolskaya, Bratsk, Nizhneudinsk, Chermkhov, Kuitun and Kachug laboratories of stations for combating animal diseases (2004-2016). A total of 7851 studies were completed. The results were processed according to standard statistical methods [20]. The incidence rate was determined per 100,000 livestock (‰).

The tendency (over the study period) of the incidence of salmonellosis was evaluated according to a parabola of the first degree and coefficient *c*. To identify the significance of this trend, the Pearson coefficient (*r*) was used. The differences were statistically significant at  $p \leq 0.05$ . The study of the tension of the epizootic situation of salmonellosis was carried out using the epizootic index (IE).

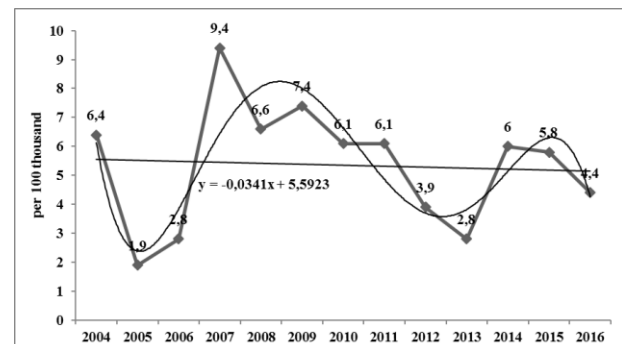
The conditional dominance scale was used to assess the degree of dominance of *Salmonella* of various species isolated from livestock. Four classes were used (absolute dominants, dominants, subdominants and rare salmonella) according to the proportion of bacteria of a certain species in the total number.

## 3 Results and discussion

It has been established that salmonellosis is a relevant disease in cattle and pigs in the Irkutsk region. This infection in livestock was recorded annually and was characterized by coverage of 30 % of the administrative areas of the study area. In this case, the average long-term incidence rate of salmonellosis of cattle in 2004–2016 turned out to be at the level of  $5.4 \pm 0.62$  ‰, pigs –  $4.0 \pm 0.04$  ‰. The incidence of cattle was characterized by variability without significant dynamics ( $r = -0.06$ ;  $p > 0.05$ ).

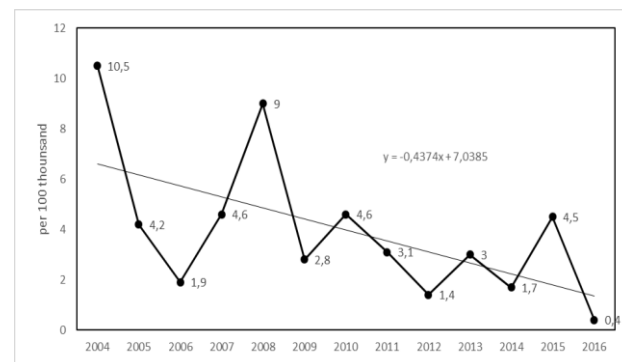
The highest incidence of salmonellosis occurred in 2007. (9.40 / 0000). Since 2008 in 2013, there was a

decrease in the incidence rate to 2.80 / 0000 in 2013. In recent years of observation in the study area, there has been a slight increase in the incidence rate of cattle salmonellosis; the indicators ranged from 6.00/0000 (2014) to 4.40 / 0000 (2016) (Fig. 1).



**Fig. 1.** Dynamics of the incidence of salmonella infection in cattle (Irkutsk region, 2004–2016)

Significant dynamics in reducing the incidence of pig salmonellosis was identified ( $r = -0,585$ ;  $p \leq 0,05$ ). The highest incidence rates of pigs were observed in 2004 (10,5 ‰) and 2008. (9,0 ‰); the lowest – at the end of the study period (0,4 ‰) (Fig. 2).



**Fig. 2.** Dynamics of the incidence of salmonella infection in pigs (Irkutsk region, 2004–2016)

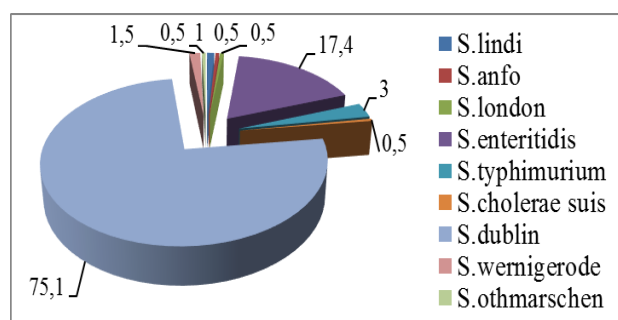
The growth rate of the incidence of salmonellosis in cattle and pigs was negative. The rate of decline in the incidence of pigs was higher compared to that in cattle («-12.8 %» and «-0.7 %»).

The tension of the epizootic situation for cattle and pigs salmonellosis is reflected by an epizootic index of 1.0.

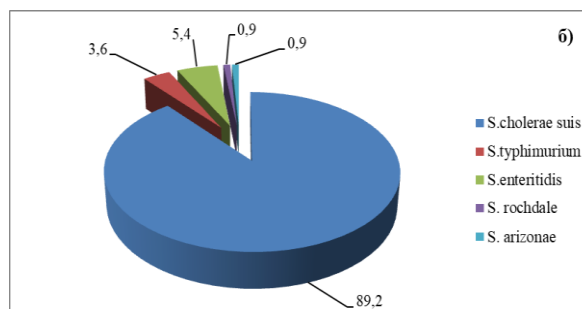
An analysis of the taxonomic spectrum of the etiological agents of salmonellosis revealed that salmonella isolated from cattle during 2004 – 2016 are represented by isolates of nine species: *Salmonella dublin*, *S. enteritidis*, *S. choleraesuis*, *S. typhimurium*, *S. london*, *S. anfo*, *S. lindi*, *S. wernigerode*, and *S. othmarschen*. *Salmonella S. dublin* ( $75.1 \pm 3.0$  %) played a major role in the incidence of cattle. Their share was significantly ( $p \leq 0.05$ ) higher than the frequency of occurrence of salmonella of other species (fig. 3).

*Salmonella* of five species (*S. choleraesuis*, *S. typhimurium*, *S. enteritidis*, *S. arizonae* and *S. rochdale*) were isolated in pigs. *Salmonella choleraesuis* were dominated at the etiology of pig

salmonellosis ( $89.2 \pm 2.9\%$ ) ( $p \leq 0,05$ ). Salmonella of other species was much less often (fig. 4).



**Fig. 3.** Taxonomic spectrum of cattle salmonellosis pathogens in the Irkutsk region, % (2004–2016)



**Fig. 4.** Taxonomic spectrum of pig salmonellosis pathogens in the Irkutsk region, % (2004–2016)

**Table 1.** Assessment of the degree of dominance of Salmonella isolated from cattle (according to the conditional scale of dominance).

Year	Class boundaries for the proportion of microorganisms of different species			
	$64 \leq N < 100$	$36 \leq N < 64$	$14 \leq N < 36$	$0 \leq N < 14$
	Class (by degree of dominance)			
	absolute dominant	dominant	subdominant	rare
2004	S.dublin	«-»	«-»	«-»
2005	S.dublin	«-»	«-»	«-»
2006	S.dublin	«-»	«-»	«-»
2007	«-»	S.dublin S.enteritidis	«-»	S.typhimurium
2008	«-»	S.dublin	S.enteritidis	S.london
2009	S.dublin	«-»	«-»	S.typhimurium
2010	S.dublin	«-»	«-»	S.lindi S.anfo S.cholerae suis
2011	S.dublin	«-»	«-»	«-»
2012	S.enteritidis	«-»	«-»	«-»
2013	«-»	S.enteritidis	S.typhimurium S.dublin	«-»
2014	S.dublin	«-»	«-»	«-»
2015	S.dublin	«-»	«-»	S.wernigerode
2016	S.dublin	«-»	«-»	S.othmarschen
For the entire observation period	S.dublin	«-»	S.enteritidis	S.lindi S.anfo S.london S.typhimurium S.cholerae suis S.wernigerode S.othmarschen

note.: «-» – were absent

The evaluation of the degree of dominance of Salmonella isolated from cattle, according to the conditional dominance scale, showed that S.dublin isolates constituted the class of absolute dominants (class boundaries in proportion from 64 to 100 %) for nine years (with a retrospection depth of 13 years). Salmonella S.enteritidis, according to the share of bacteria of this species in the total number, in some years belonged to the classes of absolute dominants (2012), dominants (2007, 2013) and subdominant (2008). In 2013, S. typhimurium strains were included in the subdominant class (class boundaries in a fraction of 14

to 36 %), and in other years they belonged to the class of rare species. Salmonella of all other species was included in the class of rare species (class boundaries in a fraction of 0 to 14 %) in the general structure of salmonellosis pathogens (table 1).

The results of the studies showed a slight change in the spectrum of salmonellosis pathogens during the observation period. Thus, strains of S.dublin, S. enteritidis and S typhimurium, as etiological agents of salmonellosis, were isolated during the entire study period. At the same time, strains S. london, S. anfo, S. lindi, S. choleraesuis were verified only in the first

observation period (2004 – 2010), and in the subsequent (2011 – 2016) isolates replaced them: *S.wernigerode* and *S.othmarschen*.

*Salmonella choleraesuis* were constituted at class of absolute dominants among *Salmonella* in pigs for 11 years. *Salmonella enteritidis* were entreated at the class of dominants in 2009; these isolates belonged to the class of subdominant and the class of rare species in other years (2006, 2014, 2015). *Salmonella typhimurium* were part of the subdominant class during the all study period.

A study of the territorial distribution of salmonellosis in cattle showed its presence in 11 districts of the Irkutsk region (Tulunsky, Osinsky, Nizhneudinsky, Bohansky, Bratsky, Zalarinsky, Irkutsky, Kuytunsky, Usolsky, Ekhirit-Bulagatsky and Cheremkhovsky). At the same time, about half of all cases of salmonella infection in the studied group of animals were detected in the Cheremkhovsky district. Every tenth case of salmonellosis is diagnosed in the Usolsky, Kuitunsky and Ekhirit-Bulagat districts. In other administrative territories of the region, salmonellosis in cattle was recorded less frequently: this indicator varied from 0.5 % to 3.0 %. Salmonellosis in pigs has been reported in nine districts (Tulunsky, Nizhneudinsky, Bohansky, Irkutsky, Usolsky, Ekhirit-Bulagatsky and Cheremkhovsky); more often the infection was diagnosed in pigs in the Nizhneudinsky, Usolsky and Tulunsky districts.

## 4 Conclusion

As a result of the studies, it was found that in 2004–2016 in the studied area, the incidence rate of salmonella infection of cattle was  $5.4 \pm 0.62$   $^{0}/_{0000}$ , pigs –  $4.0 \pm 0.04$   $^{0}/_{0000}$ . The highest rate was found in 2007 ( $9.4^{0}/_{0000}$ ). The recession took place in 2005 ( $1.9^{0}/_{0000}$ ). The incidence of pigs was characterized by a dynamics of decline: from  $10.5^{0}/_{0000}$  in 2004 to  $0.4^{0}/_{0000}$  in 2016. The increase in the incidence rate was characterized by negative rates. The tension of the epizootic situation for this infection of the studied group of animals is evidenced by a high epizootic index (1.0). Nine species of *Salmonella* were verified as etiological agents in cattle, five species were verified in pigs. *Salmonella* Dublin were prevailed ( $75.1 \pm 3.0$  %) ( $p \leq 0.05$ ) in the etiology of salmonellosis in cattle, *S.choleraesuis* in pigs ( $89.2 \pm 2.9$  %) ( $p \leq 0.01$ ), making up the class of absolute dominants. Salmonellosis in cattle was recorded in 11 regions of the study region. Most often in the Cheremkhovsky, Ekhirit-Bulagat, Kuytun and Usolsky districts. Salmonellosis in pigs was detected in nine districts (mainly in the Lower Udinsky, Usolsky and Tulunsky districts).

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