

REVIEW

Open Access



Rethinking the fight against pig-related human salmonellosis in the European union

Raúl C. Mainar-Jaime^{1*}, A. Casanova-Higes¹, María Bernad-Roche², Juan P. Vico³ and S. Andrés-Barranco⁴

Abstract

The prevalence of human salmonellosis associated with pork products remains a significant concern for public health authorities within the European Union. Despite the implementation of national programs in some member states with the objective of controlling the infection of *Salmonella* in farms, the proportion of human cases involving swine-associated *Salmonella* serotypes has remained constant in recent years, and the majority of these programs were either discontinued or reduced to biosecurity guidance. This article discusses the reasons for the lack of success of these programs, including the focus on the growing-finishing period without consideration of earlier stages of production, the structure of the pig sector, the limited and unrepresentative sampling carried out in the programs, and the use of imperfect serological tests, which have likely resulted in biased estimates of the true health status of the herds. A potential comprehensive approach is proposed, based on predicting the risk of *Salmonella* shedding prior to the arrival of pigs at the slaughter. This knowledge would be combined with the administration of on-farm additives (i.e. organic acids, bacteriophages) during the days prior to slaughter. It would help to reduce shedding in those batches with a high risk of shedding and decrease slaughter environmental contamination. Furthermore, this approach would contribute to obtain more accurate information regarding the *Salmonella* status of the pig farms.

Keywords Control programs, Foodborne zoonoses, Prediction, *Salmonella*, Slaughter, Swine

Introduction

Salmonella enterica has been infecting humans in western Eurasia for over 5,000 years [1]. Human-adapted serotypes (i.e., Typhi and Paratyphi) are the most virulent serotypes for humans, causing the so-called typhoid fever. This disease claims millions of cases and 200,000 deaths annually, mostly in low- and middle-income countries where poor sanitation and deficient hygiene infrastructure persist [2]. In high-income countries, however, *Salmonella* has found its way through the non-typhoidal *Salmonella* (NTS) serovars. Infection with these non-host-adapted serovars typically results in self-limiting diarrheal disease with a low case fatality rate.

In the EU, 77,486 cases were officially recorded in 2023 [3], but the actual number is likely to be much higher and could cost up to €3 billion per year, with significant

*Correspondence:

Raúl C. Mainar-Jaime
rcmainar@unizar.es

¹Departamento de Patología Animal, Facultad de Veterinaria, Instituto Agroalimentario de Aragón-IA2, Universidad de Zaragoza-CITA, Zaragoza 50013, Spain

²Pig and Poultry Development Department, The Irish Food and Agriculture Authority, Moorepark, Fermoy, Co. Cork P61 C996, Teagasc, Ireland

³Facultad de Ciencias Agropecuarias, IRNASUS-CONICET-Universidad Católica de Córdoba, Universidad Católica de Córdoba, Córdoba 5000, Argentina

⁴Departamento de Ciencia Animal, Centro de Investigación y Tecnología Agroalimentaria de Aragón, Instituto Agroalimentario de Aragón-IA2, CITA-Universidad de Zaragoza, Zaragoza 50059, Spain



© The Author(s) 2025. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>. The Creative Commons Public Domain Dedication waiver (<http://creativecommons.org/publicdomain/zero/1.0/>) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

costs arising from hospitalizations, productivity losses, and public health interventions [4–6]. Globally, NTS may cause over 95 million of cases and more than 50,000 deaths [7], with increasing prevalence in developed regions such as the EU, and in many developing countries [3, 8]. These trends are associated with the widespread and emergence of new virulent serotypes (i.e. the monophasic *S. Typhimurium*) as well as of multidrug-resistant and invasive strains of NTS [9–11]. This has prompted the World Health Organization (WHO) to prioritize NTS serovars within the list of pathogens that could trigger future pandemics [12].

Most NTS human infections are associated with contaminated foods of animal or plant origin, resulting directly from infected animals or indirectly contaminated by them [13]. Specific serovars, such as *S. Typhimurium* and *S. Enteritidis*, have been identified as frequently associated with these animal/plant food sources [13]. Therefore, in numerous countries, initiatives have been implemented to reduce the prevalence of infection in food-producing animals, particularly poultry, which is considered the main source of *S. Enteritidis*, the top NTS infecting humans [14]. The control of *Salmonella* in poultry, largely through targeted interventions against it (e.g., vaccination) and the implementation of comprehensive farm-to-fork strategies, has been effective in reducing the incidence of human salmonellosis by this serotype in the EU [15]. However, an increasing trend in human cases of *S. Typhimurium* and its monophasic variant has been observed in recent years, which seems to be associated with pigs [16, 17].

Recent genomic studies indicate that the development of intensive swine production in the EU and the US over the last century, together with the globalization of trade and transportation from these regions, have been pivotal in the global emergence and spread of pig-related *Salmonella* [18]. The modern swine industry has undergone a period of gradual evolution, characterized by a series of incremental improvements in productivity and the steady consolidation of smaller farms into much larger herds, increasing pig population densities and, thus, the potential for some pathogen transmission [19]. Similarly, improved travel and transportation have enabled a significant expansion in the export and import of live pigs (e.g., in 2022 the EU imported and exported 29,054,656 and 34,954,421 live pigs, respectively; 20), contributing to the dissemination of pathogens across different world regions.

It is now widely accepted that pigs and their products represent a significant source of NTS infections in humans, with serotypes such as *S. Typhimurium*, its monophasic variant (*S. 1,4,[5],12:i:-*), *S. Derby*, and *S. Rissen* being particularly problematic [13, 16]. In the EU, the proportion of human salmonellosis cases attributable to

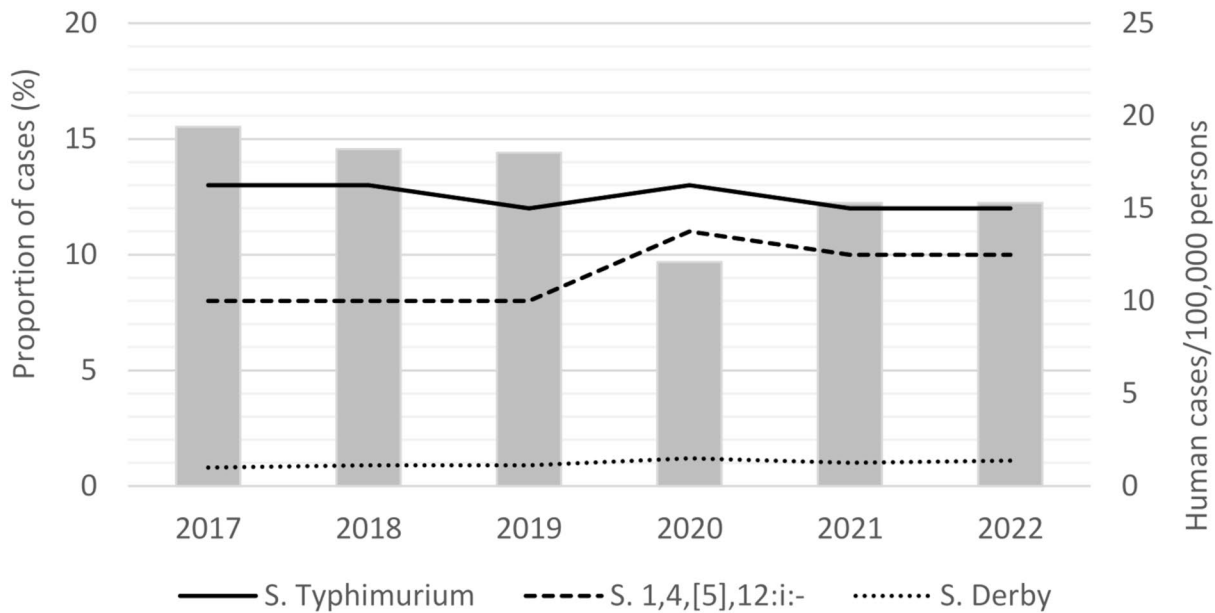
pigs, including pork and pork products, was estimated to be approximately 30%, although there were notable regional variations between Southern (43.6%) and Northern EU member states (10.6%) [21, 22]. In other developed countries such as the US, up to 12.1% of the *Salmonella* outbreaks were attributed to pork [23] and, in some regions of Australia, the proportion of cases attributed to pork increased from 20% in the period 2009–2016 to 40% in 2017–2019. This increase was likely associated with the emergence of the monophasic variant of *S. Typhimurium* [24].

In general, the proportion of cases of human salmonellosis involving swine-associated serotypes (*S. Typhimurium*, *S. 1,4,[5],12:i:-*, and *S. Derby*) has remained fairly constant in the EU during the last years (Fig. 1), suggesting that the efforts carried out to limit the spread of *Salmonella* from pigs to humans have not been successful. It is therefore apparent that new strategies need to be implemented to limit the global spread of NTS from pigs to humans, particularly from areas with a significant pig industry such as the EU. In this article, we discuss possible reasons for the lack of success of most national control programs (NCPs) within the EU and propose new approaches to this problem.

Control programs against pig salmonellosis in the EU

Following the successful implementation of NCPs targeting salmonellosis in fowls (laying hens, broilers, and turkeys) across all EU Member States (MS) in 2005, according to the framework set by the European Food Safety Authority (EFSA) and EU Regulation No. 2160/2003 [15], attention turned to pigs. It was hypothesized that a reduction in the prevalence of *Salmonella*-infected pigs would result in a decrease in the incidence of human cases associated with pork.

To determine the need for initiating these NCPs, a series of EU-wide baseline surveys were first conducted to assess the prevalence of *Salmonella* in pigs at various stages of production, from breeding to slaughter [25, 26]. These surveys provided crucial insights into the extent of *Salmonella* prevalence in breeding (average of 28.7%) and production (33.3%) holdings in the EU, as well as the prevalence of infection in pigs at slaughter (10.3%) and carcass contamination (8.3%). The results also revealed considerable variability in *Salmonella* prevalence among the MS, ranging from 0 to 64% in breeding holdings and from 0 to 29% in slaughtered pigs. The data also highlighted the dominance of specific serovars, such as *Salmonella Typhimurium* and *Salmonella Derby*, both commonly associated with human infections. These baseline figures should have served as essential data for setting reduction targets and assessing the effectiveness of control programs aimed at curbing *Salmonella* transmission in pig populations within the EU.



*After 2019 UK was not considered; in 2020 COVID19 pandemic occurred.

Data obtained from EFSA&ECDC reports, 2019, 2020, 2021, 2022 and 2023 (<https://efsa.onlinelibrary.wiley.com/>).

Fig. 1 Proportion of major swine-associated serotypes infecting humans in the EU during the period 2017–2022.*

Despite the general high levels of *Salmonella* prevalence in pig herds, and the fact that this species was considered the second most important source of human salmonellosis, no further action was taken at farm level in the EU. A comprehensive cost-benefit analysis suggested that there would be no positive economic benefit from setting targets to reduce *Salmonella* in slaughter pigs [27]. Therefore, the decision of whether or not to implement a NCP for salmonellosis in pigs was left to the discretion of each MS, but only a limited number of them decided to do so. These countries were either major producers of swine or demonstrated a high level of commitment to the control of this infection in food-producing animals.

The first comprehensive *Salmonella* NCP (encompassing the entire production chain) for pigs in Europe was already established in Sweden in the 1960s [28]. This was followed by the implementation of comprehensive NCPs in Norway, Finland, and Denmark in 1995 [29]. All of them, except the Danish program, focused on eradication and had bacteriological testing as the cornerstone of their programs. Denmark focused its NCP on *Salmonella* control and relied on both bacteriological and serological analyses [30]. In Norway and Finland the prevalence at the farm level was initially relatively low and strict

measures were enforced when *Salmonella* was detected. However, in Denmark, presenting a much larger pig population and higher *Salmonella* prevalence, actions at farm level were less restrictive [31].

Following the success of the Scandinavian action plans, and in line with the EU regulation, new NCPs followed suit in other MS: Germany and the United Kingdom in 2002, Ireland in 2003, the Netherlands in 2005, and Belgium in 2007 (Table 1). In general, these programs were modeled on the Danish approach, that is, focusing on control but using only serology for the surveillance program, which was carried out on a relatively small number of pigs per slaughter batch (from 12 to 72 per year, depending on the country) [30]. Thus, a weighted mean seroprevalence was calculated based on the most recent serological samplings on the farm, and the herds were subsequently classified into three distinct risk groups: low-risk (I), medium-risk (II), and high-risk (III) herds. Category III herds were required to implement specific on-farm measures aimed at reducing their exposure to *Salmonella* and, consequently, their *Salmonella* seroprevalence. While penalties were not typically imposed in most countries, in some cases some potential incentives (i.e. obtaining pork quality labels) were offered to farmers through certification programs. Examples of

Table 1 Simplified scheme of EU countries that have implemented National control programs against *Salmonella* in pigs and an overview of their results

Country (start year)	Key elements	Results	Success factors / Challenges
Sweden (1960) [28]	<ul style="list-style-type: none"> - Mandatory - Eradication - Emphasis on herd health and biosecurity - Regular bacteriological testing of all farms and slaughtered pigs - Special guarantees for imported meat 	Pig herd prevalence < 1% (based on lymph nodes and carcass bacteriology); stringent control measures, including vaccination, have led to almost negligible rates of <i>Salmonella</i>	Strong biosecurity measures and effective control at slaughter
Norway (1995) [38, 39]	<ul style="list-style-type: none"> - Mandatory - Eradication - Extensive bacteriological sampling of breeding herds and slaughtered pigs - Focus on low <i>Salmonella</i> prevalence through strict biosecurity 	Pig herd prevalence ≈ 0.03% (based on lymph nodes and carcass bacteriology)	Extremely low prevalence compared to other EU countries
Finland (1960) [40, 41]	<ul style="list-style-type: none"> - Mandatory - Eradication - Biosecurity, bacteriological surveillance (breeding herds and slaughtered pigs) - Special guarantees for imported meat 	Low prevalence at slaughter < 1% (based on lymph nodes and carcass bacteriology) due to extensive biosecurity practices and strict monitoring in pig farms	Exceptional biosecurity, low prevalence in imports
Denmark (1995) [42–44]	<ul style="list-style-type: none"> - Mandatory - Surveillance and control - Regular bacteriological and serological testing of whole pig production chain - Risk categorization - Strong biosecurity measures - Surveillance at slaughterhouses 	2–3% prevalence of <i>Salmonella</i> in breeding and multiplier pigs (based on serology) and 9–15% prevalence in slaughter pigs (based on serology) Low prevalence < 1% in carcass swabs (based on bacteriology)	High compliance with hygiene measures and transparent reporting systems Success but at high cost
Germany (2002) [31, 45]	<ul style="list-style-type: none"> - Mandatory - Surveillance and control - Serological testing of fattening and slaughtered pigs - Biosecurity and vaccination in high-risk areas 	Decreasing trend in <i>Salmonella</i> prevalence in fattening pigs (based on serology)	Stringent controls in farms and slaughterhouses Minimum success after 20 years
United Kingdom* (2002) [46, 47]	<ul style="list-style-type: none"> - Mandatory for QA** abattoirs - Surveillance and control - Serological testing of slaughtered pigs 	No significant improvement in seroprevalence [48]	Discontinued in 2012
Ireland (2003) (SI No. 165/2002 Abattoirs Act 1966; Veterinary Examination Amendment Regulations; SI No. 521/2009 y 522/2009)	<ul style="list-style-type: none"> - Mandatory - Surveillance and control - Monitoring via on-farm bacteriological sampling (finishers) and serological testing of slaughtered pigs - Biosecurity measures - Focus on high-risk farms 	<ul style="list-style-type: none"> - Decreasing trend in carcass <i>Salmonella</i> prevalence (probably due to hygiene practices at slaughter). - Seroprevalence of slaughter pigs has remained stable 	Ongoing focus on reducing contamination
The Netherlands (2005) [49]	<ul style="list-style-type: none"> - Mandatory - Surveillance and control - Serological testing of fattening and slaughtered pigs - Biosecurity improvements 	Current information on the <i>Salmonella</i> status of Dutch herds not available (QA-organizations do not publish results)	Difficulty eliminating persistent strains

Table 1 (continued)

Country (start year)	Key elements	Results	Success factors / Challenges
Belgium (2007) [50–52]	- Voluntary (since 2015) - Maintained for certain quality labels (e.g. Bepork) - Surveillance and control - Serological testing of fattening pigs	No significant improvement in seroprevalence has been observed	Facing ongoing challenges in collaboration with farmers and food industries. Difficulty of ensuring widespread compliance Variations in the effectiveness of biosecurity practices Differences in farm management
Estonia (2014) [35, 36]	- Mandatory for high-risk pig herds - Surveillance and control - Targeted serological and bacteriological sampling	Pig prevalence remains high (> 20%) at the farm level and around 3% at the slaughterhouse (based on bacteriology) [53]	Better farm practices and hygiene, though still facing challenges

*Pre-Brexit; ** Quality Assurance

these certification programs include Qualität und Sicherheit (QS) in Germany [32], British Quality Assured Pork (BQAP) in the United Kingdom [33], Bord Bia Quality Assurance Scheme in Ireland [34], and IKB Nederland varkens in the Netherlands [35].

Estonia was the last MS to implement a mandatory NCP in 2013, but with a different approach. The Estonian NCP is based on the detection of specific serotypes, namely, *S. Enteritidis*, *S. Typhimurium*, *S. Hadar*, *S. Infantis*, *S. Virchow*, *S. Choleraesuis*, *S. Derby*, and *S. Newport*, in pig farms, slaughterhouses, and processing plants [36, 37]. One-fifth of the farms are tested each year, including those that have tested positive for *Salmonella* in previous years and new farms that have not been previously tested. Restrictions are imposed if a farm tests positive.

The Danish NCP came to an end in January 2025. This decision was based on three key factors: the prevalence of *Salmonella* in farms has remained stable in recent years, the prevalence in pork carcasses has remained around 1%, and the number of people becoming ill from consuming Danish pork has also remained low. As a result, the responsibility for ensuring a low prevalence of *Salmonella* in Danish pork will lie solely with the slaughterhouses. (https://svineproduktion.dk/aktuelt/nyheder/2024/10/041024_Salmonella_ophoer). However, the approach taken by most of those countries that had opted for the control of the infection based on serological results from finishing pigs did not yield the expected results. Neither a significant decline in the prevalence of salmonellosis in pigs nor a reduction in human cases related to pork consumption was achieved. Consequently, most programs were discontinued [30].

What is wrong with the EU national control programs?

Among the NCP for pig salmonellosis initiated by European MS, only a few can be considered successful, often at considerable economic costs. In some cases, such as in the United Kingdom (UK) and Belgium, the programs have been discontinued or reduced to voluntary biosecurity guidance. Various reasons may explain this lack of success, including each country’s specifics—pig census, farm types, production systems and climate. Program implementation also varied, some being voluntary, others compulsory, and enforcement penalties were inconsistent [54]. Countries with strong enforcement, that is, actions to clean up contaminated farms, and penalties, had the most successful *Salmonella* programs.

Where programs were not mandatory nor penalties considered, the farmer’s perception of the problem certainly played an important part [55]. Effective *Salmonella* control requires changes in daily practices, dependent on farmers’ motivation, which in turn may depend on receiving tangible benefits from their actions [56, 57].

However, since porcine salmonellosis is often asymptomatic, direct benefits are unclear, reducing farmer interest.

Technical issues also contributed to the failure of the programs and the frustration of farmers. These issues likely relate to the structure of the pig sector and its impact on the epidemiology of the infection. In addition, the sampling methods employed and the use of imperfect serological tests (see below) to monitor *Salmonella* infection have undoubtedly contributed to the lack of success of these NCP. Understanding the importance of these factors is crucial for developing new and effective control strategies for swine salmonellosis in the future.

The structure of the pig sector and the epidemiology of the infection

The pig production system is complex, involving various production periods that may occur on different farms. For the sake of simplicity, it typically begins with sow farms, encompassing gestation, farrowing, and lactation. Following weaning, usually at three to four weeks, piglets move to nurseries until nine to ten weeks. Subsequently, the animals undergo a period of growth and fattening before being sent to slaughter at approximately 20 to 22 weeks of age. Some farms cover the entire production cycle (farrow-to-finish), while others focus on specific stages, such as breeding and nursery, nursery care alone, nursery and fattening (isowean units), or fattening alone. In some systems, batch production is also considered to prevent the spread of diseases.

In recent years, EU pig production has consolidated into large-scale farms [58]. Over 75% of pigs are raised on large commercial farms, that is, over 2,000 production (fattening) pigs [59, 60], with Denmark having the largest average herd size (4,700) and Germany the smallest (1,900). A special case is Spain, with a predominately intensive pig sector. It experienced a two-thirds drop in holdings (128,000) from 1999 to 2013 while the number of pigs per holding quadrupled. From 2014 to 2023, Spain has been the primary contributor to the growth in the EU pig census [60].

One consequence of the intensification process is the emergence of greater specialization among farming operations. An increasing number of farms have prioritized the construction of large fattening units typically situated in locations distant from the breeding-only farms from which their pigs originate. This practice is intended to enhance biosecurity [20]. Control efforts for pig salmonellosis have focused mostly on these fattening units, largely because this is the period immediately preceding slaughter. Therefore, the majority of epidemiological studies on the infection have been conducted during this period [61–69].

However, outcomes have been unsatisfactory due to *Salmonella*'s resilience and its interaction with farm- and

animal-related factors such as the type of infrastructure, farm external biosecurity, farm hygiene and disinfection, animal origin, animal management and associated stress, or concomitant infections with other enteric pathogens such as *Lawsonia intracellularis*. Risk factors associated with pig salmonellosis have been extensively described in the literature [70, 71]. The direct consequence of combining a highly versatile pathogen with the multitude of risk factors present on the farm (and at varying levels over the year) is that the presence of *Salmonella* in a herd is typically unpredictable [72]. The occurrence of *Salmonella* varies between and within age groups and pens within herds [63], and its presence is often inconsistent across batches [73].

Salmonella infections in fattening pigs may also originate from previous production phases, notably breeding sows, lactating piglets and the nursery, which have been the subject of comparatively little research [74–76]. A 2008 EFSA survey indicated that about 30% of sow holdings were *Salmonella*-positive, with higher prevalences in major pig-producing countries like Spain, the Netherlands, and Denmark [26]. The results of various serological and bacteriological surveys also indicate that the detection of high levels of *Salmonella* in sows is a common occurrence [77–80]. Infected sows and subsequent early infections occurring between birth and weaning are likely to play a pivotal role in the transmission and maintenance of *Salmonella*, increasing the overall likelihood of exposure to the bacteria in further production phases [81, 82].

There is a paucity of comprehensive studies on the prevalence of *Salmonella* in suckling piglets [76], mainly due to challenges in assessing the true infection status. Most research relies on fecal samples, finding low levels of shedding [63, 83–89]. Maternal immunity may influence these results [63, 89–91]. The limited amount of fecal matter typically obtained from rectal swabs, particularly in very young piglets, and the low sensitivity of bacteriology when performed on this matrix [92–94] may also have contributed to the underestimation of the true prevalence of infection in these animals. The prevailing view that weaning- or post-weaning-age pigs would be among the most clinically affected if they had become infected by *Salmonella* [22] has reinforced the perception that the prevalence of *Salmonella* at these ages is very low. In addition, until recently, antibiotics used to control other enteric pathogens in piglets (e.g., *E. coli*) likely masked *Salmonella* detection.

Recent studies on weaning piglets found a 36% *Salmonella* prevalence, with serotypes matching those in sows, suggesting that infected sows are the likely source for exposure and subsequent infection in piglets, which in turn would impact later production stages [78, 95]. It is therefore probable that infected sows are ultimately

responsible for infections occurring in the fattening units. Indeed, a risk assessment model indicated sow prevalence as a strong indicator of slaughter pig prevalence [96].

Salmonella transmission from sow to piglet mainly occurs via fecal contamination, but other pathways might exist. Evidence of congenital transmission of NTS from dairy cows to newborn calves was proposed in 2016 [97], and it could be happening in swine as well. If proven, new questions will be raised. For instance, whether persistently infected piglets may develop. This type of transmission, if confirmed, would have profound implications for the way this infection should be controlled on pig farms. Regardless of congenital transmission, minimizing sow-piglet transmission is crucial. Strategies to prevent *Salmonella* shedding in sows should be prioritized to control pig salmonellosis in the pre-harvest period.

Sampling procedures and the use of imperfect tests

In Europe, many commercial pig farms exceed 2,000 pigs [59, 60]. To accurately assess herd health, sampling must consider herd size. For a herd of 2,000 pigs, a perfect diagnostic test would require sampling between 70 (with 1% expected sero/prevalence) and 321 pigs (with 50% expected sero/prevalence) for a 95% confidence interval and 5% error (Win Episcopi; <http://www.winepi.net/index.php>). A typical fattening farm may market 4,000 to 6,000 pigs annually (between two and three fattening cycles per year), depending on its performance and production systems. Consequently, the number of animals sampled in NCPs often fails to yield precise health estimates.

Additionally, sampled animals (the study population) must represent the entire fattening unit (target population), as a strict all-in/all-out strategy is expected to be followed by those farms engaged in a *Salmonella* control program. Therefore, random selection is crucial to ensure equal probabilities of selection. Some NCPs exhibit significant bias in animal selection, particularly when using carcasses from slaughterhouses, which may not accurately reflect the distribution of pigs on the farm as they have been previously mixed during transportation and lairage. Given the evidence suggesting that *Salmonella* infection distribution in the herd is clustered, with the potential presence of herd subpopulations [98], results obtained from carcasses are likely giving a biased estimate of the health status of the herd. The sampling timing is also critical, as sampling at the beginning of the growing/fattening period may not reflect further changes in sero/prevalence. In contrast, sampling close to slaughter time can provide insights into the potential risk these pigs may pose for slaughter and carcass contamination.

Serological testing is the standard for monitoring *Salmonella* infection [54]. Indirect ELISA tests are quick and

cost-effective [99, 100] and can be performed on serum or meat juice samples, targeting major *Salmonella* serotypes affecting pigs [101, 102]. However, current serological tests for detecting *Salmonella*-specific antibodies are far from perfect, and their overall diagnostic accuracy is low. Studies show that serological testing often detects only a small fraction (15%) of *Salmonella*-shedding pigs [103], with sensitivities ranging from 59 to 65% depending on the ELISA used [102]. More recent studies employing Bayesian approaches reported ELISA sensitivities as low as 45% [104]. In general, the sensitivity of these tests can only be enhanced at the expense of a notable reduction in their specificities [105, 106].

Discrepancies between bacteriological and serological tests arise from various factors, including the timing of infection, serovar diversity [69, 107–109], or even the possibility of seropositive pigs becoming seronegative [63, 110, 111]. A further issue is the overall lack of agreement between serological tests at the individual level. This inconsistency appears to depend on the specific cut-off point recommended for each test [102, 105, 112, 113].

While some researchers find satisfactory herd-level agreement between serology and bacteriology [102], which may be useful for gaining insight into the circulation of *Salmonella* within a farm at a given moment, using serological tests to classify herds could misrepresent their true health status. It has been shown that when categorizing herds on different serological tests, significant discrepancies are observed [106, 113].

Focusing solely on the growing-finishing period for *Salmonella* control may be ineffective when the objective is to reduce the overall prevalence of *Salmonella* and the subsequent infection in humans attributed to pigs. The source of the infection may be in earlier stages of production that are not included in NCPs. This problem may be even more important in countries with intensive pig industries, where fattening units are separate from sow and nursery units. The limited and potentially unrepresentative sampling in many NCPs, coupled with the use of imperfect serological tests, results in biased health status estimates, leading to farmer skepticism and undermining effective on-farm *Salmonella* control measures.

New approaches for the control of pig-derived salmonellosis

The limited effectiveness and high costs of NCPs for pig salmonellosis, especially in countries with large pig populations [114, 115], highlight the need for reassessment. In the EU, pig salmonellosis is primarily a public health issue rather than an animal health concern. Thus, the main objective of any NCP should be to reduce human salmonellosis incidences linked to pork consumption. However, if consumers and public health authorities benefit from effective control programs while costs fall

solely on pig producers without compensation, voluntary farmer participation will likely be challenging. As is the case with other NCPs targeting zoonotic diseases (e.g., brucellosis, tuberculosis), penalties and/or compensation mechanisms should be considered, but given the current high prevalence of infection, costs in major pig-producing countries could be substantial.

Nevertheless, it is vital for farmers to understand the public health implications of salmonellosis in their pigs and their potential role in reducing it before engaging them in on-farm control activities. Once this awareness is established, modifying practices and implementing new ones will become easier [57]. However, given the infection's widespread nature, it may take time to observe positive outcomes from these efforts. In the meantime, new strategies are necessary to reduce human salmonellosis more rapidly. They should probably be implemented at the interface between the farm and the slaughterhouse.

People become ill after consuming contaminated pork, which often originates from pig carcasses contaminated during slaughter. The primary risk to humans arises from slaughtering pigs with high *Salmonella* concentrations in their feces [116]. It has been shown that a correlation exists between high cecal *Salmonella* loads in pigs and carcass contamination [117]. Asymptomatic *Salmonella*-infected pigs arriving at slaughterhouses are thus the main source of carcass contamination [118–120]. These pigs are particularly prone to *Salmonella* shedding due to pre-slaughter stressors like feed withdrawal, transport, and lairage [121–126]. Consequently, strategies should be developed to prevent or minimize the contamination of slaughterhouses by *Salmonella* from these animals. Such strategies may prove to be a more cost-effective short-term solution than attempting to control the infection on farms. Preferably, these strategies should be applicable regardless of the farmers' willingness to modify their production practices.

The farm-slaughterhouse interface

The relationship between herd *Salmonella* status and pig carcass contamination is well documented [25, 127, 128], but complex. The presence of *Salmonella* at the slaughterhouse may depend on several factors, such as recent pig seroconversion, pre-slaughter stressors, and lairage contamination [129–134], or the presence of *Salmonella* in the mesenteric lymph nodes of infected pigs [135]. Moreover, the implementation of proper slaughter procedures can significantly reduce the risk of contamination [128]. However, preventing carcass contamination cannot rely solely on slaughter activities, especially when *Salmonella* infection prevalence is high [128].

The only baseline study conducted in the EU indicated that, on average, 10% of slaughter pigs are infected, with some countries allowing up to a third of infected pigs into

their slaughterhouses [25]. It is expected that a significant number of pigs will shed *Salmonella* while at lairage. A recent study in Spain found that 27.3% of slaughter pigs were shedding *Salmonella* [136]. Lairage contamination may result in new infections and an increase in environmental contamination as this area cannot be adequately cleaned and disinfected during the day [137]. From there, the contamination can spread to the slaughter room, clean room, and even into the chillers, via vectors, fomites, or airborne transmission [138].

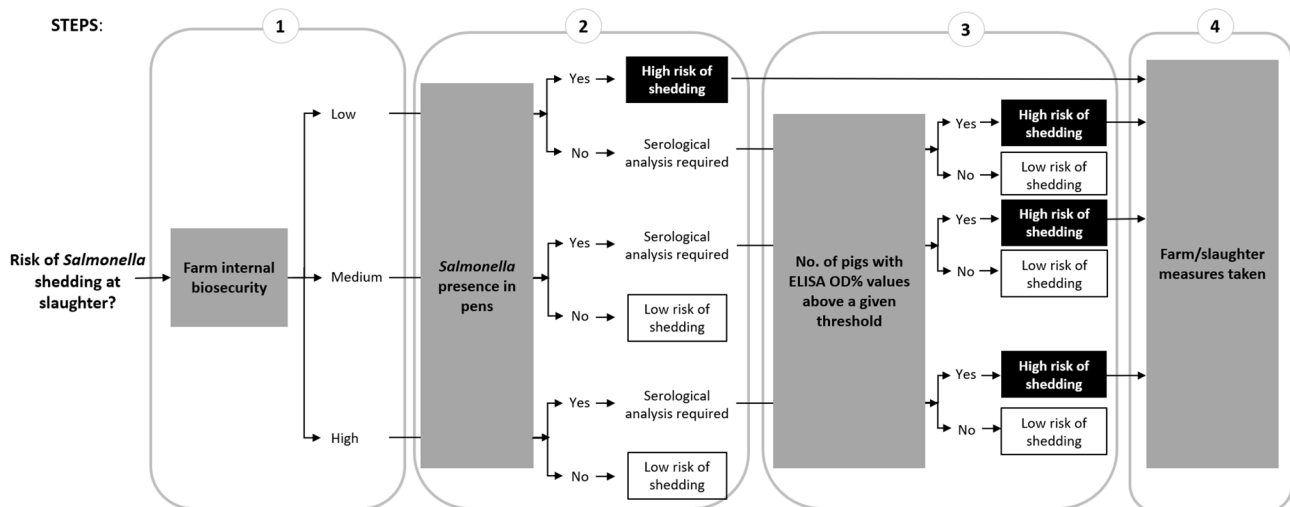
Although serology does not show a strong correlation with on-farm *Salmonella* shedding, this relationship becomes more evident when comparing serological results with shedding at the slaughterhouse [69, 135]. At the farm-slaughter interface, on-farm serology, along with two farmer-independent variables—the farm's internal biosecurity score and the prevalence of *Salmonella* in pens—has proven reliable for predicting batches of pigs with a high likelihood of shedding *Salmonella* during lairage [139]. Interestingly, serological data were not always necessary in the model by Bernad-Roche et al. Pigs from farms with low internal biosecurity and *Salmonella* in the pens were considered at high risk of shedding, regardless of serological results. Conversely, pigs from farms with good biosecurity and no detected *Salmonella* should be viewed as low risk, simplifying the approach. Consequently, a decision tree for reducing shedding risk at the slaughterhouse could be developed based on this model (Fig. 2). If a significant number of pigs from a specific batch are predicted to shed *Salmonella* prior to slaughter, both on-farm and on-slaughterhouse preventive measures can be identified to reduce the risk of slaughter contamination.

On-farm measures

In the final 2–3 weeks of fattening, measures for high-risk batches should aim at reducing *Salmonella* shedding both on the farm and at the slaughterhouse. Lowering shedding on the farm will reduce the likelihood of new infections, thus reducing contamination during transport and lairage. It will also lower infection pressure at the slaughterhouse.

At this stage, when only a few weeks or days remain before slaughter, the interventions must be capable of rapidly eliminating bacteria while remaining effective until slaughter and safe for consumers (no withdrawal period should be required). It can be reasonably assumed that the ingestion of products designed to reduce the burden of *Salmonella* in the intestinal tract is the only on-farm intervention that can be expected at this stage.

The use of coarse meal in pig feed has been associated with a decrease in *Salmonella* prevalence when compared to diets comprising finely ground or pelleted feed. The addition of coarse meal increases gastric retention



First, the internal biosecurity of the farm is determined. This could be derived for standardized questionnaires, such as those at <https://biocheckgent.com/en>. Then farms can be classified as of low, medium or high internal biosecurity according to percentiles (i.e. obtained from values from biocheckgent website for a given country) (Step 1). Analyze 10 pools of fecal samples from representative pens in the fattening unit 3-4 weeks before the time of slaughter (Step 2). The presence of *Salmonella* with low internal biosecurity will suggest high risk of shedding at slaughterhouse. On the contrary, the absence of *Salmonella* in farms with medium-to-high internal biosecurity will suggest low risk of shedding *Salmonella* at slaughter. For those farms for which a decision has not been made at step 2, a representative number of pigs from the fattening unit (50 were used for developing the model) are analyzed serologically (Step 3). The proportion of pigs above a given threshold (determined by the model) is calculated and a decision is taken regarding the risk of shedding for that batch (in the model proposed by Bernad-Roche et al, 2023, a probability of shedding >26% according to the model implied a >70% chance of being a shedder). On those batches of pigs considered of risk further on-farm or at slaughter actions may be taken (Step 4).

Fig. 2 Decision tree to assess the risk of shedding *Salmonella* at the slaughterhouse for a given fattening unit based on three variables, the farm internal biosecurity score, the presence of *Salmonella* in 10 representative pens and the serological results of 50 representative fattening pigs

time and encourages fermentation, leading to higher levels of organic acids (e.g., lactic acid), lower pH, and a richer anaerobic microbiota. These gut improvements have been shown to significantly reduce *Salmonella* survival during gastric transit and correlate with lower *Salmonella* shedding at slaughter, suggesting coarse meal could serve as a potential strategy for integrated *Salmonella* control [140].

However, given that pigs are typically subjected to a fasting period of 12 to 24 h prior to slaughter, a critical period for bacterial shedding, drinking water should probably be also considered a primary vehicle for most products. Water will be available to the animals until they are loaded onto the truck and, subsequently, in the lairage area. Prebiotics, probiotics, different types of organic acids, and more recently, postbiotics, parabiotics, and bacteriophages, have shown promising antibacterial properties and could be administrated through water (see below).

Prebiotics and probiotics

Prebiotics and probiotics can enhance animal health by influencing gut microbiota by stimulating beneficial bacteria (prebiotics) or directly increasing their populations (probiotics) [141]. This leads to a healthier digestive tract and immune system via mechanisms such as competitive exclusion and immune modulation, as well as the

production of useful metabolites, enzymes, or bacteriocins [142].

The efficacy of probiotics is contingent upon the interaction between the host (e.g. age, health, immune status) and the probiotic microorganism (e.g. strain, single or combination of several strains, dose) [143]. The impact of probiotics on *Salmonella* control has been primarily observed in piglets (suckling or postweaning), where gut flora is still in development [144–146]. In finishing pigs, where the microbiota is more stable [147], the treatment period may need to be much longer. Moreover, not all probiotics have proven effective against *Salmonella* [148], and safety remains a concern [143]. Strain selection must be carefully studied, with each probiotic requiring thorough safety and risk assessments [149]. If drinking water is used for administration, factors such as chlorination should be considered to prevent the reduction of probiotic doses.

Prebiotics should also require long periods of treatment to observe a positive effect. Some studies on piglets have shown reductions in *Enterobacteriaceae* and *Salmonella* counts after lengthy treatment periods (≈ 4 weeks) [150, 151]. In older pigs, there is only one study showing that the administration of prebiotics (a β -galacto mannan oligosaccharide) resulted in a reduction of *Salmonella* shedding and infection at the time of slaughter, but its effect was observed after two months of treatment [152].

Therefore, it remains uncertain whether probiotics and prebiotics could prove beneficial in reducing shedding during the final days of fattening. Further studies are needed to assess whether they can be used to reduce *Salmonella* shedding prior to slaughter in such a short time.

Postbiotics and parabiotics

The fields of postbiotics and parabiotics (PP) represent a novel area of research within the disciplines of animal nutrition, preventive veterinary medicine, and production [153]. Postbiotics are defined as the metabolic products secreted by probiotics, including enzymes, proteins, and peptides. In contrast, parabiotics are inactivated microbial cells containing components such as peptidoglycans, teichoic acids, surface proteins, or crude cell extracts [154]. PP are regarded as a safer alternative to probiotics, as they do not pose the same biological risks, such as bacterial translocation from the gut lumen to the bloodstream or the transfer of antibiotic resistance [155].

The available evidence suggests that PP can enhance animal performance and reduce *Enterobacteriaceae* counts and diarrhea [156]. However, most of trials have been conducted on young piglets and have lasted four to five weeks, which limits the applicability of these findings to pigs near slaughter. Despite the growing body of evidence supporting the health benefits of PP, further research is needed to elucidate their mechanisms of action, develop *Salmonella*-targeted PP, and establish international definitions for their regulation [157].

Organic acids

The addition of organic acids (OA) in feed or drinking water represents one of the most extensively researched strategies for the control of swine salmonellosis not only because of its direct antimicrobial activity on potential pathogens present in feed or water, but also because of its effects on the gastrointestinal tract of the animals. The bactericidal action of these acids is due to their ability to cross the cell membrane, dissociate inside where the pH is more alkaline, and acidify the cell cytoplasm, affecting protein and DNA synthesis and causing cell death [158]. In addition to this effect, the mechanisms of action of these acids in the gastrointestinal tract are numerous. Firstly, they lower the pH, mainly in the anterior sections of the gastrointestinal tract, as acids are normally absorbed along the small intestine, thus reinforcing the stomach as an entry barrier for *Salmonella*. Secondly, they stimulate the growth of epithelial cells [159]. It has been shown that certain acids, mainly butyric, caproic and caprylic acids, can also reduce the expression of *Salmonella* pathogenicity genes, thus limiting their capacity to colonize the intestinal epithelium of pigs [160, 161]. Nevertheless, some studies have suggested that some

short-chain fatty acids may also induce the expression of invasion genes in *Salmonella* [162, 163].

While the early use of OA faced several challenges, such as the corrosive effects on watering pipes, poor palatability, and difficulty reaching the posterior sections of the gastrointestinal tract (ileum, caecum, colon), where *Salmonella* typically colonizes, these issues have been effectively overcome through the microencapsulation of OA [159, 164].

Nevertheless, the efficacy of OA for the control of *Salmonella* in pigs has been variable, contingent upon factors such as the different study designs (e.g., piglets vs. fattening pigs, natural vs. experimental infection, different administration periods), the type of OA used or the dose applied [165–170]. In addition, a potential adverse effect is the development of acid resistance, which would reduce the efficacy of these agents [171].

In general, beneficial results have been observed after the administration of OA for at least four weeks [167, 172–175], and it appears to be a cost-effective measure to reduce *Salmonella* prevalence along the pork production chain [176, 177]. However, the need for prolonged treatment periods raises doubts about their suitability when applied to finishing pigs prior to slaughter.

A particularly interesting type of OA is that which has undergone esterification. They are short- and medium-chain fatty acids combined with glycerol and have shown enhanced antimicrobial activity against Gram-negative bacteria in both in vitro and in vivo settings [178]. Esterified OA have shown additional advantages, including reduced pH dependence and enzymatic breakdown susceptibility, which allows for activity across the entire gastrointestinal tract [179]. Furthermore, they possess an amphipathic structure that allows them to be soluble in water without altering the pH. Additionally, they are odorless and non-corrosive, which prevents any off-flavors in the water that might deter the pigs from drinking it.

A recent study using an esterified form of formic acid showed that the inclusion of 10 kg/1000L of this acid into the farm water supply for five days prior to slaughter effectively reduced the shedding of *Salmonella* by 82% [180]. The treatment also resulted in a significant reduction in the *Salmonella* loads in pigs that continued to shed the bacteria. The same dosage was observed to reduce the proportion of shedders by up to 63% when the treatment was applied exclusively in the drinking water of the lairage area [136]. These results indicate that this esterified form of formic acid may be a promising product within an overall strategy to minimize *Salmonella* shedding at slaughter.

In general, OA, and particularly those that could be easily blended with drinking water, appear to be a feasible strategy to reduce the shedding of slaughter pigs. The use

of OA is safe and could be administered even during the stay of the pigs in the lairage area, thereby increasing the likelihood of timely elimination of the bacteria from the pigs' gut.

Bacteriophages

The use of bacteriophages (phages) offers a promising strategy for the reduction of *Salmonella* loads within the intestinal tract of finishing pigs. Phages exclusively infect bacteria, thus they are not harmful to animal cells or consumers [181]. They can lyse multidrug-resistant (MDR) strains [182] and their effects are observed quickly [183, 184]. Both in-feed and in-water delivery methods are effective as phages multiply in the gastrointestinal tract while bacteria are present [185]. While a minimum bacterial presence is needed for phage propagation [186], this condition is likely to be met in most of the target pigs.

Salmonella phages are abundant in pig slurry [187] and can be easily obtained and selected for use [188]. Each year, new phages are being characterized for potential use in commercial farms [189–191]. However, there are several biological and technical obstacles that must be overcome before phages can be employed to treat *Salmonella* in finishing pigs. First, only virulent (non-lysogenic) phages should be selected to ensure bacterial elimination [192]. Phages are also highly specific, often targeting only a particular species, serotype, or subset of strains [193]. Therefore, it is of paramount importance to select the appropriate phage for the target *Salmonella* serotype. Knowledge of the most prevalent serotypes on the farm is needed, but focusing on zoonotic serotypes like Typhimurium, its monophasic variant, and Derby may prove effective. In addition, the use of phage cocktails can help broaden the host range [193].

Phages are susceptible to a range of external factors, including temperature, acidity, salinity, and ions [194]. Therefore, delivery methods must ensure phage survival. Water or feed are anticipated as vehicles for the on-farm administration of phages, and factors such as water composition, chlorination, feed pelleting, and stomach acidity can reduce their survival [182]. Solutions to this challenge include the use of phages in buffer solutions, encapsulation, and freeze- or spray-drying [182, 195, 196]. Ensuring phage stability remains a key challenge for the industry [197].

A potential risk associated with the use of phages is the emergence of phage-resistant *Salmonella* variants, which typically occurs through spontaneous mutations [198]. Experimental studies show that resistance can develop within hours after exposure to a single phage, primarily through mutations in lipopolysaccharide (LPS) biosynthetic genes [199, 200]. This can be also mitigated by the use of phage cocktails [201–203]. Interestingly, the development of phage resistance can render bacteria

more susceptible to environmental factors and antibiotics [200].

The use of phages has significantly reduced *Salmonella* colonization and shedding in post-weaned pigs [183–185, 204, 205]. It is expected that similar results would be observed in older animals, thus making them a suitable intervention for the treatment of pigs at high risk of shedding *Salmonella* at the time of slaughter. Although a comprehensive regulatory framework for phage therapy in veterinary medicine is still lacking in most countries, progress has been made. Phages are not yet authorized in the EU, but the European Medicines Agency (EMA) has recently issued guidelines on the quality, safety, and efficacy of veterinary medicinal products for phage therapy [206]. These guidelines provide clear regulatory, technical, and scientific requirements for phage-based veterinary medicines.

Additional on-farm interventions

While the implementation of these strategies may assist in reducing the shedding of *Salmonella* at the slaughterhouse and subsequent contamination of the pig carcasses, the original sources of *Salmonella* infection will remain unaddressed, contributing to the sustained high *Salmonella* prevalence in many farms. However, the routine sampling of a representative number of pens and animals on the farm and the identification of batches of high risk of *Salmonella* shedding would provide accurate information to properly identify risk farms. This information could be used to prompt further investigations into the sources of infection in these farms and the implementation of additional, more general, on-farm interventions.

Most activities on pig farms should focus on preventing new *Salmonella* infections. This objective can be accomplished through three fundamental interventions: the sanitary control of feed and breeding animals, as well as the enhancement of biosecurity measures.

Feed contamination is recognized as a significant route for the introduction of *Salmonella* into pig farms. It has been estimated to account for up to 14.2% of the total infection risk [207]. All Member States (MS) runs a rather equal risk because the high risk feed ingredients (vegetable protein, in particular soybeans- and rapeseed meal) are equally used in all MS, and when studied found to be frequently contaminated. The risk may be higher in countries using animal derived proteins. The relative importance of the risk of introducing *Salmonella* by feed into farms is higher in low prevalence countries where other sources largely are minimized. Those countries generally also apply special measures to minimize the risk for introducing *Salmonella* by contaminated feed. In the absence of such measures, contaminated feed will jeopardize efforts to improve the *Salmonella* status also in high prevalence countries [208–210].

Swine producer should ensure that feed mills are implementing Hazard Analysis and Critical Control Points (HACCP)-based program according to EU Regulation (183/2005 EC), which in detail presents the elements involved, and the role of the competent authority and the feed operator. Control measures include heat treatment (e.g., pelleting at high temperatures), the use of organic acids or other antimicrobial additives, protocols for hygiene in feed production facilities, and regular microbiological monitoring of both raw materials and finished products and, most important, that corrective interventions are undertaken when *Salmonella* is isolated. Furthermore, the sourcing of certified *Salmonella*-free ingredients is emphasized [42]. Studies have shown that when implemented effectively, HACCP systems can be successful in reducing and even eliminating *Salmonella* contamination in feed and feed ingredients, consequently leading to a decline in the number of infected farms [209, 211]. The production of *Salmonella* safe feed requires that all feed business operators have implement the EU regulation and can document that prescribed procedures are maintained. However, the efficacy of this mandate largely depends on enforcement at the national level, and compliance may vary among feed producers, particularly in regions with limited official control. Consequently, farmers may rely on feed mills that choose to implement rigorous HACCP-based programs [212].

Effective feed management on pig farms is also crucial to prevent *Salmonella* recontamination, particularly when feed is mixed or processed on-site. Even when employing ingredients that have been certified as safe, inadequate on-farm practices, including improper storage, a lack of pest control, or insufficient cleaning of mixers and silos, can jeopardize the safety of the feed. Farms that produce their own feed should use heat-treated components and refrain from long storage periods. Sealed storage, rodent control, equipment hygiene, and strict biosecurity during handling are all key to reducing the risk of infection [211].

Breeding sows also play an important role in the transmission of *Salmonella* to piglets. Therefore, in breeding farms, the efforts should be focused on minimizing the shedding of *Salmonella* in sows. As with finishing pigs, the routine administration to the sows' feed or drinking water of some organic acids or phages with a rapid bacterial killing effect during the days prior to farrowing could be an effective strategy to decrease *Salmonella* contamination in maternal crates.

Salmonella vaccination has been an optimal strategy for the eradication of *Salmonella* in laying hens and to prevent egg contamination [213]. The same approach could be used in swine to limit animal, environmental, and food contamination [79]. Sow vaccination prior to farrowing to boost passive immunity of the suckling piglets [214],

which could be combined with piglet vaccination during the nursery period to induce its own immune response to *Salmonella*, would help to reduce *Salmonella* infection and shedding in this initial period of the pig production [215].

Nonetheless, many factors, such as the intracellular mode of *Salmonella* infection, its antigenic diversity and the lack of vaccine cross protection, the high prevalence of infection even from the early life of the animals, and the persistence of the bacteria in the environment, challenge the efficacy of vaccination [215]. A meta-analysis study showed that, irrespective of the type of vaccine used (attenuated or inactivated), this strategy was effective in reducing the number of *Salmonella*-positive samples on the farm, therefore contributing to the reduction of within-farm transmission, but its overall efficacy was limited (<30%) [216]. Vaccination should therefore be considered as another intervention that could be implemented along with other on-farm measures, but not to rely solely on it. In addition, depending on the vaccine strategy used (e.g. only vaccination of sows, vaccination of piglets or growers), vaccines should be neither used in farms undergoing routine serological monitoring because of their interference with test results [215].

Biosecurity, and particularly internal biosecurity, has been positively associated with reduced *Salmonella* shedding at slaughter [139]. Internal biosecurity is described as those measures intended to prevent the within-herd spread of pathogens as opposed to external biosecurity, which includes measures to avoid the introduction of pathogens from outside the farm [217, 218]. Thus, implementing and maintaining activities such as disease containment, strict hygiene protocols, and proper use of working lines and manure handling, should contribute to a reduction of *Salmonella* circulation within the farm. However, internal and external biosecurity are highly correlated and should be treated as a common strategy if effectiveness is to be maximized [219].

Cleaning and disinfection (C&D) are critical activities to prevent bacteria from remaining in the facilities and the subsequent infection of new animals entering them [220]. *Salmonella* is known for its ability to persist in the environment [221], which is enhanced by the production of biofilms. Consequently, many cleaning and disinfection (C&D) protocols may prove inadequate for eradicating the bacteria [222]. Phages may serve as a supplementary measure alongside C&D protocols for farm facilities, given their capacity to produce lytic compounds and enzymes that disrupt biofilms [223, 224]. The use of autophages, that is, phages isolated from the same farm where the target bacterium has been isolated [225] appears to be a promising strategy to eradicate recalcitrant *Salmonella* strains in farms [226].

Interventions at the slaughterhouse

While on-farm *Salmonella* infection represents a significant source of slaughterhouse environmental contamination, the potential for *Salmonella* contamination exists at any stage of pork production. The ultimate carcass status depends upon the slaughtering conditions [118], thereby underscoring the pivotal role of the slaughterhouse.

Slaughters are subject to considerable pressure due to the high prevalence of *Salmonella*-infected pigs received. Consequently, rigorous measures must be implemented along the slaughter line to prevent contamination at any stage. The available evidence indicates that the failure to implement effective control measures at critical points in the slaughter process, such as scalding, dehairing, singeing, evisceration, and facility cleaning, is associated with an increased risk of carcass contamination. Reviews of the efficacy of different slaughter interventions are available elsewhere [227, 228]. Despite the efforts of many slaughterhouses, contamination of pig carcasses persists [6, 118].

One of the primary sources of *Salmonella* contamination in the slaughterhouse is the lairage area. During lairage, stressed infected pigs are more likely to shed *Salmonella*, thereby facilitating its transmission to other pigs within hours [129]. Implementing measures such as reducing lairage time and cleaning between batches may prove an effective means of mitigating the risk of carcass contamination [128, 129, 137]. However, the practical feasibility of such measures may present a significant challenge. Logistic slaughter of *Salmonella*-positive batches is another proposed intervention, but its efficacy is still a matter of debate [228]. The effectiveness of logistic slaughter could be enhanced by segregating high-risk pig batches before delivery [137]. Furthermore, holding these pigs separately and providing water treated with esterified organic acids may reduce *Salmonella* shedding before slaughter [136].

Implementing more straightforward slaughter interventions to approach a zero policy for *Salmonella*-positive carcasses is also a possibility. Methods include physical and chemical decontamination treatments on carcasses, such as chlorine, electrolyzed oxidizing water, and organic acids, which have proven effective [229–232]. However, these can result in side effects such as the proliferation of acid-resistant bacteria [233]. In addition, chlorine-based disinfectants pose safety concerns [234] and they are not widely accepted by European consumers [235]. Physical techniques such as irradiation or pulsed-light UV have also been shown to significantly reduce bacterial counts, but they raise consumer concerns [236, 237]. Bacteriophage cocktails also offer a promising avenue for decontamination, though more research is needed before they can be recommended for widespread use [238]. Furthermore, although these methods

are regarded as cost-effective for reducing microbiological contamination on pig carcasses [239–241], their impact should not be overstated. Overreliance may lead to a relaxation of hygiene practices [233]. Therefore, they should be part of a comprehensive food safety system [242]. In light of these concerns, the EU has adopted a cautious stance, permitting only water for carcass decontamination (Regulation (EC) No 853/2004). In contrast, the US allows the use of alternative treatments, including organic acids [243].

Until new interventions are approved, slaughterhouses must continue implementing strict measures along the slaughter line in order to effectively manage the high number of infected pigs that they receive. Accurate knowledge of the *Salmonella* status of incoming pigs would greatly improve the efficiency of these control measures, particularly at the lairage stage, helping to reduce environmental contamination during slaughter.

Future directions and conclusions

Previous attempts to reduce human salmonellosis in the EU through NCPs targeting pig salmonellosis have largely been unsuccessful, likely due to the difficulty in lowering *Salmonella* infection rates on pig farms. The focus of these programs has primarily been on fattening pigs and slaughterhouses, while earlier stages in the pig production chain, where the infection is also prevalent, have been neglected.

The role of infected sows in the transmission of *Salmonella* to piglets has been the subject of little research. A recent study suggests *Salmonella* could be transmitted congenitally in mammals such as cattle. However, no studies have been conducted on swine in this regard, an animal species that exhibits higher infection rates than cattle. Furthermore, it was previously assumed that infected suckling piglets would become sick, but recent evidence shows they can appear healthy while shedding *Salmonella*, thus spreading it to the nursery. This issue is further aggravated by the ban on antimicrobials like colistin [244], an antibiotic previously used to control gram-negative bacteria. These findings underscore the need for effective salmonellosis control measures to be initiated at the sow farm.

The rapid growth of the pig industry, with more large-scale farms and concentrated production, must be considered when designing effective monitoring programs. Sampling procedures should better reflect the herd's true *Salmonella* status to foster trust among farmers. Without it, the implementation of new on-farm biosecurity measures will be challenging. The diagnostic accuracy of serological tests is constrained by the nature of this infection, rendering results difficult to interpret. Furthermore, vaccination is only effective in conjunction with strict biosecurity measures.

If there is a real commitment to reducing human salmonellosis linked to pork, there are different strategies that could be used at farm level. It may also be possible to predict the risk of *Salmonella* shedding before pigs arrive at the slaughterhouse. Batches identified as high-risk could be treated on-farm with in-water additives in the days preceding slaughter, in order to reduce shedding and environmental contamination during processing. This approach is likely to be economically feasible, provided that the public health benefit of reducing human salmonellosis is properly recognized. At present, OA represent the most effective short-term treatment option, but alternative methods, such as bacteriophages, may soon emerge. Furthermore, routine on-farm sampling of a representative number of finishing pigs and pens could assist in a more accurate classification of farms according to their *Salmonella* risk, thereby enabling slaughterhouses to optimize slaughter procedures and targeted interventions.

Abbreviations

C&D	Cleaning and disinfection
DNA	Deoxyribonucleic acid
ECDC	European centre for disease prevention and control
EFSA	European food safety Authority
ELISA	Enzyme-linked immunosorbent assay
EMA	European medicines agency
EU	European union
QA	Quality assurance
QS	Qualität und sicherheit
LPS	Lipopolysaccharide
MS	Member state
NCPs	National control programs
NTS	Non-typhoidal salmonella
OA	Organic acids
PP	Postbiotics and parabiotics
UK	United Kingdom
US	United States
WHO	World health organization

Acknowledgements

Not applicable.

Author contributions

RCM-J: Conceptualization, Supervision, Contribution to writing original draft preparation, Writing—reviewing and editing. AC-H: Contribution to writing original draft preparation, Writing—reviewing and editing. MB-R: Contribution to writing original draft preparation. JPV: Contribution to writing original draft preparation. SA-B: Contribution to writing original draft preparation.

Funding

this research did not receive any specific grant from funding agencies in the public, commercial, or not-for-profit sectors.

Data availability

No datasets were generated or analysed during the current study.

Declarations

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare no competing interests.

Received: 2 June 2025 / Accepted: 22 August 2025

Published online: 21 October 2025

References

1. Key FM, Posth C, Esquivel-Gomez LR, Hübner R, Spyrou MA, Neumann GU, et al. Emergence of human-adapted *Salmonella* enterica is linked to the neolithization process. *Nat Ecol Evol*. 2020;4(3):324–33. <https://doi.org/10.1038/s41559-020-1106-9>
2. Als D, Radhakrishnan A, Arora P, Gaffey MF, Campisi S, Velummailum R, et al. Global trends in typhoidal salmonellosis: A systematic review. *Am J Trop Med Hyg*. 2018;99(3Suppl):10–9. <https://doi.org/10.4269/ajtmh.18-0034>
3. European Food Safety Authority (EFSA), European Centre for Disease Prevention and Control (ECDC). The European union one health 2023 zoonoses report. *EFSA J*. 2024;22:e9106. <https://doi.org/10.2903/j.efsa.2024.9106>
4. European Food Safety Authority (EFSA). 2014. Fact sheet *Salmonella*. Available online: https://www.efsa.europa.eu/sites/default/files/corporate_publication/files/factsheetSalmonella.pdf Accessed 3 October 24.
5. Országh E, Pitter J, Kaló Z, Vokó Z, Jóźwiak Á. Retrospective cost-utility analysis of the Non-typhoidal *Salmonella* control programme in Hungary. *Food Control*. 2021;120:107529. <https://doi.org/10.1016/j.foodcont.2020.107529>
6. European Food Safety Authority (EFSA), European Centers for Disease Control (ECDC). The European union one health 2022 zoonoses report. *EFSA J*. 2023;21:e8442. <https://doi.org/10.2903/j.efsa.2023.8442>
7. Non-Typhoidal GBD *Salmonella*, Collaborators ID. 2019. The global burden of non-typhoidal *Salmonella* invasive disease: a systematic analysis for the Global Burden of Disease Study 2017. *Lancet Infect Dis*. 2017;19(12):1312–1324. [https://doi.org/10.1016/S1473-3099\(19\)30418-9](https://doi.org/10.1016/S1473-3099(19)30418-9)
8. Teklemariam A, Al-Hindi R, Albiheyri R, Alharbi M, Alghamdi M, Filimban AM, et al. Human salmonellosis: A continuous global threat in the Farm-to-Fork food safety continuum. *Foods*. 2023;12. <https://doi.org/10.3390/foods12091756>
9. Glynn MK, Bopp C, Dewitt W, Dabney P, Mokhtar M, Angulo FJ. Emergence of multidrug-resistant *Salmonella* enterica serotype typhimurium DT104 infections in the United States. *N Engl J Med*. 1998;338(19):1333–8. <https://doi.org/10.1056/NEJM199805073381901>
10. Feasey NA, Dougan G, Kingsley RA, Heyderman RS, Gordon MA. Invasive non-typhoidal *Salmonella* disease: an emerging and neglected tropical disease in Africa. *Lancet*. 2012;379(9835):2489–99. [https://doi.org/10.1016/S0140-6736\(11\)61752-2](https://doi.org/10.1016/S0140-6736(11)61752-2)
11. Lund S, Tahir M, Vohra LI, Hamdana AH, Ahmad S. Outbreak of monophasic *Salmonella* typhimurium sequence type 34 linked to chocolate products. *Ann Med Surg*. 2022;82:104597. <https://doi.org/10.1016/j.jamsu.2022.104597>
12. World Health Organization (WHO). Pathogens prioritization. A scientific framework for epidemic and pandemic research preparedness. *Health Emergence Program* 2024;72.
13. Ferrari R, Rosario D, Cunha-Neto A, Mano S, Figueiredo E, Conte-Junior C. Worldwide epidemiology of *Salmonella* serovars in Animal-Based foods: a Meta-analysis. *Appl Environ Microbiol*. 2019;85(14):e00591–19. <https://doi.org/10.1128/AEM.00591-19>
14. Li Q, Wang X, Yin K, Hu Y, Xu H, Xie X, et al. Genetic analysis and CRISPR typing of *Salmonella* enterica serovar enteritidis from different sources revealed potential transmission from poultry and pig to human. *Int J Food Microbiol*. 2018;266:119–25. <https://doi.org/10.1016/j.jfoodmicro.2017.11.025>
15. Cota JB, Langkabel N, Barco L, Olsen A, Bonardi S, Vieira-Pinto M, et al. Comparison of European surveillance and control programs for *Salmonella* in broiler and Turkey chains. *Food Control*. 2024;165:110656. <https://doi.org/10.1016/j.foodcont.2024.110656>
16. Campos J, Mourão J, Peixe L, Antunes P. Non-typhoidal *Salmonella* in the pig production chain: A comprehensive analysis of its impact on human health. *Pathogens*. 2019;8. <https://doi.org/10.3390/pathogens8010019>
17. Merlotti A, Manfreda G, Munck N, Hald T, Litrup E, Nielsen E, et al. Network approach to source attribution of *Salmonella* enterica serovar typhimurium and its monophasic variant. *Front Microbiol*. 2020;11. <https://doi.org/10.3389/fmicb.2020.01205>
18. Li H, Wu Y, Feng D, Jiang Q, Li S, Rong J, et al. Centralized industrialization of pork in Europe and America contributes to the global spread of *Salmonella*

- enterica. *Nat Food*. 2024;5(5):413–22. <https://doi.org/10.1038/s43016-024-00968-8>
19. VanderWaal K, Deen J. Global trends in infectious diseases of swine. *PNAS*. 2018;115(45):11495–500. <https://doi.org/10.1073/pnas.1806068115>
 20. Zimmerman J. In furtherance of Dr. Tom Alexander's legacy of innovation in swine health: Innovation in surveillance. Proceedings of the 27th International Pig Veterinary Society Congress & 15th European Symposium of Porcine Health Management 2024. June 4th–7th. Leipzig, Germany, pp. XV–XVIII.
 21. Pires SM, Vieira AR, Hald T, Cole D. Source attribution of human salmonellosis: an overview of methods and estimates. *Foodborne Pathog Dis*. 2024;11(9):667–76. <https://doi.org/10.1089/fpd.2014.1744>
 22. Bonardi S. *Salmonella* in the pork production chain and its impact on human health in the European union. *Epidemiol Infect*. 2017;145(8):1513–26. <https://doi.org/10.1017/S095026881700036X>
 23. Interagency Food Safety Analytics Collaboration. Foodborne illness source attribution estimates for 2021 for *Salmonella*, *Escherichia coli* O157, and *Listeria monocytogenes* using multi-year outbreak surveillance data, United States. GA and D.C.: U.S. Department of Health and Human Services, Centers for Disease Control and Prevention, Food and Drug Administration, U.S. Department of Agriculture's Food Safety and Inspection Service 2023.
 24. McLure A, Shadbolt C, Desmarchelier PM, Kirk MD, Glass K. Source attribution of salmonellosis by time and geography in new South Wales, Australia. *BMC Infect Dis*. 2022;22(1):14. <https://doi.org/10.1186/s12879-021-06950-7>
 25. European Food Safety Authority (EFSA). Report of the task force on zoonoses data collection on the analysis of the baseline survey on the prevalence of *Salmonella* in slaughter pigs. Part A. *EFSA J*. 2018;135:1–111. <https://doi.org/10.2903/j.efsa.2009.1377>
 26. European Food Safety Authority (EFSA). Analysis of the baseline survey on the prevalence of *Salmonella* in holdings with breeding pigs in the EU, 2008-Part A: *Salmonella* prevalence estimates. *EFSA J*. 2009;7(12):1377. <https://doi.org/10.2903/j.efsa.2009.1377>
 27. FCC Consortium. Analysis of the Costs and Benefits of Setting a Target for the Reduction of *Salmonella* in Breeding Pigs for European Commission Health and Consumers Directorate-General Final Report SANCO/2008/E2/056. 2010. https://food.ec.europa.eu/system/files/2016-10/biosafety_food-borne-disease_Salmonella_fattening-pigs_slaughterhouse-analysis-costs.pdf Accessed 12 December 2024.
 28. Swedish Board of Agriculture. Regulations on control of *Salmonella* in animals (in Swedish). *SJVFS* 2004;2 Saknr K 102.
 29. Hopp P, Wahlström H, Hirn J. A common *Salmonella* control programme in Finland, Norway and Sweden. *Acta Vet Scand Suppl*. 1999;91:45–9.
 30. Correia-Gomes C, Leonard F, Graham D. Description of control programmes for *Salmonella* in pigs in Europe. *Progress Date?? J Food Saf* 2021;41(5):e12916.
 31. European Food Safety Authority (EFSA). Opinion of the scientific panel on biological hazards (BIOHAZ) related to risk assessment and mitigation options of *Salmonella* in pig production. *EFSA J*. 2006;341:1–131. <https://doi.org/10.2903/j.efsa.2006.341>
 32. QS (Qualität und Sicherheit). 2024. Guideline *Salmonella* Monitoring Pigs. https://www.q-s.de/services/files/downloadcenter/h-salmonellenmonitoring/2024/leitfaden/englisch/Guideline_Salmonella_Monitoring_Pigs_01.01.2024.pdf. Accessed 12 December 2024.
 33. British Meat Processors Association (BMPA). BMPA Pork Scheme. 2024. <https://britishmeatindustry.org/our-work/bmpa-pork-scheme/> Accessed 5 November 24.
 34. Bord Bia Irish Food Board. Pigmeat Quality Assurance Scheme (PQAS). 2024. <https://www.bordbia.ie/farmers-growers/get-involved/become-quality-assured/pigmeat-quality-assurance-scheme-pqas/> Accessed 5 November 24.
 35. IKB Nederland. IKB Nederland varkens. 2024. <https://www.ikbnederland.nl/varkens/> Accessed 5 November 24.
 36. STE. Regulation No. 39 Salmonellooside tõrje eeskiri for the control/eradication of salmonellosis of Minister of Agriculture of Estonia (STE), Riigi Teataja (RT) I, 24.05.2013. 2021. Available online: <https://www.riigiteataja.ee/akt/10412020050> Accessed 12 December 2024.
 37. VTA (Veterinaar-ja Toiduamet/Veterinary and Food Board of Estonia). Salmonelloosi kontrollprogramm aastateks 2020–2021 (Estonian Salmonellosis control program for years 2020–2021) 2019;1–26.
 38. Jore S, Lyngstad TM, Hofshagen M, Bergsjø B, Bruheim T, Falck M et al. The surveillance and control programmes for *Salmonella* in live animals, eggs and meat in Norway. En: Brun, E., Hellberg, H., Mørk, T. y Jordsmyr, H.M. (Coord.). Surveillance and control programmes for terrestrial and aquatic animals in Norway. Annual report. Oslo (Norway): National Veterinary Institute 2007;21–8.
 39. Heier BT, Norström M, Bergsjø B, Sæbø KS, Kalberg S, Linaker M, et al. The surveillance programme for *Salmonella* in live animals, eggs and meat in Norway 2014. Oslo (Noruega): Norwegian Veterinary Institute; 2015.
 40. Maijala R, Ranta J, Seuna E, Peltola J. The efficiency of the Finnish *Salmonella* control programme. *Food Control*. 2005;16(8):669–75. <https://doi.org/10.1016/j.foodcont.06003>
 41. Ministry of Agriculture and Forestry of Finland (MMMEEO). Zoonoses in Finland in 1995–1999 (in Finnish: Zoonootit Suomessa 1995–1997). Helsinki (Finlandia). 2000. https://www.ruokavirasto.fi/globalassets/teemat/zoonootit/eskus/zoonootit/zoonootit00_2.pdf. Accessed 12 December 2024.
 42. Hald T, Wingstrand A, Pires SM, Vieira A, Domingues AR, Lundsby K et al. Assessment of the human-health impact of *Salmonella* in animal feed. Copenhagen (Denmark): DTU Food. 2012. https://backend.orbit.dtu.dk/ws/portalfiles/portal/84060316/Report_Assessment_of_the_human_health_impact_of_Salmonella_in_animal_feed.pdf. Accessed 12 December 2024.
 43. Anonymous. Annual report on zoonoses in Denmark 2018. Kongens Lyngby (Denmark): National Food Institute, Technical University of Denmark. 2019.
 44. SEGES. Health status management. 2020. <http://spsus.dk/en>. accessed 12 December 2024.
 45. Merle R, Kösters S, May T, Ports U, Blaha T, Kreienbrock L. Serological *Salmonella* monitoring in German pig herds: results of the years 2003–2008. *Prev Vet Med*. 2011;99(2–4):229–33. <https://doi.org/10.1016/j.prevetmed.2011.02.007>
 46. Hill AA, Snary EL, Arnold ME, Alban L, Cook AJC. Dynamics of *Salmonella* transmission on a British pig grower-finisher farm: A stochastic model. *Epidemiol Infect*. 2008;136(3):320–33. <https://doi.org/10.1017/S0950268807008485>
 47. Snary EL, Munday DK, Arnold ME, Cook AJC. Zoonoses action plan *Salmonella* monitoring programme: an investigation of the sampling protocol. *J Food Prot*. 2012;73(3):488–94. <https://doi.org/10.4315/0362-028X-73.3.488>
 48. Simons R, Berriman A, Hill A, Gavin C, Smith R. A cost-benefit assessment of *Salmonella*-control strategies in pig herds within the United Kingdom. Proceedings of the 12th International Symposium on the Epidemiology and Control of Biological, Chemical and Physical Hazards in Pigs and Pork (SafePork), Foz do Iguaçu (Brasil), 2017;197–200. <https://doi.org/10.31274/safepork-180809-356>
 49. van der Wolf PJ. Monitoring for *Salmonella* in Swine in the Netherlands. 2017. https://www.pig333.com/articles/monitoring-for-salmonella-in-swine-in-the-netherlands_12866/ Accessed 12 December 2024.
 50. Méroc E, Strubbe M, Vangroenweghe F, Czaplicki G, Vermeersch K, Hooyberghs J, et al. Evaluation of the *Salmonella* surveillance program in Belgian pig farms. *Prev Vet Med*. 2012;105:309–14. <https://doi.org/10.1016/j.prevetmed.03006>
 51. Anonymous. Koninklijk besluit tot opheffing van het koninklijk en het ministerieel besluit van 27 april 2007 betreffende de bewaking van *Salmonella* bij varkens. 2015. https://etaamb.openjustice.be/nl/koninklijk-besluit-van-29-mei-2015_n2015018197.html. Accessed 12 December 2024.
 52. Peeters L. On-farm control measures against *Salmonella* Typhimurium infections in pigs: focus on vaccination with an attenuated vaccine and the application of a probiotic feed additive. Ghent University. Faculty of Veterinary Medicine, Merelbeke, Belgium. 2019. <http://hdl.handle.net/1854/LU-8639749>. Accessed 13 November 2024.
 53. Kuus K, Kramarenko T, Sögel J, Mäesaar M, Fredriksson-Ahomaa M, Roasto M. Prevalence and serotype diversity of *Salmonella* enterica in the Estonian meat production chain in 2016–2020. *Pathogens*. 2021;10(12):1622. <https://doi.org/10.3390/pathogens10121622>
 54. Carvajal A, Kramer M, Argüello H. *Salmonella* control in swine: A thoughtful discussion of the Pre- and Post-Harvest control approaches. *Industrialized Ctries Anim*. 2024;14(7):1035. <https://doi.org/10.3390/ani14071035>. PMID: 38612274; PMCID: PMC11010990.
 55. Houben MAM, Caekbeke N, van den Hoogen A, Ringenier M, Tobias TJ, Jonquiere FJ, et al. The ADKAR® change management model for farmer profiling with regard to antimicrobial stewardship in livestock production. *Vlaams Diergeneeskundig Tijdschrift*. 2020;89:309–14.
 56. Fraser RW, Williams NT, Powell LF, Cook AJ. Reducing *Campylobacter* and *Salmonella* infection: two studies of the economic cost and attitude to adoption of on-farm biosecurity measures. *Zoonoses Public Health*. 2010;57(7–8):e109–15. <https://doi.org/10.1111/j.1863-2378.2009.01295.x>
 57. Marier E, Piers R, Ellis-Iversen J, Watson E, Armstrong D, Hogeveen H, et al. Changes in perceptions and motivators that influence the implementation

- of on-farm *Salmonella* control measures by pig farmers in England. *Prev Vet Med.* 2016;133:22–30. <https://doi.org/10.1016/j.prevetmed.2016.09.009>
58. Bellini S. The pig sector in the European Union. In: Understanding and combatting African Swine Fever. A European perspective (Ed: L. Iacolina, M.-L. Penrith, S. Bellini, E. Chenais, F. Jori, M. Montoya, K. Ståhl and D. Gavner-Widén). Wageningen Academic Pub. 2021;183–95. <https://doi.org/10.3920/978-90-8686-910-7>
59. Augère-Granier ML. The EU pig meat sector. European Parliamentary Research Service -EPRS-, PE 652.044. 2020. [https://www.europarl.europa.eu/R egData/etudes/BRIE/2020/652044/EPRS_BRI\(2020\)652044_EN.pdf](https://www.europarl.europa.eu/R egData/etudes/BRIE/2020/652044/EPRS_BRI(2020)652044_EN.pdf). Accessed 24 May 2024.
60. Mateos GG, Corrales NL, Tategón G, Aguirre L. Pig meat production in the European Union-27: current status, challenges, and future trends. *Anim Biosci.* 2024;37(4):755–74. <https://doi.org/10.5713/ab.23.0496>
61. Davies PR, Morrow WE, Jones FT, Deen J, Fedorka-Cray PJ, Harris IT. Prevalence of *Salmonella* in finishing swine Raised in different production systems in North caroline, USA. *Epidemiol Infect.* 1997;119–237:244. <https://doi.org/10.1017/s095026889700784x>.
62. van der Wolf PJ, Bongers JH, Elbers AR, Franssen FM, Hunneman WA, van Exsel AC, et al. *Salmonella* infections in finishing pigs in the netherlands: bacteriological herd prevalence, serogroup and antibiotic resistance of isolates and risk factors for infection. *Vet Microbiol.* 1999;67(4):263–75. [https://doi.org/10.1016/s0378-1135\(99\)00054-1](https://doi.org/10.1016/s0378-1135(99)00054-1)
63. Kranker S, Alban L, Boes J, Dahl J. Longitudinal study of *Salmonella* enterica serotype typhimurium infection in three Danish farrow-to-finish swine herds. *J Clin Microbiol.* 2003;41:2282–8. <https://doi.org/10.1128/JCM.41.6.2282-2288.2003>
64. Rajic A, Keenlside J, McFall ME, Deckert AE, Muckle AC, O'Connor BP, et al. Longitudinal study of *Salmonella* species in 90 Alberta swine finishing farms. *Vet Microbiol.* 2005;105(1):47–56. <https://doi.org/10.1016/j.vetmic.2004.10.005>
65. Sanchez J, Dohoo I, Christensen J, Rajic A. Factors Influencing the prevalence of *Salmonella* spp. In swine farms: A metaanalysis approach. *Prev Vet Med.* 2077;81(1–3):148–77. <https://doi.org/10.1016/j.prevetmed.2007.04.005>
66. Farzan A, Friendship RM, Dewey CE, Muckle AC, Gray JT, Funk J. Distribution of *Salmonella* serovars and phage types on 80 Ontario swine farms in 2004. *Can J Vet Res.* 2008;72(1):1–6.
67. Vico JP, Rol I, Garrido V, San Román B, Grilló MJ, Mainar-Jaime RC. Salmonellosis in finishing pigs in Spain: prevalence, antimicrobial agent susceptibilities, and risk factor analysis. *J Food Prot.* 2011;74(7):1070–8. <https://doi.org/10.4315/0362-028X.JFP-10-515>
68. Pires AF, Funk JA, Bolin CA. Longitudinal study of *Salmonella* shedding in naturally infected finishing pigs. *Epidemiol Infect.* 2013;141(9):1928–36. <https://doi.org/10.1017/S0950268812002464>
69. Nair S, Farzan A, O'Sullivan TL, Friendship RM. Time course of *Salmonella* shedding and antibody response in naturally infected pigs during grower-finisher stage. *Can J Vet Res.* 2018;82(2):139–45.
70. Fosse J, Seegers H, Magras C. Prevalence and risk factors for bacterial food-borne zoonotic hazards in slaughter pigs: a review. *Zoonoses Public Health.* 2009;56(8):429–54. <https://doi.org/10.1111/j.1863-2378.2008.01185.x>
71. De Busser EV, De Zutter L, Dewulf J, Houf K, Maes D. *Salmonella* control in live pigs and at slaughter. *Vet J.* 2013;196(1):20–7. <https://doi.org/10.1016/j.tvjl.2013.01.002>
72. Eddington TS, Brown TR. A commentary on *Salmonella* from a Pre-Harvest perspective. *Front Anim Sci.* 2022;3:877392. <https://doi.org/10.3389/fanim.2022.877392>
73. Henry A, Letellier A, Côté JC, Desmarais G, Lachapelle V, Bergeron N, et al. *Salmonella* contamination in a network of 10 pig farms interconnected within the same cooperative. *Vet Rec Open.* 2019;6(1):e000269. <https://doi.org/10.1136/vetreco-2017-000269>
74. Nollet N, Houf K, Dewulf J, De Kruif A, De Zutter L, Maes D. *Salmonella* in sows: a longitudinal study in farrow-to-finish pig herds. *Vet Res.* 2005;36(4):645–56. <https://doi.org/10.1051/vetres:2005022>
75. Wilkins W, Rajić A, Waldner C, McFall M, Chow E, Muckle A, Rosengren L. Distribution of *Salmonella* serovars in breeding, nursery, and grow-to-finish pigs, and risk factors for shedding in ten farrow-to-finish swine farms in Alberta and Saskatchewan. *Can J Vet Res.* 2010;74(2):81–90. PMID: 20592836; PMCID: PMC2851729.
76. Wales AD, Cook AJ, Davies RH. Producing *Salmonella*-free pigs: a review focusing on interventions at weaning. *Vet Rec.* 2011;168(10):267–76. <https://doi.org/10.1136/vr.d1125>
77. Bessire BC, Thomas M, Gehring KB, Savell JW, Griffin DB, Taylor TM, et al. National survey of *Salmonella* prevalence in lymph nodes of sows and market hogs. *Transl Anim Sci.* 2018;2(4):365–71. <https://doi.org/10.1093/tas/txy072>
78. Casanova-Higes A, Marín-Alcalá CM, Andrés-Barranco S, Cebollada-Solanas A, Alvarez J, Mainar-Jaime RC. Weaned piglets: another factor to be considered for the control of *Salmonella* infection in breeding pig farms. *Vet Res.* 2019;50(1):45. <https://doi.org/10.1186/s13567-019-0666-7>
79. Bearson SMD. *Salmonella* in swine: prevalence, multidrug resistance, and vaccination strategies. *Annu Rev Anim Biosci.* 2022;10:373–93. <https://doi.org/10.1146/annurev-animal-013120-043304>
80. Hollmann I, Lingens JB, Wilke V, Homann C, Teich K, Buch J, et al. Epidemiological study on *Salmonella* prevalence in Sow herds using direct and indirect detection methods. *Microorganisms.* 2022;10(8):1532. <https://doi.org/10.3390/microorganisms10081532>
81. Nollet N, Houf K, Dewulf J, Duchateau L, De Zutter L, De Kruif A, et al. Distribution of *Salmonella* strains in farrow-to-finish pig herds: A longitudinal study. *J Food Prot.* 2005;68:2012–21.
82. Lurette A, Touzeau S, Lamboni M, Monod H. Sensitivity analysis to identify key parameters influencing *Salmonella* infection dynamics in a pig batch. *J Theor Biol.* 2009;258(1):43–52. <https://doi.org/10.1016/j.jtbi.2009.01.026>
83. Funk J, Davies P, Nichols M. Longitudinal study of *Salmonella* enterica in growing pigs reared in multiple-site swine production systems. *Vet Microbiol.* 2001;83:45–60. [https://doi.org/10.1016/s0378-1135\(01\)00404-7](https://doi.org/10.1016/s0378-1135(01)00404-7)
84. Barber DA, Bahnsen PB, Isaacson R, Jones CJ, Weigel RM. Distribution of *Salmonella* in swine production ecosystems. *J Food Prot.* 2022;65:1861–8. <https://doi.org/10.4315/0362-028X-65.12.1861>
85. Beloeil PA, Chauvin C, Proux K, Rose N, Queguiner S, Eveno E et al. Longitudinal serological responses to *Salmonella* enterica of growing pigs in a subclinically infected herd. *Prev Vet Med.* 2003;60:207–26. [https://doi.org/10.1016/s0167-5877\(03\)00126-0](https://doi.org/10.1016/s0167-5877(03)00126-0). PMID: 12900159.
86. Roesler U, Vonaltrock A, Heller P, Bremerich S, Arnold T, Lehmann J, et al. Effects of fluoroquinolone treatment acidified feed, and improved hygiene measures on the occurrence of *Salmonella* typhimurium DT104 in an integrated pig breeding herd. *J Vet Med B Infect Dis Vet Public Health.* 2005;52:69–74. <https://doi.org/10.1111/j.1439-0450.2005.00825.x>
87. Dors A, Czyżewska-Dors E, Wasyl D, Pomorska-Mól M. Prevalence and factors associated with the occurrence of bacterial enteropathogens in suckling piglets in farrow-to-finish herds. *Vet Rec.* 2016;179(23):598. <https://doi.org/10.1136/vr.103811>
88. Lynch H, Wallia K, Leonard FC, Lawlor PG, Manzanilla EG, Grant J, et al. *Salmonella* in breeding pigs: shedding pattern, transmission of infection and the role of environmental contamination in Irish commercial farrow-to-finish herds. *Zoonoses Public Health.* 2018;65:e196–206. <https://doi.org/10.1111/zph.12428>
89. Schut CH, Farzan A, Ainslie-Garcia MH, Friendship RM, Lillie B. Antibody responses to *Salmonella* in pigs from weaning up to marketing and presence of *Salmonella* at slaughter. *Foodborne Pathog Dis.* 2019;16(3):187–94. <https://doi.org/10.1089/fpd.2018.2454>
90. Rooke JA, Bland IM. The acquisition of passive immunity in the new-born piglet. *Livest Prod Sci.* 2002;78(1):13–23. [https://doi.org/10.1016/S0301-6226\(02\)00182-3](https://doi.org/10.1016/S0301-6226(02)00182-3)
91. Parada J, Carranza AI, Pichel M, Tamiozzo PJ, Pelliza BR, Ambrogi A. *Salmonella* transmission from the gilt to her offspring. *Livest Sci.* 2013;157:605–11. <https://doi.org/10.1016/j.livsci.2013.09.010>
92. Hurd HS, Stabel TJ, Carlson S. August. Sensitivity of various fecal sample collections techniques for detection of *Salmonella* Typhimurium in finish hogs. In: Proceeding of the third international symposium on the epidemiology and control of *Salmonella* in Pork, Washington DC, 1999.
93. Funk JA, Davies P, Nichols MA. The effect of fecal sample weight on detection of *Salmonella* enterica in swine feces. *J Vet Diagn Invest.* 2000;12:412–18. <https://doi.org/10.1177/104063870001200504>
94. Sangvatanakul P. Prevalence of *Salmonella* in piglets and in the fattening period in Chiang Mai, Thailand. Master Thesis, Veterinary Public Health, Chiang Mai University and Freie Universität Berlin. 2007. <https://cmudc.library.cmu.ac.th/frontend/info/item/dc:108396>. Accessed 19 November 24.
95. Bernad-Roche M, Casanova-Higes A, Marín-Alcalá CM, Cebollada-Solanas A, Mainar-Jaime RC. *Salmonella* Infection in Nursery Piglets and Its Role in the Spread of Salmonellosis to Further Production Periods. *Pathogens.* 2021;10(2):123. <https://doi.org/10.3390/pathogens10020123>
96. Martínez JM, McCarthy C, Taylor RA, Animal and Plant Health Agency (APHA), Biomathematics and Risk Research work group, United Kingdom. Livestock

- health and food chain risk assessment. EFSA J. 2020;18(S1):e181111–pp11. <https://doi.org/10.2903/j.efsa.2020.e181111>
97. Hanson DL, Loneragan GH, Brown TR, Nisbet DJ, Hume ME, Edrington TS. Evidence supporting vertical transmission of *Salmonella* in dairy cattle. Epidemiol Infect. 2016;144(5):962–7. <https://doi.org/10.1017/S0950268815002241>
98. Rao S, Kitron U, Weigel RM. Spatial and genotypic clustering of *Salmonella* over time in a swine production unit. Prev Vet Med. 2010;97:90–9. <https://doi.org/10.1016/j.prevetmed.2010.09.005>
99. Proux K, Cariolet R, Fravallo P, Houdayer C, Keranflech A, Madec F. Contamination of pigs by nose-to-nose contact or airborne transmission of *Salmonella* typhimurium. Vet Res. 2001;32(6):591–600. <https://doi.org/10.1051/vetres:2001148>
100. Bohaychuk VM, Gensler GE, King RK, Wu JT, McMullen LM. Evaluation of detection methods for screening meat and poultry products for the presence of foodborne pathogens. J Food Prot. 2005;68(12):2637–47. <https://doi.org/10.4315/0362-028X-68.12.2637>
101. Mousing J, Jensen PT, Halgaard C, Bager F, Feld N, Nielsen B, Nielsen JP, et al. Nation-wide *Salmonella* enterica surveillance and control in Danish slaughter swine herds. Prev Vet Med. 1997;29(4):247–61. [https://doi.org/10.1016/s0167-5877\(96\)01082-3](https://doi.org/10.1016/s0167-5877(96)01082-3)
102. Farzan A, Friendship RM, Dewey CE. Evaluation of enzyme-linked immunosorbent assay (ELISA) tests and culture for determining *Salmonella* status of a pig herd. Epidemiol Infect. 2007;135(2):238–44. <https://doi.org/10.1017/S0950268806006868>
103. Sibley J, Yue B, Huang F, Harding J, Kingdon J, Chirino-Trejo M, et al. Comparison of bacterial enriched-broth culture, enzyme linked immunosorbent assay, and broth culture-polymerase chain reaction techniques for identifying asymptomatic infections with *Salmonella* in swine. Can J Vet Res. 2003;67(3):219–24. PMID: 12889729; PMCID: PMC227056.
104. Arnold M, Smith RP, Martelli F, Davies R. Bayesian evaluation of meat juice ELISA for detecting *Salmonella* in slaughtered pigs without specifying a cut-off. Zoonoses Public Health. 2024;71(4):369–80. <https://doi.org/10.1111/zph.13109>
105. Mainar-Jaime RC, Atashparvar N, Chirino-Trejo M, Blasco JM. Accuracy of two commercial enzyme-linked immunosorbent assays for the detection of antibodies to *Salmonella* spp. In slaughter pigs from Canada. Prev Vet Med. 2008;85:41–51. <https://doi.org/10.1016/j.prevetmed.2007.12.015>
106. Vico JP, Engel B, Buist WG, Mainar-Jaime RC. Evaluation of three commercial enzyme-linked immunosorbent assays for the detection of antibodies against *Salmonella* spp. In meat juice from finishing pigs In Spain. Zoonoses Public Health. 2010;57:107–14. <https://doi.org/10.1111/j.1863-2378.2010.01364.x>
107. Nielsen B, Baggesen D, Bager F, Haugegaard J, Lind P. The serological response to *Salmonella* serovars typhimurium and infantis in experimentally infected pigs. The time course followed with an indirect anti-LPS ELISA and bacteriological examinations. Vet Microbiol. 1995;47:205–18. [https://doi.org/10.1016/0378-1135\(95\)00113-1](https://doi.org/10.1016/0378-1135(95)00113-1)
108. Jacob J, Rachel T, Shankar B, Gunasekaran K, Iyadurai R, Anandan S, et al. MLST based serotype prediction for the accurate identification of Non typhoidal *Salmonella* serovars. Mol Biol Rep. 2020. <https://doi.org/10.1007/s11033-020-05856-y>. 47,7797 – 803.
109. Nair S, Farzan A, Poljak Z, Friendship R. Identifying active *Salmonella* infections in swine nurseries using serology and bacterial culture and evaluating associated risk factors. Animals. 2020;27(9):1517. <https://doi.org/10.3390/ani10091517>
110. Dahl J, Wingstrand A, Nielsen B, Baggesen DL. Elimination of *Salmonella* typhimurium infection by the strategic movement of pigs. Vet Rec. 1997;140:679–81.
111. van der Wolf PJ, Fo L, Wong DM, Wolbers WB, Elbers AR, van der Heijden HM, et al. A longitudinal study of *Salmonella* enterica infections in high- and low-seroprevalence finishing swine herds in the Netherlands. Vet Q. 2001;23:116–21. <https://doi.org/10.1080/01652176.2001.9695096>
112. Mejia W, Casal J, Mateu E, Martín M. Comparison of two commercial ELISAs for the serological diagnosis of salmonellosis in pigs. Vet Rec. 2005;157(2):47–8. <https://doi.org/10.1136/vr.157.2.47>
113. Szabó I, Scherer K, Roesler U, Appel B, Nöckler K, Hensel A. Comparative examination and validation of ELISA test systems for *Salmonella* typhimurium diagnosis of slaughtering pigs. Int J Food Microbiol. 2008;124(1):65–9. <https://doi.org/10.1016/j.jfoodmicro.2008.02.022>
114. Anonymous FCC, Consortium F. Report. Analysis of the costs and benefits of setting a target for the reduction of *Salmonella* in breeding pigs for European Commission Health and Consumers Directorate-General SANCO/2008/E2/036. 2010. https://food.ec.europa.eu/system/files/2016-10/biosafety_food-borne-disease_Salmonella_fattening-pigs_slaught-house-analysis-costs.pdf. Accessed 14 June 2024.
115. Gavin C, Simons RRL, Berriman ADC, Moorhouse D, Snary EL, Smith RP, et al. A cost-benefit assessment of *Salmonella*-control strategies in pigs reared in the United Kingdom. Prev Vet Med. 2018;160:54–62. <https://doi.org/10.1016/j.prevetmed.2018.09.022>
116. Snary EL, Swart AN, Simons RR, Domingues AR, Vigre H, Evers EG, et al. A quantitative Microbiological risk assessment for *Salmonella* in pigs for the European union. Risk Anal. 2016;36(3):437–49. <https://doi.org/10.1111/risa.12586>
117. Pesciaroli M, Cucco L, De Luca S, Massacci FR, Maresca C, Medici L, et al. Association between pigs with high caecal *Salmonella* loads and carcass contamination. Int J Food Microbiol. 2017;242:82–6. <https://doi.org/10.1016/j.jfoodmicro.2016.11.021>
118. Argüello H, Alvarez-Ordoñez A, Carvajal A, Rubio P, Prieto M. Role of slaughtering in *Salmonella* spreading and control in pork production. J Food Prot. 2013;76:899–911. <https://doi.org/10.4315/0362-028X.JFP-12-404>
119. Swart AN, Evers EG, Simons RL, Swanenburg M. Modeling of *Salmonella* contamination in the pig slaughterhouse. Risk Anal. 2016;36:498–515. <https://doi.org/10.1111/risa.12514>
120. Marin C, Chinillac MC, Cerdà-Cuellar M, Montoro-Dasi L, Sevilla-Navarro S, Ayats T, Marco-Jimenez F, et al. Contamination of pig carcass with *Salmonella* enterica serovar typhimurium monophasic variant 1,4[5],12:i:- originates mainly in live animals. Sci Total Environ. 2020;10703:134609. <https://doi.org/10.1016/j.scitotenv.2019.134609>
121. Williams LP Jr, Newell KW. *Salmonella* excretion in joy-riding pigs. Am J Public Health. 1970;60:926–9. <https://doi.org/10.2105/AJPH.60.5.926>
122. Isaacson RE, Firkins LD, Weigel RM, Zuckerman FA, DiPietro JA. Effect of transportation and feed withdrawal on shedding of *Salmonella* typhimurium among experimentally infected pigs. Am J Vet Res. 1999;60:1155–8.
123. Marg H, Scholz HC, Arnold T, Rösler U, Hensel A. Influence of long-time transportation stress on re-activation of *Salmonella* typhimurium DT104 in experimentally infected pigs. Berl Munch Tierarztl Wochenschr. 2001;114:385–8.
124. Martín-Peláez S, Peralta B, Creus E, Dalmau A, Velarde A, Pérez JF. Different feed withdrawal times before slaughter influence caecal fermentation and faecal *Salmonella* shedding in pigs. Vet J. 2009;182:469–73. <https://doi.org/10.1016/j.tvjl.2008.08.002>
125. Eicher SD, Rostagno MH, Lay DC. Feed withdrawal and transportation effects on *Salmonella* enterica levels in market-weight pigs. J Anim Sci. 2017;95(7):2848–58. <https://doi.org/10.2527/jas.2017.1454>
126. Massacci FR, Morelli A, Cucco L, Castinel A, Ortenzi R, Tofani S, et al. Transport to the slaughterhouse affects the *Salmonella* shedding and modifies the fecal microbiota of finishing pigs. Animals. 2020;10(4):676. <https://doi.org/10.3390/ani10040676>
127. Beloeil PA, Chauvin C, Proux K, Madec F, Fravallo P, Alioum A. Impact of the *Salmonella* status of market-age pigs and the pre-slaughter process on *Salmonella* caecal contamination at slaughter. Vet Res. 2004;35(5):513–30. <https://doi.org/10.1051/vetres:2004028>
128. Alban L, Stärk KD. Where should the effort be put to reduce the *Salmonella* prevalence in the slaughtered swine carcass effectively? Prev Vet Med. 2005;68(1):63–79. <https://doi.org/10.1016/j.prevetmed.2005.01.001>
129. Hurd HS, McKean JD, Wesley IV, Karriker LA. The effect of lairage on *Salmonella* isolation from market swine. J Food Prot. 2001;64(7):939–44. <https://doi.org/10.4315/0362-028X-64.7.939>
130. Dorr PM, Tadesse DA, Zewde BM, Fry P, Thakur S, Gebreyes WA. Longitudinal study of *Salmonella* dispersion and the role of environmental contamination in commercial swine production systems. Appl Environ Microbiol. 2009;75(6):1478–86. <https://doi.org/10.1128/AEM.01632-08>
131. Hernández M, Gómez-Laguna J, Luque I, Herrera-León S, Maldonado A, Reguillo L, et al. *Salmonella* prevalence and characterization in a free-range pig processing plant: tracking in trucks, lairage, slaughter line and quartering. Int J Food Microbiol. 2013;162(1):48–54. <https://doi.org/10.1016/j.jfoodmicro.2012.12.026>
132. Simons RR, Hill AA, Swart A, Kelly L, Snary EL. A transport and lairage model for *Salmonella* transmission between pigs applicable to EU member States. Risk Anal. 2016;36(3):482–97. <https://doi.org/10.1111/risa.12390>
133. Buder C, Meemken D, Fürstenberg R, Langforth S, Kirse A, Langkabel N. Drinking pipes and nipple drinkers in pig abattoir lairage Pens-A source of zoonotic pathogens as a hazard to meat safety. Microorganisms. 2023;11(10):2554. <https://doi.org/10.3390/microorganisms11102554>

134. Dias Costa R, Silva V, Leite A, Saraiva M, Lopes TT, Themudo P et al. *Salmonella* spp., *Escherichia coli* and Enterobacteriaceae Control at a Pig Abattoir: Are We Missing Lairage Time Effect, Pig Skin, and Internal Carcass Surface Contamination? *Foods*. 2023;12(15):2910. <https://doi.org/10.3390/foods12152910>
135. Casanova-Higes A, Andrés-Barranco S, Mainar-Jaime RC. Influence of On-farm pig *Salmonella* status on *Salmonella* shedding at slaughter. *Zoonoses Public Health*. 2017;64(5):328–36. <https://doi.org/10.1111/zph.12301>
136. Bernad-Roche M, Casanova-Higes A, Marín-Alcalá CM, Mainar-Jaime RC. *Salmonella* Shedding in Slaughter Pigs and the Use of Esterified Formic Acid in the Drinking Water as a Potential Abattoir-Based Mitigation Measure. *Animals* 2022;12(13):1620. <https://doi.org/10.3390/ani12131620>
137. Berriman AD, Clancy D, Clough HE, Armstrong D, Christley RM. Effectiveness of simulated interventions in reducing the estimated prevalence of *Salmonella* in UK pig herds. *PLoS ONE*. 2013;8(6):e66054. <https://doi.org/10.1371/journal.pone.0066054>
138. Pearce RA, Sheridan JJ, Bolton DJ. Distribution of airborne microorganisms in commercial pork slaughter processes. *Int J Food Microbiol*. 2006;107(2):186–91. <https://doi.org/10.1016/j.jfoodmicro.2005.08.029>
139. Bernad-Roche M, Marín-Alcalá CM, Cebollada-Solanas A, de Blas I, Mainar-Jaime RC. Building a predictive model for assessing the risk of *Salmonella* shedding at slaughter in fattening pigs. *Front Microbiol*. 2023;14:1232490. <https://doi.org/10.3389/fmicb.2023.1232490>
140. Mikkelsen LL, Naughton PJ, Hedemann MS, Jensen BB. Effects of physical properties of feed on microbial ecology and survival of *Salmonella enterica* serovar typhimurium in the pig Gastrointestinal tract. *Appl Environ Microbiol*. 2004;70(6):3485–92. <https://doi.org/10.1128/AEM.70.6.3485-3492.2004>
141. Berge AC, Wierup M. Nutritional strategies to combat *Salmonella* in monogastric food animal production. *Animal*. 2012;6(4):557–64. <https://doi.org/10.1017/S1751731111002217>
142. Liao SF, Nyachoti M. Using probiotics to improve swine gut health and nutrient utilization. *Anim Nutr*. 2017;3(4):331–43. <https://doi.org/10.1016/j.aninu.2017.06.007>
143. Bajagai YS, Klieve AV, Dart PJ, Bryden WL. Probiotics in animal nutrition – Production, impact and regulation by. Editor Harinder P.S. Makkar. *FAO Animal Production and Health Paper No. 179*. Rome. 2016.
144. Genovese KJ, Anderson RC, Harvey RB, Callaway TR, Poole TL, Edrington TS, et al. Competitive exclusion of *Salmonella* from the gut of neonatal and weaned pigs. *J Food Prot*. 2003;66(8):1353–9. <https://doi.org/10.4315/0362-028x-66.8.1353>
145. Casey PG, Gardiner GE, Casey G, Bradshaw B, Lawlor PG, Lynch PB, et al. A five-strain probiotic combination reduces pathogen shedding and alleviates disease signs in pigs challenged with *Salmonella enterica* serovar typhimurium. *Appl Environ Microbiol*. 2007;73(6):1858–63. <https://doi.org/10.1128/AEM.01840-06>
146. Yin F, Farzan A, Wang QC, Yu H, Yin Y, Hou Y, et al. Reduction of *Salmonella enterica* serovar typhimurium DT104 infection in experimentally challenged weaned pigs fed a lactobacillus-fermented feed. *Foodborne Pathog Dis*. 2014;11(8):628–34. <https://doi.org/10.1089/fpd.2013.1676>
147. Zhao W, Wang Y, Liu S, Huang J, Zhai Z, He C, et al. The dynamic distribution of Porcine microbiota across different ages and Gastrointestinal tract segments. *PLoS ONE*. 2015;10(2):e0117441. <https://doi.org/10.1371/journal.pone.0117441>
148. Kreuzer S, Janczyk P, Assmus J, Schmidt MF, Brockmann GA, Nöckler K. No beneficial effects evident for *Enterococcus faecium* NCIMB 10415 in weaned pigs infected with *Salmonella enterica* serovar typhimurium DT104. *Appl Environ Microbiol*. 2012;78(14):4816–25. <https://doi.org/10.1128/AEM.00395-12>
149. Shanahan F. A commentary on the safety of probiotics. *Gastroenterol. Clin North Am*. 2012;41(4):869–76. <https://doi.org/10.1016/j.gtc.2012.08.006>
150. Letellier A, Messier S, Lessard L, Quessy S. Assessment of various treatments to reduce carriage of *Salmonella* in swine. *Can J Vet Res*. 2000;64(1):27–31.
151. Smith AG, O'Doherty JV, Reilly P, Ryan MT, Bahar B, Sweeney T. The effects of laminarin derived from laminaria digitata on measurements of gut health: selected bacterial populations, intestinal fermentation, mucin gene expression and cytokine gene expression in the pig. *Br J Nutr*. 2011;105(5):669–77. <https://doi.org/10.1017/S0007114510004277>
152. Andrés-Barranco S, Vico JP, Grilló MJ, Mainar-Jaime RC. Reduction of subclinical *Salmonella* infection in fattening pigs after dietary supplementation with a β -galactomannan oligosaccharide. *J Appl Microbiol*. 2015;118(2):284–94. <https://doi.org/10.1111/jam.12713>
153. Hosseini SH, Farhangfar A, Moradi M, Dalir-Naghadeh B. Beyond probiotics: exploring the potential of postbiotics and parabiotics in veterinary medicine. *Res Vet Sci*. 2024;167:105133. <https://doi.org/10.1016/j.rvsc.2023.105133>
154. Nataraj BH, Ali SA, Behare PV, Yadav H. Postbiotics-parabiotics: the new horizons in microbial biotherapy and functional foods. *Microb Cell Fact*. 2020;19:168. <https://doi.org/10.1186/s12934-020-01426-w>
155. Piqué N, Berlanga M, Miñana-Galbis D. Health benefits of Heat-Killed (Tyndalized) probiotics: an overview. *Int J Mol Sci*. 2019;20(10):2534. <https://doi.org/10.3390/ijms20102534>
156. Zhong Y, Wang S, Di H, Deng Z, Liu J, Wand H. Gut health benefit and application of postbiotics in animal production. *J Anim Sci Biotechnol*. 2022. <https://doi.org/10.1186/s40104-022-00688-1>. 13,38.
157. Collado MC, Vinderola G, Salminen S. Postbiotics: facts and open questions. A position paper on the need for a consensus definition. *Benef Microbes*. 2019;10(7):711–9. <https://doi.org/10.3920/BM2019.0015>. PMID: 31965850.
158. Van Immerseel F, Russell JB, Flythe MD, Gantois I, Timbermont L, Pasmans F, et al. The use of organic acids to combat *Salmonella* in poultry: a mechanistic explanation of the efficacy. *Avian Pathol*. 2006;35(3):182–8. <https://doi.org/10.1080/03079450600711045>
159. Mroz Z. Organic acids as potential alternatives to antibiotic growth promoters for pigs. *Adv Pork Prod*. 2005;16:169–82.
160. Gantois I, Ducatelle R, Pasmans F, Haesebrouck F, Hautefort I, Thompson A, et al. Butyrate specifically down-regulates *Salmonella* pathogenicity Island 1 gene expression. *Appl Environ Microbiol*. 2006;72(1):946–9. <https://doi.org/10.1128/AEM.72.1.946-949.2006>
161. Boyen F, Haesebrouck F, Vanparys A, Volf J, Mahu M, Van Immerseel F, et al. Coated fatty acids alter virulence properties of *Salmonella* typhimurium and decrease intestinal colonization of pigs. *Vet Microbiol*. 2008;132(3–4):319–27. <https://doi.org/10.1016/j.vetmic.2008.05.008>
162. Durant JA, Corrier DE, Ricke SC. Short-chain volatile fatty acids modulate the expression of the HliA and InvF genes of *Salmonella* typhimurium. *J Food Prot*. 2000;63(5):573–8. <https://doi.org/10.4315/0362-028x-63.5.573>
163. Huang Y, Suyemoto M, Garner CD, Cicconi KM, Altier C. Formate acts as a diffusible signal to induce *Salmonella* invasion. *J Bacteriol*. 2008;190(12):4233–41. <https://doi.org/10.1128/jb.00205-08>
164. Piva A, Pizzamiglio V, Morlacchini M, Tedeschi M, Piva G. Lipid microencapsulation allows slow release of organic acids and natural identical flavors along the swine intestine. *J Anim Sci*. 2007;85:486–93. <https://doi.org/10.2527/jas.2006-323>
165. Friendship RM, Mounchili A, McEwen S, Rajic A. Critical Review of On-farm Intervention Strategies Against *Salmonella*. 2009. https://www.researchgate.net/publication/242671033_Critical_review_of_on-farm_intervention_strategies_against_Salmonella#fullTextFileContent. Accessed 08 November 24.
166. De Lange CFM, Pluske J, Gong J, Nyachoti CM. Strategic use of feed ingredients and feed additives to stimulate gut health and development in young pigs. *Livest Sci*. 2010;134:124–34. <https://doi.org/10.1016/j.livsci.2010.06.117>
167. Wilhelm B, Rajić A, Parker S, Waddell L, Sanchez J, Fazil A, et al. Assessment of the efficacy and quality of evidence for five on-farm interventions for *Salmonella* reduction in grow-finish swine: a systematic review and meta-analysis. *Prev Vet Med*. 2012;107(1–2):1–20. <https://doi.org/10.1016/j.prevetmed.2012.07.011>
168. Walia K, Argüello H, Lynch H, Leonard FC, Grant J, Yearsley D, et al. Effect of feeding sodium butyrate in the late finishing period on *Salmonella* carriage, seroprevalence, and growth of finishing pigs. *Prev Vet Med*. 2016;131:79–86. <https://doi.org/10.1016/j.prevetmed.2016.07.009>
169. Fabá LI, Litjens R, Allaart J, Roubos-, van den Hil P. Feed additive blends fed to nursery pigs challenged with *Salmonella*. *J Anim Sci*. 2020;1–10. <https://doi.org/10.1093/jas/skz382>
170. Gomes da Silva D, Gonçalves de Oliveira Moura EA, Carnevalli Sanches TV, Turco CH, Belloni Zambotti B, Moreira Petri FA, et al. Use of organic acids to reduce *Salmonella* typhimurium excretion in swine. *Braz J Vet Res Anim Sci*. 2023;60. <https://doi.org/10.11606/issn.1678-4456.bjvras.2023.198402>
171. Álvarez-Ordóñez A, Fernández A, Bernardo A, López M. Comparison of acids on the induction of an acid tolerance response in *Salmonella* typhimurium, consequences for food safety. *Meat Sci*. 2009;81(1):65–70. <https://doi.org/10.1016/j.meatsci.2008.06.019>
172. Creus E, Pérez JF, Peralta B, Baucells F, Mateu E. Effect of acidified feed on the prevalence of *Salmonella* in market-age pigs. *Zoonoses Public Health*. 2007;54(8):314–9. <https://doi.org/10.1111/j.1863-2378.2007.01069.x>
173. Argüello H, Carvajal A, Costillas S, Rubio P. Effect of the addition of organic acids in drinking water or feed during part of the finishing period on

- the prevalence of *Salmonella* in finishing pig. *Foodborne Pathog Dis.* 2013;10(10):842–9. <https://doi.org/10.1089/fpd.2013.1497>
174. De Ridder L, Maes D, Dewulf J, Pasmans F, Boyen F, Haesebrouck F, et al. Evaluation of three intervention strategies to reduce the transmission of *Salmonella* typhimurium in pigs. *Vet J.* 2013;197(3):613–8. <https://doi.org/10.1016/j.tvjl.2013.03.026>
175. Casanova-Higes A, Andrés-Barranco S, Mainar-Jaime RC. Effect of the addition of protected sodium butyrate to the feed on *Salmonella* spp. Infection dynamics in fattening pigs. *Anim Feed Sci Tech.* 2017;231:12–8. <https://doi.org/10.1016/j.anifeedsci.2017.06.008>
176. Lynch H, Leonard FC, Walia K, Lawlor PG, Duffy G, Fanning S, et al. Investigation of in-feed organic acids as a low cost strategy to combat *Salmonella* in grower pigs. *Prev Vet Med.* 2017;139:50–7. <https://doi.org/10.1016/j.prevetmed.2017.02.008>
177. Bester C, Käsbohrer A, Wilkins N, Carreira GC, Marschik T. Identification of cost-effective biosecurity measures to reduce *Salmonella* along the pork production chain. *Front Vet Sci.* 2024;11. <https://doi.org/10.3389/fvets.2024.1380029>
178. Cantini F. Compositions Containing C1 to C7 Organic Acid Monoglycerides and Glycerol, Their Preparation and Use as Antibacterials and Anti-Mould Agents. Google patents. 2015. <https://patents.google.com/patent/WO2010106488A2/en>. Accessed 01 November 24.
179. Gomez-Osorio LM, Yepes-Medina V, Ballou A, Parini M, Angel R. Short and medium chain fatty acids and their derivatives as a natural strategy in the control of necrotic enteritis and microbial homeostasis in broiler chickens. *Front Vet Sci.* 2021;8:773372. <https://doi.org/10.3389/fvets.2021.773372>
180. Bernad-Roche M, Marín-Alcalá CM, Vico JP, Mainar-Jaime R. *Salmonella* control in fattening pigs through the use of esterified formic acid in drinking water shortly before slaughter. *Animals.* 2023;13(18):2814. <https://doi.org/10.3390/ani13182814>
181. Huff WE, Huff GR, Rath NC, Balog JM, Donoghue AM. Alternatives to antibiotics: utilization of bacteriophage to treat colibacillosis and prevent foodborne pathogens. *Poult Sci.* 2005;84(4):655–9. <https://doi.org/10.1093/ps/84.4.655>
182. Thanki AM, Clavijo V, Healy K, Wilkinson RC, Sicheritz-Pontén T, Millard AD, et al. Development of a phage cocktail to target *Salmonella* strains associated with swine. *Pharmaceutics.* 2022;15(1):58. <https://doi.org/10.3390/ph15010058>
183. Callaway TR, Edrington TS, Brabban A, Kutter B, Karriker L, Stahl C, et al. Evaluation of phage treatment as a strategy to reduce *Salmonella* populations in growing swine. *Foodborne Pathog Dis.* 2011;8(2):261–6. <https://doi.org/10.1089/fpd.2010.0671>
184. Saez AC, Zhang J, Rostagno MH, Ebner PD. Direct feeding of microencapsulated bacteriophages to reduce *Salmonella* colonization in pigs. *Foodborne Pathog Dis.* 2011;8(12):1269–74. <https://doi.org/10.1089/fpd.2011.0905>
185. Thanki AM, Mignard G, Atterbury RJ, Barrow P, Millard AD, Clokie MRJ. Prophylactic delivery of a bacteriophage cocktail in feed significantly reduces *Salmonella* colonization in pigs. *Microbiol Spectr.* 2022;10(3):e0042222. <https://doi.org/10.1128/spectrum.00422-22>
186. Wiggins BA, Alexander M. Minimum bacterial density for bacteriophage replication: implications for significance of bacteriophages in natural ecosystems. *Appl Environ Microbiol.* 1985;49(1):19–23. <https://doi.org/10.1128/aem.49.1.19-23.1985>
187. Switt AI, den Bakker HC, Vongkamjan K, Hoelzer K, Warnick LD, Cummings KJ, et al. *Salmonella* bacteriophage diversity reflects host diversity on dairy farms. *Food Microbiol.* 2013;36(2):275–85. <https://doi.org/10.1016/j.fm.2013.06.014>
188. Zhang J, Li Z, Cao Z, Wang L, Li X, Li S, et al. Bacteriophages as antimicrobial agents against major pathogens in swine: a review. *J Anim Sci Biotechnol.* 2015;6(1):39. <https://doi.org/10.1186/s40104-015-0039-7>
189. Yousefi MH, Wagemans J, Shekarforoush SS, Vallino M, Pozhydaieva N, Höfer K, et al. Isolation and molecular characterization of the *Salmonella* typhimurium orphan phage Arash. *BMC Microbiol.* 2023;23(1):297. <https://doi.org/10.1186/s12866-023-03056-9>
190. Arista-Regalado AD, Viera-Segura O, de Oca SA, Hernández-Hernández L, González-Aguilar DG, León JB. Characterization and efficacy of *Salmonella* phage cocktail PHA46 in the control of *Salmonella* Newport and typhimurium internalized into Cherry tomatoes. *Int J Food Microbiol.* 2024;164:110745. <https://doi.org/10.1016/j.jifoodmicro.2024.110745>
191. Li L, Fan R, Chen Y, Zhang Q, Zhao X, Hu M, et al. Characterization, genome analysis, and therapeutic evaluation of a novel *Salmonella* phage vB_SalS_JNS02: a candidate bacteriophage for phage therapy. *Poult Sci.* 2024;103(7):103845. <https://doi.org/10.1016/j.psj.2024.103845>
192. Naureen Z, Malacarne D, Anpilogov K, Dautaj A, Camilleri G, Cecchin S, et al. Comparison between American and European legislation in the therapeutic and alimentary bacteriophage usage. *Acta Biomed.* 2020;9(13–S):e2020023. <https://doi.org/10.23750/abm.v9i13-S.10815>
193. Chan BK, Abedon ST. Phage therapy Pharmacology phage cocktails. *Adv Appl Microbiol.* 2012;78:1–23. <https://doi.org/10.1016/B978-0-12-394805-2.00001-4>
194. Ackermann HW, Tremblay D, Moineau S. Long-term bacteriophage preservation. *Journal: World Fed Cult Collect Newslett.* 2004.
195. Dlamini SB, Gigante AM, Hooton SPT, Atterbury RJ. Efficacy of different encapsulation techniques on the viability and stability of diverse phage under simulated gastric conditions microorganisms. 2023;11(10):2389. <https://doi.org/10.3390/microorganisms11102389>
196. Vila MMDC, Balcão LMN, Balcão VM. Phage Delivery Strategies for Biocontrolling Human, Animal, and Plant Bacterial Infections: State of the Art. *Pharmaceutics* 2024;16,374. <https://doi.org/10.3390/pharmaceutics16030374>
197. Jończyk E, Klak M, Międzybrodzki R, Górski A. The influence of external factors on bacteriophages—review. *Folia Microbiol.* 2011;56(3):191–200. <https://doi.org/10.1007/s12223-011-0039-8>
198. Oechslin F. Resistance development to bacteriophages occurring during bacteriophage therapy. *Viruses.* 2018;10(7):351. <https://doi.org/10.3390/v10070351>
199. Yu J, Zhang H, Ju Z, Huang J, Lin C, Wu J, et al. Increased mutations in lipopolysaccharide biosynthetic genes cause time-dependent development of phage resistance in *Salmonella*. *Antimicrob Agents Chemother.* 2024;68(2):e0059423. <https://doi.org/10.1128/aac.00594-23>
200. Zeng Y, Shen M, Liu S, Zhou X. Characterization and resistance mechanism of phage-resistant strains of *Salmonella* enteritidis. *Poult Sci.* 2024;103(6):103756. <https://doi.org/10.1016/j.psj.2024.103756>
201. Pereira C, Moreirinha C, Lewicka M, Almeida P, Clemente C, Cunha Â, et al. Bacteriophages with potential to inactivate *Salmonella* typhimurium: use of single phage suspensions and phage cocktails. *Virus Res.* 2016;220:179–92. <https://doi.org/10.1016/j.virusres.2016.04.020>
202. Acton L, Pye HV, Thilliez G, Kolenda R, Matthews M, Turner AK, et al. Collateral sensitivity increases the efficacy of a rationally designed bacteriophage combination to control *Salmonella* enterica. *J Virol.* 2024;19(3):e0147623. <https://doi.org/10.1128/jvi.01476-23>
203. Martinez-Soto CE, McClelland M, Kropinski AM, Lin JT, Khursigara CM, Anany H. Multireceptor phage cocktail against *Salmonella* enterica to circumvent phage resistance. *Microlife.* 2024;5:uqae003. <https://doi.org/10.1093/femsml/uqae003>
204. Gebru E, Lee JS, Son JC, Yang SY, Shin SA, Kim B, et al. Effect of probiotic-, bacteriophage-, or organic acid-supplemented feeds or fermented soybean meal on the growth performance, acute-phase response, and bacterial shedding of grower pigs challenged with *Salmonella* enterica serotype typhimurium. *J Anim Sci.* 2010;88(12):3880–6. <https://doi.org/10.2527/jas.2010-2939>
205. Seo BJ, Song ET, Lee K, Kim JW, Jeong CG, Moon SH, et al. Evaluation of the broad-spectrum lytic capability of bacteriophage cocktails against various *Salmonella* serovars and their effects on weaned pigs infected with *Salmonella* typhimurium. *J Vet Med Sci.* 2018;80(6):851–60. <https://doi.org/10.1292/jvms.17-0501>
206. European Medicines Agency (EMA). Guideline on quality, safety and efficacy of veterinary medicinal products specifically designed for phage therapy. Committee for Veterinary Medicinal Products (CVMP); 2023. EMA/CVMP/NTWP/32862/2022.
207. Hill A, Simons R, Ramnial V, Tennant J, Denman S, Cheney et al. Quantitative Microbiological Risk Assessment on *Salmonella* in Slaughter and Breeder pigs: Final Report. EFSA Supporting Publications 2011;EN-46. <https://doi.org/10.2903/sp.efsa.2010.EN-46>
208. Binter C, Straver JM, Häggblom P, Bruggeman G, Lindqvist PA, Zentek J, et al. Transmission and control of *Salmonella* in the pig feed chain: a conceptual model. *Int J Food Microbiol.* 2011;1(145 Suppl 1):S7–17. <https://doi.org/10.1016/j.jifoodmicro.2010.09.001>
209. Wierup M, Kristoffersen T. Prevention of *Salmonella* contamination of finished soybean meal used for animal feed by a Norwegian production plant despite frequent *Salmonella* contamination of Raw soy beans, 1994–2012. *Acta Vet Scand.* 2014;56:41. <https://doi.org/10.1186/s13028-014-0041-7>
210. Parker EM, Parker AJ, Short G, O'Connor AM, Wittum TE. *Salmonella* detection in commercially prepared livestock feed and the Raw ingredients and equipment used to manufacture the feed: A systematic review and meta-analysis.

- Prev Vet Med. 2022;198:105546. <https://doi.org/10.1016/j.prevetmed.2021.105546>
211. Wierup M. The importance of hazard analysis by critical control point for effective pathogen control in animal feed: assessment of *Salmonella* control in feed production in Sweden, 1982–2005. *Foodborne Pathog Dis*. 2023;20(12):545–52. <https://doi.org/10.1089/fpd.2023.0067>
212. European Commission, Directorate-General for Health and Food Safety. Final report of an audit carried out in the Czech Republic from 6 to 17 March 2023 in order to evaluate the implementation of official controls on feed hygiene. DG(SANTE) 2023–7697. Brussels: European Commission. 2023 Jun 29. https://ec.europa.eu/food/audits-analysis/rep_details_en.cfm?rep_inspection_ref=2023-7697. Accessed 02 July 2025.
213. Shaji S, Selvaraj RK, Shanmugasundaram R. *Salmonella* infection in poultry: A review on the pathogen and control strategies. *Microorganisms*. 2023;11(11):2814. <https://doi.org/10.3390/microorganisms11112814>
214. Smith RP, Andres V, Martelli F, Gosling B, Marco-Jimenez F, Vaughan K, et al. Maternal vaccination as a *Salmonella* typhimurium reduction strategy on pig farms. *J Appl Microbiol*. 2018;124(1):274–85. <https://doi.org/10.1111/jam.13609>
215. Wales AD, Davies RH. *Salmonella* vaccination in pigs: A review. *Zoonoses Public Health*. 2017;64(1):1–13. <https://doi.org/10.1111/zph.12256>
216. de la Cruz ML, Conrado I, Nault A, Perez A, Dominguez L, Alvarez J. Vaccination as a control strategy against *Salmonella* infection in pigs: A systematic review and meta-analysis of the literature. *Res Vet Sci*. 2017;114:86–94. <https://doi.org/10.1016/j.rvsc.2017.03.005>
217. Gelaude P, Schlepers M, Verlinden M, Laanen M, Dewulf J. Biocheck. UGent: a quantitative tool to measure biosecurity at broiler farms and the relationship with technical performances and antimicrobial use. *Poult Sci*. 2014;93(11):2740–51. <https://doi.org/10.3382/ps.2014-04002>
218. Backhans A, Sjölund M, Lindberg A, Emanuelson U. Biosecurity level and health management practices in 60 Swedish farrow-to-finish herds. *Acta Vet Scand*. 2015;57(1):14. <https://doi.org/10.1186/s13028-015-0103-5>
219. Postma M, Backhans A, Collineau L, Loesken S, Sjölund M, Belloc C, et al. The biosecurity status and its associations with production and management characteristics in farrow-to-finish pig herds. *Animal*. 2016;10(03):478–89. <https://doi.org/10.1017/S1751731115002487>
220. Corcoran M, Morris D, De Lappe N, O'Connor J, Lalor P, Dockery P, et al. Commonly used disinfectants fail to eradicate *Salmonella* enterica biofilms from food contact surface materials. *Appl Environ Microbiol*. 2014;80:1507–14. <https://doi.org/10.1128/AEM.03109-13>
221. Winfield MD, Groisman EA. Role of nonhost environments in the lifestyles of *Salmonella* and *Escherichia coli*. *Appl Environ Microbiol*. 2003;69(7):3687–94. <https://doi.org/10.1128/AEM.69.7.3687-3694.2003>
222. Ohashi I, Kobayashi S, Tamamura-Andoh Y, Arai N, Takamatsu D. Disinfectant resistance of *Salmonella* in vitro contaminated poultry house models and investigation of efficient disinfection methods using these models. *J Vet Med Sci*. 2002;84(12):1633–44. <https://doi.org/10.1292/jvms.22-0311>
223. Sevilla-Navarro S, Catalá-Gregori P, García C, Cortés V, Marin C. *Salmonella* infantis and *Salmonella* enteritidis specific bacteriophages isolated from poultry faeces as a complementary tool for cleaning and disinfection against *Salmonella*. *Comp Immunol Microbiol Infect Dis*. 2020;68:101405. <https://doi.org/10.1016/j.cimid.2019.101405>
224. D'accolti M, Soffritti I, Mazzacane S, Caselli E. Bacteriophages as a potential 360-degree pathogen control strategy. *Microorganisms*. 2021;9:1–15. <https://doi.org/10.3390/microorganisms9020261>
225. Sevilla-Navarro S, Marin C, Cortés V, García C, Vega S, Catalá-Gregori P. Autophagy as a control measure for *Salmonella* in laying hens. *Poult Sci*. 2018;97:4367–73. <https://doi.org/10.3382/ps/pey294>
226. Sevilla-Navarro S, Torres-Boncompagni J, García-Llorens J, Bernabéu-Gimeno M, Domingo-Calap P, Catalá-Gregori P. Fighting *Salmonella* infantis: bacteriophage-driven cleaning and disinfection strategies for broiler farms. *Front Microbiol*. 2024;15:1401479. <https://doi.org/10.3389/fmicb.2024.1401479>
227. Zdolec N, Kotsiri A, Houf K, Alvarez-Ordóñez A, Blagojevic B, Karabasil N, et al. Systematic review and Meta-Analysis of the efficacy of interventions applied during primary processing to reduce microbial contamination on pig carcasses. *Foods*. 2022;11(14):2110. <https://doi.org/10.3390/foods11142110>
228. Viltrop A, Niine T, Tobias T, Sassu EL, Bartolo ID, Pavoni E, et al. A review of slaughter practices and their effectiveness to control Microbial - esp. *Salmonella* spp. - Contamination of pig carcasses. *J Food Prot*. 2023;86(11):100171. <https://doi.org/10.1016/j.jfp.2023.100171>
229. van Netten P, Mossel DA, Huis in 't Veld J. Lactic acid decontamination of fresh pork carcasses: a pilot plant study. *Int J Food Microbiol*. 1995;25(1):1–9. [https://doi.org/10.1016/0168-1605\(94\)00039-9](https://doi.org/10.1016/0168-1605(94)00039-9)
230. Pipek P, Houška M, Hoke K, Jeleníková J, Kýhos K, Šikulová M. Decontamination of pork carcasses by steam and lactic acid. *J Food Eng*. 2006;74(2):224–31. <https://doi.org/10.1016/j.jfoodeng.2005.03.015>
231. Gellynck X, Messens W, Halet D, Grijspeerd K, Hartnett E, Viaene J. Economics of reducing *Campylobacter* at different levels within the Belgian poultry meat chain. *J Food Prot*. 2008;71(3):479–85. <https://doi.org/10.4315/0362-028x-71.3.479>
232. Loretz M, Stephan R, Zweifel C. Antibacterial activity of decontamination treatments for pig carcasses. *Food Control*. 2011;22:1121–5. <https://doi.org/10.1016/j.foodcont.2011.01.013>
233. Hugas M, Tsigarida E. Pros and cons of carcass decontamination: the role of the European food safety authority. *Meat Sci*. 2008;78(1–2):43–52. <https://doi.org/10.1016/j.meatsci.2007.09.001>
234. Wang H, Duan D, Wu Z, Xue S, Xu X, Zhou G. Primary concerns regarding the application of electrolyzed water in the meat industry. *Food Control*. 2019;95:50–6. <https://doi.org/10.1016/j.foodcont.2018.07.049>
235. Piriou P, Devesa R, Puget S, Thomas-Danguin T, Zraïck F. Evidence of regional differences in Chlorine perception by consumers: sensitivity differences or habituation? *J Water Supply: Res Technology-Aqua*. 2015;64(7):783–92. <https://doi.org/10.2166/aqua.2014.097>
236. Albert T, Braun PG, Saffaf J, Wiacek C. Physical methods for the decontamination of meat surfaces. *Curr Clin Micro Rpt*. 2021;8:9–20. <https://doi.org/10.1007/s40588-021-00156-w>
237. Indarto R, Irawan AN, Subroto E. Meat irradiation: A comprehensive review of its impact on food quality and safety. *Foods*. 2023;12(9):1845. <https://doi.org/10.3390/foods12091845>
238. Abd-El Wahab A, Basiouni S, El-Seedi HR, Ahmed MFE, Bielke LR, Hargis B, et al. An overview of the use of bacteriophages in the poultry industry: successes, challenges, and possibilities for overcoming breakdowns. *Front Microbiol*. 2023;21:14:1136638. <https://doi.org/10.3389/fmicb.2023.1136638>
239. Miller GY, Liu X, McNamara PE, Barber DA. Influence of *Salmonella* in pigs preharvest and during pork processing on human health costs and risks from pork. *J Food Prot*. 2005;68(9):1788–98. <https://doi.org/10.4315/0362-028x-68.9.1788>
240. Goldbach SG, Alban L. A cost-benefit analysis of *Salmonella*-control strategies in Danish pork production. *Prev Vet Med*. 2006;77(1–2):1–14. <https://doi.org/10.1016/j.prevetmed.2005.10.008>
241. Lawson LG, Jensen JD, Christiansen P, Lund M. Cost-effectiveness of *Salmonella* reduction in Danish abattoirs. *Int J Food Microbiol*. 2009;134(1–2):126–32. <https://doi.org/10.1016/j.jfoodmicro.2009.03.024>
242. Hochreutener M, Zweifel C, Corti S, Stephan R. Effect of a commercial steam-vacuuming treatment implemented after slaughtering for the decontamination of cattle carcasses. *Ital J Food Saf*. 2017;6(3):6864. <https://doi.org/10.4081/jifs.2017.6864>
243. Totton SC, Glanville JM, Dzikanunhenga RS, Dickson JS, O'Connor AM. Systematic review of the magnitude of change in prevalence and quantity of *Salmonella* after administration of pathogen reduction treatments on pork carcasses. *Anim Health Res Rev*. 2016;17(1):39–59. <https://doi.org/10.1017/S1466252316000025>
244. Rhouma M, Madec JY, Laxminarayan R. Colistin: from the shadows to a one health approach for addressing antimicrobial resistance. *Int J Antimicrob Agents*. 2023;61(2):106713. <https://doi.org/10.1016/j.ijantimicag.2023.106713>

Publisher's note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.