

1 Running title: *Y. enterocolitica* in pigs

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3 Original research paper

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5 Prevalence and dynamics of pathogenic *Yersinia enterocolitica* 4/O:3 among Finnish piglets,
6 fattening pigs, and sows

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23 ABSTRACT

24

25 Pigs are considered the main reservoir of *Yersinia enterocolitica*, hence understanding the ecology of
26 this foodborne pathogen at the farm level is crucial. We calculated Bayesian estimates for the ability
27 of a commercial ELISA diagnostic test kit to detect antibodies against pathogenic *Yersinia* in pigs. The
28 sensitivity and specificity of the test were 75.4% and 98.1%, respectively. We also studied the
29 dynamics of *Y. enterocolitica* infection in three farrow-to-finish pig farms by following the same 30
30 pens of pigs through their lifetime from farrowing unit to slaughterhouse. Each farm was sampled four
31 times, and 864 fecal and 730 serum samples were collected altogether.
32 Pathogenic *Y. enterocolitica* 4/O:3 was isolated from 31.6% of the fecal samples by culturing,
33 and *Yersinia* antibodies were detected in 38.2% of the serum samples with the commercial ELISA test.
34 The pathogen was not isolated from farrowing units or all-in/all-out weaning units. However, in the
35 weaning and fattening units using continuous management systems, the pathogen was isolated from
36 every pen at some point of the study. After the pigs were transported into slaughterhouse, 150 tonsils
37 were collected and 96.7% were positive by culturing. Among the strains isolated from feces and
38 tonsils, 56 different genotypes of pathogenic *Y. enterocolitica* 4/O:3 were found by multiple-locus
39 variable-number tandem repeat analysis (MLVA). Finally, we collected tonsils of 266 sows from 115
40 farrowing farms, and *Y. enterocolitica* 4/O:3 was detected in 6.0% of the samples by the culture
41 method, while 77.1% of the tonsils were serologically positive; the estimate for true seroprevalence
42 was 95.8%. In conclusion, sows may not be the main source of *Y. enterocolitica* for piglets, although
43 sows may still play a role in maintaining *Y. enterocolitica* in pig farms. Instead, pigs appear to get this
44 foodborne pathogen mainly during the fattening period, especially if continuous management is
45 applied.

- 46 Key words: antibody, ecology, ELISA, enteropathogenic *Yersinia*, foodborne pathogen, MLVA, farm,
- 47 reservoir, slaughterhouse, swine

50 Yersiniosis is the third most commonly reported bacterial zoonosis in the European Union and is primarily
51 caused by pathogenic *Y. enterocolitica* (EFSA and ECDC, 2018). Yersiniosis manifests as febrile gastrointestinal
52 disease but may also lead to other symptoms and complications such as erythema nodosum or reactive
53 arthritis (Fredriksson-Ahomaa et al., 2010; Bottone et al., 2015). Pigs are considered to be significant
54 reservoirs for *Y. enterocolitica*, and the main source of human infections. *Y. enterocolitica* contaminates
55 carcasses and offal during slaughter processes (Fredriksson-Ahomaa et al., 2000; Laukkanen et al., 2009) and
56 contaminated pork and other food of swine origin have been associated with yersiniosis (Tauxe et al., 1987;
57 Lee et al., 1990; Ostroff et al., 1994; Huovinen et al., 2010). Moreover, genotypically similar strains of
58 *Y. enterocolitica* have been isolated from yersiniosis patients and pigs (Fredriksson-Ahomaa et al., 2001;
59 Fredriksson-Ahomaa et al., 2006; Virtanen et al., 2013).

60 Pigs are symptomless carriers of enteropathogenic *Yersinia*, especially *Y. enterocolitica*, but
61 the prevalence is variable depending on factors such as age, sampling and detection methodology, farm
62 management, and biosecurity level (reviewed by Laukkanen-Ninios et al., 2014). Despite some geographical
63 variation, bioserotype 4/O:3 is the type most frequently isolated in pigs (Laukkanen-Ninios et al., 2014).
64 Newborn piglets are negative for *Y. enterocolitica*. Piglets start excreting the pathogen in feces around the
65 age of 1–3 months with peak of excretion around the age of 2–5 months, the fecal prevalence starts reducing
66 thereafter, and pigs tend to remain seropositive for longer periods (Fukushima et al., 1983; Nielsen et al.,
67 1996; Gürtler et al., 2005; Nesbakken et al., 2006; Bowman et al., 2007; Wehebrink et al., 2008; Virtanen et
68 al., 2012; Vilar et al., 2013).

69 Based on numerous studies, variable number of slaughter pigs carry enteropathogenic
70 *Y. enterocolitica* in tonsils and feces; 2–93% and 0.5–76%, respectively (reviewed by Laukkanen-Ninios et al.
71 2014). Usually most pigs are seropositive at slaughter age (Skjerve et al., 1998; Thibodeau et al., 2001;
72 Nesbakken et al., 2006; Virtanen et al., 2012; Vilar et al., 2013, Vilar et al., 2015; Bonardi et al., 2016;
73 Lorencova et al., 2016). Only a few carrier pigs are needed to spread the infection within and between pig

74 farms (Virtanen et al., 2012; Virtanen et al., 2014). In contrast to fattening pigs, the prevalence of
75 *Y. enterocolitica* in sows is less studied. The reported prevalence has been relatively low, varying between
76 0% and 14% (Fukushima et al., 1983; Korte et al., 2004; Bowman et al., 2005; Gürtler et al., 2005; Wehebrink
77 et al., 2008; Farzan et al., 2010; Vilar et al., 2013). However, most sows still appear to be seropositive (Vilar
78 et al., 2013; Vilar et al., 2015) suggesting development of immunity against *Yersinia*. More studies are needed
79 to understand the dynamics of *Y. enterocolitica* at the farm level, especially the role of sows and piglets in
80 *Yersinia* ecology.

81 A commercially available enzyme-linked immunosorbent assay (ELISA) has been used to
82 determine antibodies against pathogenic *Yersinia* in pigs. However, no diagnostic test should be considered
83 fully sensitive and specific. Therefore, we have calculated Bayesian estimations for the sensitivity and
84 specificity of the ELISA test (Vilar et al., 2015). The updated estimates are needed, because the manufacturer
85 has revised the test and increased the recommended cut-off value.

86 The aim of our study was to assess the dynamics of *Y. enterocolitica* infection in three farrow-
87 to-finish pig farms. In addition, we studied the prevalence of pathogenic *Y. enterocolitica* and the
88 seroprevalence of *Yersinia* antibodies in the tonsils of sows collected at slaughterhouses.

90

91 *Experimental plan and sampling*

92 To assess the dynamics of pathogenic *Y. enterocolitica* in pig farms (study I), three Finnish farrow-to-finish
93 farms (A-C) with known *Yersinia* positivity were followed in a longitudinal study. From each farm, pigs from
94 six pens (mean 7.3 pigs per pen) were followed throughout their lifetime (i.e. from farrowing units to weaning
95 and fattening units, and finally the slaughterhouse). In farm B, two subsequent longitudinal follow-ups were
96 performed at a nine-month interval to study the persistence of pathogenic *Y. enterocolitica* strains in the
97 farm; 12 pens were included in the second follow-up. Pigs from different pens were not mixed, except in the
98 fattening unit of farm C, where the pigs were regrouped in two larger pens of approximately 20 pigs each.
99 Conventional straw bedding was used in all the farms and units, except the fattening unit of farm C, which
100 used deep peat bedding. All-in/all-out management systems were used in the weaning units of farms B and
101 C to avoid any contact between pigs from different groups. In contrast, the weaning unit of farm A and the
102 fattening units of all the three farms used continuous management systems.

103 At the farms, pens were sampled four times, approximately once per month. On average, feces
104 and blood samples were collected from 96% and 88% of the pigs from every pen, respectively. Furthermore,
105 fecal samples were collected from the mother sows ($n = 24$) during the first sampling at all but farm A. Rectal
106 swabs were used for piglets (first sampling), and also for older pigs if feces could not be collected. Finally,
107 150 tonsils were collected at the slaughterhouse (last sampling).

108 To assess the prevalence of pathogenic *Y. enterocolitica* and the seroprevalence of *Yersinia*
109 antibodies in sows, tonsils of 266 sows from 115 farrowing farms were collected from two Finnish
110 slaughterhouses (study II).

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112 *Isolation and identification of pathogenic Y. enterocolitica*

113 Fecal and tonsil samples from study I and tonsil samples from study II were screened for the presence of
114 pathogenic *Y. enterocolitica* according to the method of the Department of Food Hygiene and Environmental
115 Health (University of Helsinki) (Laukkanen et al. 2010). Briefly, for the fecal samples (study I), a three-step
116 isolation was used before the plating on cefsulodin-irgasan-novobiosin agar (Schiemann, 1979): (i)
117 immediately, and (ii) after one week and (iii) after two weeks of cold enrichment at 4°C. In the case of (iii),
118 an alkali treatment with 0.25% potassium hydroxide solution (Aulisio et al., 1980) was applied after the cold
119 enrichment. For the tonsil samples (studies I and II), a four-step isolation was used, as an additional
120 enrichment in irgasan-ticarcillin-potassium chloride broth (Wauters et al., 1988) at 25°C for two days was
121 applied, but for the sow tonsils (study II) the immediate plating on CIN agar was omitted. After the culturing,
122 presumptive colonies were identified with an urea hydrolysis test, API 20E test (BioMérieux, Marcy l'Etoile,
123 France), biotyping, serotyping and a multiplex PCR targeting the virulence genes *ail* (Nakajima et al., 1992)
124 located on the chromosome, and *virF* (Kaneko et al., 1996) located on the virulence plasmid pYV Laukkanen
125 et al. (2010). To investigate the spread of pathogenic *Y. enterocolitica* among pigs, pathogenic isolates from
126 the three farrow-to-finish farms were genotyped using a multiple-locus variable-number tandem repeat
127 analysis (MLVA) method (Alakurtti et al. 2016).

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129 *Enumeration of Yersinia antibodies and performance of the ELISA test*

130 The serum samples collected from the farrow-to-finish farms (study I), as well as the tissue fluid samples from
131 the tonsils of sows (study II), were screened for the presence of antibodies against *Yersinia* outer membrane
132 proteins by using a commercial ELISA test (formerly Pigtype Yopscreen, Labor Diagnostik, Leipzig, Germany;
133 currently Pigtype Yersinia Ab, Qiagen, Leipzig, Germany) according to the manufacturer's instructions (short
134 protocol). Samples with activity values (S/P ratios) of 0.3 or higher are considered positive.

135 Estimates of true prevalence were calculated by using a Bayesian approach as follows: prior
136 distributions of sensitivity and specificity were defined as beta probability distributions using a test with
137 imperfect sensitivity and specificity (Thrusfield, 2007). For the ELISA test, the estimates of sensitivity and

138 specificity were recalculated from Vilar et al. (2015) using the current recommended cut-off value (S/P ratio)
139 of 0.3 instead of 0.2. These sensitivity and specificity estimations were used to calculate the prior beta
140 probability distributions, $Se \sim \text{beta}(\alpha_{Se}, \beta_{Se})$, $Sp \sim \text{beta}(\alpha_{Sp}, \beta_{Sp})$, thus sensitivity was beta (106.79, 35.52)
141 and the specificity was beta (46.81, 1.89). For the isolation using the culture method, the sensitivity (77.9%)
142 reported by Laukkanen et al. (2010) was used to calculate the prior beta probability distribution beta (39.79,
143 12.01), whereas a non-informative beta (1,1) was used for the specificity. Models were constructed in
144 OpenBugs 3.2.2 as described by Vilar et al. (2015). Briefly, inferences were based on 50,000 iterations after
145 a burn-in for convergence of 1,000 iterations. For the apparent seroprevalence in the tonsils of sows (study
146 II), exact binomial 95% confidence intervals described by Agresti and Coull (1998) were calculated in
147 Microsoft Excel 2016 (Microsoft Corporation, Redmond, WA).

148

149 *Statistical analysis*

150 To investigate pathogenic *Y. enterocolitica* in the three farrow-to-finish farms (study I), the pen was the
151 experimental unit and considered positive for *Y. enterocolitica* by culturing, if any fecal sample tested
152 positive, or by serology, if any blood sample tested positive.

153 The statistical difference between isolation of pathogenic *Y. enterocolitica* and the presence
154 of *Yersinia* antibodies in the tonsils of sows (study II) was assessed by McNemar's test in IBM SPSS Statistics
155 24.0 (IBM, Armonk, NY).

Results

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158 *Performance of the ELISA test*

159 The recalculated estimates for the sensitivity and specificity of the commercial ELISA test were 75.4% and
160 98.1%, respectively (Table 1).

161

162 *Transmission of Y. enterocolitica in farrow-to-finish farms (study I)*

163 Among the 1014 samples cultured, 418 (41.2%) were positive for pathogenic *Y. enterocolitica* 4/O:3: 273 of
164 864 (31.6%) of the fecal samples were positive, and 145 of 150 (96.7%) of the tonsil samples. Of the 418
165 strains genotyped, all were *ail* positive, and 402 were *virF* positive. Antibodies against pathogenic *Yersinia*
166 were found in a total of 38.2% of the blood samples (279 of 730). In the farrowing units (first sampling), no
167 pathogenic *Y. enterocolitica* was detected, although some piglets were seropositive (Table 2). In addition, all
168 the 24 fecal samples taken from sows in the farrowing units were negative. In the weaning units, the
169 pathogen was detected in all of the pens of farm A but not in pigs of farms B and C, which used all-in/all-out
170 management systems (Table 2). In the fattening units using continuous management systems, most of the
171 pens were positive. At slaughter age, 96.3% of the pigs carried pathogenic *Y. enterocolitica* in their tonsils
172 (Table 2).

173 From the three farms studied, a total of 56 different MLVA types of pathogenic
174 *Y. enterocolitica* 4/O:3 were found (Table 3, Supplemental figure). During the follow-ups of A, B-I, B-II, and C,
175 a total of 9, 17, 21, and 15 different MLVA types were detected, respectively. The farms had their own MLVA
176 types, except the most frequently isolated type in farm C which was also found at the slaughterhouse in one
177 pig of farm B. Of the MLVA types isolated from farm B at the farm level excluding the slaughterhouse
178 sampling, eight genotypes were detected during only the first follow-up and 10 genotypes during only the
179 second follow-up. Four genotypes were detected during both follow-ups. The highest count of unique
180 genotypes ($n = 14$) was found in the fattening unit of farm C (Table 3). During the follow-ups of A, B-I, B-II and
181 C, 2, 5, 8 and 1 new MLVA types were found in the slaughterhouse, respectively.

182

183 *Prevalence of Y. enterocolitica in the tonsils of sows (study II)*

184 Pathogenic *Y. enterocolitica* was detected in 6.0% (95% CI: 3.5–9.6%) of the tonsil samples of sows. All the 16
185 positive sows carried *Y. enterocolitica* bioserotype 4/O:3 positive for the *ail* gene, whereas 10 strains were
186 positive for the *virF* gene. *Yersinia* antibodies were detected in 77.1% of the sows, and according to our
187 Bayesian model, the true seroprevalence was 95,8% (Table 4). The sows that tested positive by culturing and
188 serology originated from 14 and 104 farms, respectively. Antibodies against *Yersinia* were detected from
189 tonsils significantly more frequently than pathogenic *Y. enterocolitica* was detected by culturing (McNemar's
190 test $p < 0.001$).

191

192

Discussion

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194 As expected, the modification of the ELISA test's cut-off activity value (S/P ratio) from 0.2 to 0.3 resulted in a
195 decrease in sensitivity (75,4%; previously 79.5%) and increase in specificity (98,1%; previously 96.9%) (Vilar
196 et al., 2015). According to the manufacturer's instructions, the antigens used in the test are produced by
197 pathogenic *Yersinia* strains only, hence no cross reactions to nonpathogenic species or other enterobacteria
198 should exist. However, there are cross reactions to other pathogenic *Yersinia*, especially
199 *Y. pseudotuberculosis*. Nevertheless, the prevalence of *Y. enterocolitica* in pigs is much higher than the
200 prevalence of *Y. pseudotuberculosis* (Laukkanen et al., 2008; Laukkanen et al., 2009, Ortiz Martínez et al.,
201 2009; Ortiz Martínez et al., 2010; Ortiz Martínez et al., 2011).

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203 The seropositivity of some piglets already at farrowing units is most likely due to colostral
204 maternal antibodies. Seropositivity levels generally increased later than fecal prevalence, which was
205 expected, since antibodies take time to develop (Nielsen et al., 1996). With the exception of farm C, the
206 seropositivity levels were high by the fourth sampling, which indicates a wide transmission of pathogenic
Y. enterocolitica in the pig populations. Pigs from farm C had been moved from an all-in/all-out weaning unit

207 into a continuously filled fattening unit later than pigs from the other farms. This could explain why only
208 some of the pigs were seropositive despite being fecal carriers by the fourth sampling.

209 In the fattening unit of farm C 14 different MLVA types were found. The use of deep peat
210 bedding that can provide a reservoir for pathogenic *Y. enterocolitica* or the high number of pigs per pen
211 might explain the highest amount of detected MLVA types. The pigs were regrouped from six smaller pens
212 into two larger pens, which possibly enhanced the spread of the pathogen. A similar amount ($n = 13$) of
213 MLVA types were found in the fattening unit and slaughterhouse during the second follow-up of farm B (B-
214 II). However, 12 pens (instead of six, as in the other farms) were studied, which may explain this result.
215 MLVA has a relatively high discriminatory power and the results should be interpreted carefully (Virtanen et
216 al., 2013; Alakurtti et al., 2016). The strains from a single farm often had multiple different VNTR loci, and
217 even in the least discriminatory VNTR loci V4 and V9 (Virtanen et al., 2013; Alakurtti et al., 2016) variation
218 was observed between the strains from different farms (Table 3, Supplemental figure), indicating that there
219 were several different strains. Our subsequent studies of farm B reveal that several genotypes might persist
220 – and perhaps dominate – on farms. New MLVA types were found at the slaughterhouse, which could
221 indicate mutations, or more likely cross-contamination during transportation and slaughter processes.

222 Management system is a key factor in controlling the spread of pathogenic *Y. enterocolitica* in
223 pig farms, as highlighted in our study. Once pigs were moved to units where all-in/all-out systems were not
224 used, the infection vigorously spread among the pig population. Earlier studies support our findings, as mixing
225 pigs from different groups has been identified as a risk factor for the spread of pathogenic *Y. enterocolitica*
226 (Virtanen et al., 2012; Virtanen et al., 2014). Other factors such as pig movement, biosecurity level, water
227 source, feed and bedding, have also been associated with the prevalence of pathogenic *Yersinia* in pigs
228 (Skjerve et al., 1998; Nowak et al., 2006; Laukkanen et al., 2009; Virtanen et al., 2011; von Altrock et al., 2011;
229 Novoslavskij et al., 2013; Vilar et al., 2013, Vanantwerpen et al., 2017).

230 According to our prevalence study of tonsil fluid in sows, pathogenic *Y. enterocolitica* is widely
231 spread among sows. However, almost all sows appear to be seropositive only, and just a few sows seem to
232 carry the pathogen in their tonsils – at least to the extent that the pathogen can be isolated by culturing.

233 Sows may have developed immunity against pathogenic *Yersinia*, as also suggested in earlier studies. Vilar et
234 al. (2013) found that only 5% of sows excreted pathogenic *Y. enterocolitica* in feces, while as many as 67%
235 were serologically positive. The true prevalence of *Yersinia* antibodies in serum of Finnish sows was 74%
236 (Vilar et al., 2015).

237 Sows may not be the main source of *Y. enterocolitica* for piglets, because pigs appear to get
238 infected in weaning and fattening units, especially if all-in/all-out management systems are not used.
239 However, sows may still be an important reservoir for pathogenic *Y. enterocolitica*. The high prevalence of
240 antibodies indicates regular contact with the pathogen. Albeit seldom carriers, sows may still sporadically
241 transfer *Y. enterocolitica* to some piglets that later start excreting the pathogen in feces. When piglets get
242 older, levels of maternal antibodies appear to decrease, as indicated in our study. Although pigs primarily get
243 *Y. enterocolitica* in weaning and fattening units, this main reservoir may partly be maintained by sows.
244 Pathogenic *Y. enterocolitica* is rarely isolated from the pig farm environment (Laukkanen-Ninios et al., 2014,
245 Vilar et al., 2013), indicating that the main route of spread is most likely between pigs, specifically their
246 secretions and snout contacts. Nevertheless, it has been possible to maintain pig herds free from
247 *Y. enterocolitica* in closed, specific pathogen-free herds (Nesbakken et al., 2007). To control this foodborne
248 pathogen at the farm level, comprehensive prevention strategies are needed, for example purchasing
249 uninfected animals only, employing all-in/all-out management systems, and avoiding mixing pig groups.

250

251

Conclusions

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253 Pathogenic *Y. enterocolitica* is widely spread in fattening pigs, and several genotypes are found on each farm.
254 The main reservoir for pathogenic *Y. enterocolitica* appears to be fattening pigs, especially if all-in/all-out
255 management systems are not used. Based on the antibodies present in tonsils, essentially all sows have been
256 in contact with this pathogen. Although pathogenic *Y. enterocolitica* is rarely isolated from tonsils of sows,
257 they may nevertheless be an important reservoir for the pathogen, by maintaining it in pig farms.

258 Comprehensive prevention, especially all-in/all-out management systems, are needed to control this
259 foodborne pathogen at the farm level.

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There are no competing financial interests to declare that are related to this publication.

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1 Table 1. Estimates^a for the sensitivity and specificity of the diagnostic test (Pigtype
2 Yopscreen^b, Labor Diagnostik, Leipzig, Germany) used for the detection of *Yersinia* antibodies.

Parameter	Median (95% probability interval)
Sensitivity	75.4 (68.9–81.4)
Specificity	98.1 (90.8–99.9)

3 ^aCalculated from the data originally published by Vilar et al. (2015), but using a revised cut-off
4 activity value (S/P ratio) of 0.3 (instead of 0.2 as used earlier) for the ELISA test according to the
5 instructions by the manufacturer (short protocol).

6 ^bCurrently known as Pigtype *Yersinia* Ab (Qiagen, Leipzig, Germany).

1 Table 2. Prevalence of *Yersinia enterocolitica* 4/O:3 in feces of pigs collected at farrow-to-finish farms and tonsils of pigs collected at abattoirs; prevalence of *Yersinia* antibodies in serum of pigs collected at farrow-to-finish farms.

Farm and method	No. pens	No. pigs per pen ± SD ^a	No. pigs sampled per pen ± SD ^b	Sampling time at farm												Abattoir	2		
				3-6 w			8-10 w			12-15 w			16-20 w					21-26 w	3
				No. positive pens	True prevalence of positive pigs (95% PI) ^c	Unit ^d	No. positive pens	True prevalence of positive pigs (95% PI) ^c	Unit ^d	No. positive pens	True prevalence of positive pigs (95% PI) ^c	Unit ^d	No. positive pens	True prevalence of positive pigs (95% PI) ^c	Unit ^d				
Culturing																	7		
A	6	7.3 ± 1.7	6.7 ± 1.8	0	1.5 (0.1–7.9)	Farr.	6	96.1 (87.8–99.4)	Wean.	6	77.4 (62.7–88.6)	Fatt.	5	16.5 (7.3–29.9)	Fatt.	31 (89.6%, 77.1–96.8)	8		
B-I	6	7.3 ± 1.8	7.2 ± 1.8	0	1.5 (0.1–7.9)	Farr.	0	1.6 (0.1–8.1)	Wean. ^e	6	98.4 (91.9–99.9)	Fatt.	6	96.2 (88.0–99.5)	Fatt.	22 (93.2%, 78.9–99.0)	9		
B-II	12	7.3 ± 0.9	7.0 ± 1.0	0	0.8 (0.0–4.1)	Farr.	0	0.8 (0.0–4.0)	Wean. ^e	12	67.2 (56.6–76.7)	Fatt.	9	22.9 (14.8–32.8)	Fatt.	74 (97.8%, 92.9–99.7)	10		
C	6	7.3 ± 0.7	7.0 ± 0.8	0	1.5 (0.1–7.9)	Farr.	0	1.6 (0.1–8.2)	Wean. ^e	0	1.6 (0.1–8.2)	Wean. ^e	2 ^f	100 (100–100)	Fatt.	18 (96.4%, 82.3–99.9)	11		
Total	30	7.3 ± 1.3	7.0 ± 1.4	0	0.3 (0.0–1.7)		6	19.3 (14.4–24.8)		24	62.3 (55.5–68.8)		26	52.2 (45.4–59.0)		145 (96.3%, 92.4–98.5)	12		
Serology																	13		
A	6	7.3 ± 1.7	6.2 ± 1.8	0	1.7 (0.1–8.4)	Farr.	5	32.3 (19.6–47.0)	Wean.	6	98.2 (90.5–99.9)	Fatt.	6	97.8 (88.9–99.9)	Fatt.		15		
B-I	6	7.3 ± 1.8	6.6 ± 1.6	4	31.5 (19.1–46.1)	Farr.	1	6.6 (1.6–16.9)	Wean. ^e	4	22.6 (11.5–37.3)	Fatt.	no data	no data	Fatt.		16		
B-II	12	7.3 ± 0.9	6.7 ± 1.2	9	57.8 (47.5–67.8)	Farr.	0	0.8 (0.0–4.4)	Wean. ^e	8	53.3 (42.1–64.3)	Fatt.	12	94.1 (87.5–97.9)	Fatt.		17		
C	6	7.3 ± 0.7	5.9 ± 1.8	2	8.1 (2.5–18.2)	Farr.	0	1.8 (0.1–9.0)	Wean. ^e	1	4.8 (0.7–15.0)	Wean. ^e	2 ^f	31.7 (16.5–50.2)	Fatt.		18		
Total	30	7.3 ± 1.3	6.4 ± 1.6	15	31.3 (25.3–37.7)		6	7.8 (4.6–12.1)		19	46.7 (39.6–53.9)		23	83.2 (76.4–88.9)			19		

21 ^aAverage number of pigs per pen at the beginning of the study ± standard deviation.

22 ^bAverage number of pigs sampled per pen during the study ± standard deviation.

23 ^cEstimate for the true prevalence of positive pigs ± 95% probability interval using Bayesian methods as described by Vilar et al. (2015). Note that revised sensitivity and specificity values for the commercial ELISA test were used (Table 1).

24 ^dUnit where the pigs were sampled: Farr. = farrowing unit; Wean. = weaning unit; Fatt. = fattening unit.

25 ^eAll-in/all-out management system was used.

26 ^fIn the fattening unit of farm C, the pigs were regrouped into two larger pens of approximately 20 pigs.

1 Table 3. Genotypic distribution of *Yersinia enterocolitica* 4/O:3 strains isolated from feces and tonsils of pigs in farrow-to-finish farms and abattoirs, based on multiple-locus variable-number
2 tandem repeat analysis (MLVA).

Farm	Sampling time at farm												Abattoir	
	8–10 w				12–15 w				16–20 w				21–26 w	
	MLVA type	No. (%)	Pens ^a	Unit ^b	MLVA type	No. (%)	Pens ^a	Unit ^b	MLVA type	No. (%)	Pens ^a	Unit ^b	MLVA type	No. (%)
A	5-2-6-9-6-3	22 (53.7)	4	Wean.	5-2-6-9-6-3	13 (44.8)	4	Fatt.	5-2-6-9-6-3	3 (50.0)	3	Fatt.	5-2-6-10-7-3	11 (35.5)
	5-2-6-10-7-3	13 (31.7)	3	Wean.	5-2-6-9-7-3	11 (37.9)	4	Fatt.	5-2-6-9-7-3	1 (16.7)	1	Fatt.	5-2-6-9-6-3	9 (29.0)
	5-2-6-9-7-3	5 (12.2)	3	Wean.	5-2-6-10-7-3	3 (10.3)	2	Fatt.	5-2-6-10-7-3	1 (16.7)	1	Fatt.	6-2-6-9-6-3	5 (16.1)
	5-2-6-10-6-3	1 (2.4)	1	Wean.	5-2-6-11-7-3	1 (3.4)	1	Fatt.	5-2-6-10-9-3	1 (16.7)	1	Fatt.	5-2-6-9-7-3	4 (12.9)
					5-2-7-9-6-3	1 (3.4)	1	Fatt.					6-2-6-8-6-3	1 (3.2)
												5-2-6-10-9-3	1 (3.2)	
B-I	SN ^e		Wean. ^c	13-2-7-7-6-3	25 (58.1)	4	Fatt.	13-2-7-7-6-3	25 (59.5)	5	Fatt.	13-2-7-7-6-3	14 (63.6)	
				11-2-7-8-6-3	11 (25.6)	2	Fatt.	11-2-7-8-6-3	11 (26.2)	3	Fatt.	3-6-7-11-13-5	1 (4.5)	
				13-2-5-7-6-3	2 (4.7)	2	Fatt.	13-2-7-7-22-3	2 (4.8)	2	Fatt.	11-2-7-7-6-3	1 (4.5)	
				13-2-8-7-6-3	2 (4.7)	2	Fatt.	11-2-7-8-20-3	1 (2.4)	1	Fatt.	11-2-7-8-6-3	1 (4.5)	
				11-2-7-7-6-3	1 (2.3)	1	Fatt.	12-2-7-7-6-3	1 (2.4)	1	Fatt.	11-2-7-9-6-3	1 (4.5)	
				11-15-7-8-6-3	1 (2.3)	1	Fatt.	12-2-7-8-6-3	1 (2.4)	1	Fatt.	13-2-7-7-14-3	1 (4.5)	
				13-3-7-7-6-3	1 (2.3)	1	Fatt.	13-2-7-7-5-3	1 (2.4)	1	Fatt.	13-2-8-7-6-3	1 (4.5)	
												13-2-7-9-6-3	1 (4.5)	
												13-2-7-9-22-3	1 (4.5)	
B-II	SN ^e		Wean. ^c	11-2-7-7-6-3	30 (55.6)	9	Fatt.	11-2-7-7-6-3	13 (72.2)	5	Fatt.	11-2-7-7-6-3	38 (51.4)	
				9-2-7-7-6-3	9 (16.7)	4	Fatt.	12-2-7-7-6-3	4 (22.2)	3	Fatt.	9-2-7-7-6-3	12 (16.2)	
				12-2-7-7-6-3	3 (5.6)	1	Fatt.	9-2-7-7-6-3	1 (5.6)	1	Fatt.	12-2-7-7-6-3	12 (16.2)	
				10-2-7-7-6-3	2 (3.7)	2	Fatt.					13-2-7-7-6-3	3 (4.1)	
				13-2-7-7-6-3	2 (3.7)	1	Fatt.					9-2-7-7-18-3	1 (1.4)	
				9-2-5-7-6-3	1 (1.9)	1	Fatt.					9-2-7-9-6-3	1 (1.4)	
				9-2-7-7-8-3	1 (1.9)	1	Fatt.					9-4-17-12-12-3	1 (1.4)	
				9-2-7-9-6-3	1 (1.9)	1	Fatt.					11-2-7-8-6-3	1 (1.4)	
				9-2-7-15-8-3	1 (1.9)	1	Fatt.					12-2-7-7-21-3	1 (1.4)	
				9-12-9-4-6-3	1 (1.9)	1	Fatt.					12-2-8-7-6-3	1 (1.4)	
				11-2-3-7-6-3	1 (1.9)	1	Fatt.					12-2-12-23-8-2	1 (1.4)	
				11-2-7-9-6-3	1 (1.9)	1	Fatt.					12-2-12-23-21-2	1 (1.4)	
11-2-9-7-6-3	1 (1.9)	1	Fatt.					13-2-6-7-6-3	1 (1.4)					
C	SN ^e		Wean. ^c	SN ^e			Wean. ^c	3-6-7-11-13-5	11 (27.5)	2 ^d	Fatt.	3-6-7-11-13-5	9 (50.0)	
								3-6-7-11-12-5	9 (22.5)	2	Fatt.	3-6-7-9-8-5	2 (11.1)	
								3-6-8-11-16-5	4 (10.0)	1	Fatt.	3-6-7-9-12-5	2 (11.1)	
								3-6-8-12-16-5	3 (7.5)	2	Fatt.	3-6-7-11-12-5	2 (11.1)	
								3-6-8-11-12-5	2 (5.0)	2	Fatt.	3-6-8-12-16-5	2 (11.1)	
								3-6-8-12-12-5	2 (5.0)	1	Fatt.	3-6-7-9-13-5	1 (5.6)	
								3-6-9-12-16-5	2 (5.0)	1	Fatt.			
								3-6-7-4-2-5	1 (2.5)	1	Fatt.			
								3-6-7-9-12-5	1 (2.5)	1	Fatt.			
								3-6-7-9-13-5	1 (2.5)	1	Fatt.			
								3-6-7-9-18-5	1 (2.5)	1	Fatt.			
								3-6-8-10-12-5	1 (2.5)	1	Fatt.			
								3-6-8-12-2-5	1 (2.5)	1	Fatt.			
								3-6-9-12-12-5	1 (2.5)	1	Fatt.			

3 ^a Number of pens from which the MLVA type was found.

4 ^b Unit where the pigs were sampled: Farr. = farrowing unit; Wean. = weaning unit; Fatt. = fattening unit.

5 ^c All-in/all-out management system was used.

6 ^d In the fattening unit of farm C, the pigs were regrouped into two larger pens of approximately 20 pigs.

7 ^e Samples negative for pathogenic *Y. enterocolitica*.

1 Table 4. Presence of *Yersinia* antibodies and isolation of *Yersinia enterocolitica* 4/O:3 in tonsils ($n = 266$) collected from sows at abattoirs.

Serology	Isolation by culturing		Apparent seroprevalence (95% CI)	True seroprevalence (95% PI) ^a
	No. isolated	No. not isolated		
No. positive	11	194	77.1% (71.7–81.7)	95.8% (87.4–99.7)
No. negative	5	56		

2 ^aEstimate for the true prevalence of positive pigs using Bayesian methods as described by Vilar et al. (2015). Note that revised sensitivity and specificity

3 values for the commercial ELISA test were used (Table 1).