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Van Damme, I.; De Zutter, L.; Jacxsens, L.; Nauta, Maarten

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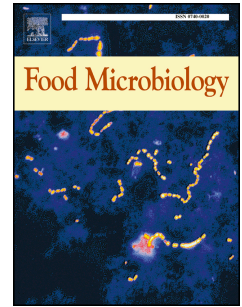
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# Accepted Manuscript

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I. Van Damme, L. De Zutter, L. Jacxsens, M.J. Nauta



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1                   **Control of human pathogenic *Yersinia enterocolitica* in minced meat:**  
2                   **Comparative analysis of different interventions using a risk assessment approach**

3

4 Van Damme I. <sup>a\*</sup>, De Zutter L. <sup>a</sup>, Jacxsens L. <sup>b</sup>, Nauta M.J. <sup>c</sup>

5

6 <sup>a</sup> Department of Veterinary Public Health and Food Safety, Faculty of Veterinary Medicine, Ghent  
7 University, Salisburylaan 133, B-9820 Merelbeke, Belgium ([inge.vandamme@ugent.be](mailto:inge.vandamme@ugent.be);  
8 [lieven.dezutter@ugent.be](mailto:lieven.dezutter@ugent.be))

9 <sup>b</sup> Laboratory of Food Microbiology and Food Preservation, Department of Food Safety and Food  
10 Quality, Faculty of Bioscience Engineering, Ghent University, Coupure Links 653, B-9000 Ghent,  
11 Belgium ([liesbeth.jacxsens@ugent.be](mailto:liesbeth.jacxsens@ugent.be))

12 <sup>c</sup> National Food Institute, Technical University of Denmark, Research Group for Risk-Benefit, Mørkhøj  
13 Bygade 19, DK-2860 Søborg, Denmark ([maana@food.dtu.dk](mailto:maana@food.dtu.dk))

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15 \* Corresponding author: Inge Van Damme; Department of Veterinary Public Health and Food Safety,  
16 Faculty of Veterinary Medicine, Ghent University, Salisburylaan 133, B-9820 Merelbeke, Belgium; e-  
17 mail: [inge.vandamme@ugent.be](mailto:inge.vandamme@ugent.be); Tel.: +32 9 264 73 41

18 **Abstract**

19 This study aimed to evaluate the effect of different processing scenarios along the farm-to-fork chain  
20 on the contamination of minced pork with human pathogenic *Y. enterocolitica*. A modular process  
21 risk model (MPRM) was used to perform the assessment of the concentrations of pathogenic *Y.*  
22 *enterocolitica* in minced meat produced in industrial meat processing plants. The model described  
23 the production of minced pork starting from the contamination of pig carcasses with pathogenic *Y.*  
24 *enterocolitica* just before chilling. The endpoints of the assessment were (i) the proportion of 0.5 kg  
25 minced meat packages that contained pathogenic *Y. enterocolitica* and (ii) the proportion of 0.5 kg  
26 minced meat packages that contained more than  $10^3$  pathogenic *Y. enterocolitica* at the end of  
27 storage, just before consumption of raw pork or preparation. Comparing alternative scenarios to the  
28 baseline model showed that the initial contamination and different decontamination procedures of  
29 carcasses have an important effect on the proportion of highly contaminated minced meat packages  
30 at the end of storage. The addition of pork cheeks and minimal quantities of tonsillar tissue into  
31 minced meat also had a large effect on the endpoint estimate. Finally, storage time and temperature  
32 at consumer level strongly influenced the number of highly contaminated packages.

33 **Keywords:** *Yersinia enterocolitica*, minced meat, risk assessment, pork, interventions

## 34 1. Introduction

35 As pork is the second most consumed meat worldwide (OECD, 2016), an effective control of zoonotic  
36 agents transferred via pork is of major importance to limit the public health risk of zoonotic diseases.  
37 Due to the frequent finding of human pathogenic *Yersinia enterocolitica* in pigs and pork compared  
38 to other food producing animals and food products, and the high genetic relatedness of human and  
39 porcine strains, pork is considered the main source of human pathogenic *Y. enterocolitica*. As such,  
40 77% of *Y. enterocolitica* cases in Europe may be attributed to the consumption of pork (Fosse et al.,  
41 2008). The consumption of raw minced meat may be of particular importance in transmitting  
42 pathogenic *Y. enterocolitica* to humans as Rosner et al. (2012) found that 34% of yersiniosis cases in  
43 Germany had consumed raw minced pork in the seven days preceding illness compared to 12% of  
44 the control group.

45 With 6,471 confirmed cases in 2013, yersiniosis remains the third most commonly reported zoonosis  
46 in the European Union. Over 98% of cases is caused by human pathogenic *Yersinia enterocolitica*  
47 (EFSA and ECDC, 2015), the majority of strains belonging to bioserotype 4/O:3 (EFSA, 2009). The main  
48 reservoirs of these strains are domestic pigs, which can asymptotically carry the pathogens in  
49 lymph nodes, tonsils and the intestinal tract (Laukkanen-Ninios et al., 2014a), resulting in the spread  
50 to the carcass during different steps in the slaughter process (Borch et al., 1996). The presence of  
51 pathogenic *Y. enterocolitica* in the intestines and especially the tonsils is strongly associated with  
52 carcass contamination (Van Damme et al., 2015; Vilar et al., 2015) and carcass contamination has  
53 been shown to differ according to the location on the carcass, with more positive samples found near  
54 the head region and sternum than other areas of the carcass (Laukkanen et al., 2010; Van Damme et  
55 al., 2015).

56 Although the species *Y. enterocolitica* is very heterogeneous, the presence of virulence genes in the  
57 most common types of pathogenic *Y. enterocolitica* seems to be homogeneous (Murros et al., 2016;  
58 Schneeberger et al., 2015). As a result, exposure to these pathogenic types may be more relevant for  
59 public health, rather than specific virulence traits of certain strains. Therefore, identification of the  
60 process steps along the farm-to-fork pathway that have the largest influence on this exposure may  
61 be the most effective way in reducing the public health risk of yersiniosis, prospecting the  
62 development of targeted control measures. Quantitative microbial risk assessment (QMRA) has  
63 emerged in the area of food safety as a comprehensive and systematic approach for addressing the  
64 risk of microbial hazards in the food chain and can be used to assess the impact of control strategies  
65 or interventions (Havelaar et al., 2008; Møller et al., 2015). Using the Modular Process Risk Model  
66 (MPRM) methodology as proposed by Nauta (2008), the food production pathway is described by  
67 subdividing the chain in different modules that each represent a basic process. These basic processes

68 include microbial (growth or inactivation) and food handling processes (cross-contamination,  
69 removal, partitioning and mixing), by which the changes in prevalence, concentration and unit can be  
70 modelled. The output of one module then serves as the input for the following module. This  
71 structured approach allows a structured analysis of the food chain, which gives new insights in the  
72 complex process of food production and can identify crucial data gaps.

73 The objective of this study was to model the spread of pathogenic *Y. enterocolitica* contamination  
74 during the production of minced meat and to evaluate the effect of different intervention scenarios  
75 during minced meat production on human exposure via raw minced pork. Therefore, a food chain  
76 modelling approach was applied to assess the exposure of human pathogenic *Y. enterocolitica*  
77 through industrially produced minced meat using the MPRM methodology. First a baseline model  
78 was built describing the current processing practices and changes in prevalence and concentrations  
79 during the process. Next, alternative scenarios were defined to evaluate the effects of potential  
80 interventions. As, to our knowledge, there is no dose response model available for *Y. enterocolitica*  
81 and no accurate data on raw minced meat consumption could be found, the endpoint of the  
82 assessment was not the exposure or the health risk but (A) the proportion of contaminated 0.5 kg  
83 minced meat packages with pathogenic *Y. enterocolitica* and (B) the proportion of 0.5 kg minced  
84 meat packages that contained more than  $10^3$  pathogenic *Y. enterocolitica* at the end of storage, just  
85 before consumption of raw minced pork or preparation. To identify the most important data gaps,  
86 uncertainties were studied by comparative scenario analyses.

## 87 **2. Material and Methods**

### 88 *2.1. Description of the food pathway and model implementation*

89 An overview of the pathway used in the model is shown in Figure 1. A general overview of the model  
90 and a detailed description of the distributions and parameters used are shown in Table 1 and 2,  
91 respectively.

92 The entire model was simulated with Monte Carlo techniques (100,000 iterations) using @Risk  
93 software (version 7.5.0., Palisade Corporation, Newfield, NY, US). By the lack of a health risk estimate,  
94 the alternative main outputs of the model were point estimates of the prevalence (proportion of 0.5-  
95 kg packages containing one or more pathogenic *Y. enterocolitica*) and/or the proportion of highly  
96 contaminated minced meat packages (containing  $> 10^3$  pathogenic *Y. enterocolitica* per 0.5-kg  
97 package). To evaluate the effect of alternative scenarios, the value of one or more model parameters  
98 was changed and the corresponding endpoint estimate was compared to that of the baseline  
99 scenario. Different scenarios were compared by calculating the  $\log_{10}$  of the relative proportions (the

100 quotient of the endpoint estimate of an alternative scenario and the endpoint estimate of the  
101 baseline scenario), as e.g. in Møller et al. (2015).

## 102 2.2. The baseline model

### 103 2.2.1. Input data - initial contamination of carcasses

104 The prevalence and concentration of human pathogenic *Y. enterocolitica* on pig carcasses were used  
105 as input for the model and were based on the results of a Belgian study describing the contamination  
106 of pork carcasses with pathogenic *Y. enterocolitica* after evisceration before cooling (Van Damme et  
107 al., 2015). The study detected *Y. enterocolitica* bioserotype 4/O:3 on the sternal region (breast cut  
108 and surrounding skin) of 16.4% of the carcasses, which was the value used as the initial prevalence of  
109 carcasses ( $P_{\text{initial}}$ ). Quantitative and semi-quantitative concentration data of pathogenic *Y.*  
110 *enterocolitica* at the sternal region were obtained by analysing different subsamples with different  
111 isolation methods. The R package “fitdistrplus” was used to fit a normal distribution to the censored  
112 data using the “fitdistcens” function (Pouillot and Delignette-Muller, 2010). The resulting normal  
113 distribution of the *Y. enterocolitica* concentration on pork carcasses was used as input for the model  
114 ( $C_{\text{initial}} \sim \text{Normal}(-2.565; 0.736)$  in  $\log_{10}$  CFU/cm<sup>2</sup>, with  $\sim$  meaning that it is a random sample from the  
115 distribution). As  $P_{\text{initial}}$  was based on the combined results of different detection methods from which  
116 the  $C_{\text{initial}}$  distribution was derived, the distribution was truncated at a minimum value of  $-1.85 \log_{10}$   
117 CFU/cm<sup>2</sup>, which was the limit of detection of the most sensitive detection method. The final  
118 (truncated) distribution had a mean of  $-1.46 \log_{10}$  CFU/cm<sup>2</sup> and standard deviation of 0.33.

### 119 2.2.2. Inactivation and growth during carcass chilling and cold storage

120 Blast chilling, during which the carcass surface is frozen, was considered to cause a  $0.6 \log_{10}$   
121 reduction in pathogenic *Y. enterocolitica* concentrations ( $I_{\text{cc}}$ ), according to data of King et al. (2012)  
122 who evaluated the effect of freezing on *Y. enterocolitica* numbers on pig organs. When the  
123 concentration after inactivation ( $N_{\text{cci}}$ ) was below 1 CFU/2000 cm<sup>2</sup>, the carcass was considered to be  
124 pathogenic *Y. enterocolitica* negative and growth after the blast chilling step was not allowed in the  
125 model.

126 After inactivation during blast chilling, *Y. enterocolitica* was assumed to grow during conventional air  
127 chilling and cold storage of carcasses at 4°C. The doubling time for the growth model during carcass  
128 cold storage ( $D_{\text{ccg}}$ ) was set at 10.0 h, based on ComBase Predictor results (<http://combase.cc>) using a  
129 pH of 5.8,  $A_w$  value of 0.997, and temperature of 4°C as input values. The lag phase ( $\lambda_{\text{ccg}}$ ) for the  
130 growth model was set at 24h and the maximum growth was never allowed to result in  
131 concentrations higher than  $7 \log_{10}$  CFU/cm<sup>2</sup> (van Netten et al., 1997). Carcasses from pigs that were  
132 slaughtered on Mondays to Thursdays were assumed to be processed the next day and pigs

133 slaughtered on Fridays were processed on Monday, resulting in a cold storage time ( $Time_{ccg}$ ) of  
 134 respectively 20h and 68h in 80% and 20% of the iterations. The concentration of pathogenic *Y.*  
 135 *enterocolitica* on carcasses after growth during cold storage,  $N_{ccg}$ , was determined:

$$N_{ccg} = N_{cci} \times 2^{\frac{Time_{ccg} - \lambda_{ccg}}{D_{ccg}}}$$

136 When  $\lambda_{ccg}$  was higher than  $Time_{ccg}$ , no growth was allowed, so  $N_{ccg}$  was equal to the number of CFU  
 137 after blast chilling ( $N_{cci}$ ).

### 138 2.2.3. Cutting, derinding, grinding and packaging at the meat processing plant

139 The model for grinding was based on practices of a representative large minced meat producing  
 140 company in Belgium. In the baseline model, a batch consisted of 900 kg minced meat and contained  
 141 34% pork bellies (weight/weight percent, w:w). The remaining ingredients (which may be beef, eggs,  
 142 herbs, and/or other pork cuts) were assumed to have no contribution to contamination with  
 143 pathogenic *Y. enterocolitica*. Although other pork cuts, such as shoulder cuts, are also frequently  
 144 used for the production of minced meat, the contribution of these cuts was not included in the  
 145 model due to the lack of sufficient reliable data. The baseline model thus assumed that bellies were  
 146 the sole source of pathogenic *Y. enterocolitica* contamination.

147 The number of pathogenic *Y. enterocolitica* on a contaminated belly cut ( $N_{bc}$ ) was determined using  
 148 the number of CFU on the carcass after growth during cold storage and assuming a total surface of  
 149 2000 cm<sup>2</sup> (approximately 20 cm x 50 cm on both sides). After derinding, the baseline model assumed  
 150 that half of the bacteria were removed. The prevalence of pathogenic *Y. enterocolitica* on belly cuts  
 151 was assumed to be the same as the initial contamination of carcasses ( $P_{bdr} = P_{initial}$ ).

152 Assuming a weight of pork bellies of  $W_{bc} = 7.5$  kg each, the number of pork bellies within one batch  
 153 was calculated ( $n_{bb}$ ). The number of pathogenic *Y. enterocolitica* contaminated pork bellies per batch  
 154 was determined using  $n_{pbb} \sim \text{Binomial}(n_{bb}; P_{bdr})$ . The total number of bacteria per contaminated pork  
 155 belly ( $N_{bdr,i}$ ) was simulated for each positive belly  $i$  ( $i = 1.. n_{pbb}$ ) included in the batch (taking a random  
 156 sampling from  $C_{initial}$  for each positive belly). All bellies that were used within one batch of minced  
 157 meat were assumed to originate from pigs slaughtered on the same day, so the time between  
 158 slaughter and cooling ( $Time_{ccg}$ ) remained constant for all bellies within the same batch. The numbers  
 159 of pathogenic *Y. enterocolitica* on each of the positive bellies were added to determine the total  
 160 number of pathogenic *Y. enterocolitica* in a batch of minced meat ( $N_{mb}$ ):

$$N_{mb} = \sum_{i=1}^{n_{pbb}} N_{bdr,i}$$



161 The weight of individual minced meat packages ( $W_{mp}$ ) was assumed to be 0.5 kg. Pathogenic *Y. enterocolitica* were assumed to be homogeneously distributed in a batch to calculate the number of  
 162 pathogenic *Y. enterocolitica* in one 0.5-kg minced meat package ( $N_{mp}$ ) (Nauta, 2005).  
 163

#### 164 2.2.4. Storage at the meat processing plant, retail and consumer level

165 As there is no specific secondary growth model available for pathogenic *Y. enterocolitica* in minced  
 166 meat at different temperatures, the growth at retail and consumer level was modelled using  
 167 ComBase data ([www.combase.cc](http://www.combase.cc)). Hereby, the maximum growth rate (in  $\log_{10}$  CFU/h) was  
 168 determined for temperatures varying between 0 and 15°C (using 1°C steps) for a pH of 5.8 and NaCl  
 169 concentration of 1%. The percentage of CO<sub>2</sub> was set at 30% to represent MAP packaging. Fitting a  
 170 regression line through the temperature – growth rate values obtained ( $R^2 = 0.9992$ ), resulted in an  
 171 equation that was used to calculate  $\mu_{max}$  according to the temperature (Table 2).

172 To represent storage in the meat processing plant, transport and retail, the temperature ( $Temp_{rg}$ )  
 173 and time ( $Time_{rg}$ ) was set at 4°C and 24h, respectively. To represent storage at consumer level, the  
 174 temperature ( $Temp_{cg}$ ) was based on data from the Belgian Food Consumption Survey of 2004, in  
 175 which the temperature of home refrigerators was determined (Devriese et al., 2006), resulting in a  
 176 Pert distribution defined by the quartiles, 5, 7 and 9°C. Pathogenic *Y. enterocolitica* were considered  
 177 not to grow below 0°C. The time during which minced meat was stored ( $Time_{cg}$ ) was based on results  
 178 of Swedish consumers (Marklinder et al., 2004), resulting in a Pert distribution with most likely one  
 179 day, a minimum of zero and maximum of four days. The final number of pathogenic *Y. enterocolitica*  
 180 in 0.5-kg minced meat packages just before consumption/preparation was calculated as

$$N_{cg} = N_{mp} \times 10^{(\mu_{max,cg} \times Time_{cg} + \mu_{max,rg} \times Time_{rg})}$$

181 The endpoint estimates were the proportion of 0.5-kg minced meat packages that contained  $\geq 1$   
 182 pathogenic *Y. enterocolitica* and the proportion of packages that contained  $\geq 1000$  pathogenic *Y.*  
 183 *enterocolitica* per 0.5-kg minced meat package.

### 184 2.3. Alternative scenarios

185 Alternative scenarios of the model were run and compared to the baseline model. Some of these  
 186 alternative scenarios represent realistic modifications of processing, which can for example be  
 187 implemented as interventions (2.3.1 – 2.3.3). Other alternative scenarios are evaluated in an  
 188 uncertainty analysis, to study the uncertainty attending parameter values and model assumptions  
 189 (2.3.4; as e.g. in Nauta et al. (2007)). An overview of the different parameters that were modified to  
 190 evaluate alternative scenarios is shown in Tables 3 to 6.

#### 191 2.3.1. Initial contamination, chilling and decontamination procedures of carcasses

192 Alternative scenarios for initial carcass contamination were analysed using a prevalence ( $P_{\text{initial}}$ ) of 7.5%  
193 and 37.5% and concentrations ( $C_{\text{initial}}$ ) that had a mean concentration of 0.5  $\log_{10}$  lower or higher than  
194 in the baseline model, to represent the 'best' and 'worst' slaughterhouses regarding pathogenic *Y.*  
195 *enterocolitica* contamination, respectively (Van Damme et al., 2015). Six different scenarios were  
196 evaluated: a lower prevalence (7.5%) but baseline concentrations (scenario A1); a lower  
197 concentration but baseline prevalence (scenario A2); a lower prevalence and a lower concentration  
198 (scenario A3); a higher prevalence but baseline concentrations (scenario A4); a higher concentration  
199 but baseline prevalence (scenario A5); and a higher prevalence and higher concentration (scenario  
200 A6).

201 To simulate a slaughterhouse that only applied conventional air chilling (no prior blast chilling;  
202 scenario A7), a 0.1  $\log_{10}$  reduction during chilling was assumed ( $I_{\text{cc}}$ ), which is based on the mean  
203 reduction of *Y. enterocolitica* after chilling of pig organs to a an internal temperature of 4°C (King et  
204 al., 2012). The use of steam condensation was evaluated based on the reductions observed by  
205 Smulders et al. (2012) when applying steam of 65°C for 18 s on pork skin, and was followed by a  
206 reduction to simulate either conventional chilling (scenario A8) or blast chilling (scenario A9).

207 The effect of applying lactic acid treatment (2% for 10 s at 40-50°C), combined with blast chilling or  
208 conventional air chilling, was simulated using a reduction of 0.7 and 1.6, respectively (King et al.,  
209 2012) (scenario A10 and A11). The reduced growth during carcass cold storage after lactic acid  
210 treatment was simulated using a lag phase ( $\lambda_{\text{ccg}}$ ) of 48h and doubling time ( $D_{\text{ccg}}$ ) of 12.4h based on  
211 results of van Netten et al. (1997), after applying 2% lactic acid (at 37°C for 120s) on pork skin.

212 The cold storage time of carcasses ( $\text{Time}_{\text{ccg}}$ ) was set at either 68h or 20h to represent the production  
213 of minced meat on Monday (from carcasses slaughtered on Friday; scenario A12) or minced meat  
214 produced on Tuesday-Friday (from carcasses slaughtered on Monday-Thursday; scenario A13).

### 215 2.3.2. Addition of head meat and tonsillar tissue during grinding and batch size effect

216 The effect of the inclusion of head meat for the production of minced meat was simulated at  
217 different levels (1%, 10%, and 50% w:w; scenarios B1, B2, and B3, respectively). As input data,  
218 prevalence and count data of human pathogenic *Y. enterocolitica* on the mandibular region of  
219 carcasses before chilling were obtained from Van Damme et al. (2015). A distribution was fitted  
220 through the censored count data (see 2.2.1), resulting in a lognormal distribution for  $C_{\text{initial},m}$  with a  
221 mean of -0.578 and standard deviation of 1.26  $\log_{10}$  CFU/100cm<sup>2</sup>. The distribution was truncated at  
222 0.15  $\log_{10}$  CFU/100cm<sup>2</sup> (the lower limit of the most sensitive isolation method), yielding a new  
223 distribution with a mean of 0.93  $\log_{10}$  CFU/100cm<sup>2</sup> and standard deviation of 0.64. All pathogenic *Y.*  
224 *enterocolitica* on one head meat cut were assumed to originate from the carcass at the surface (100

225 cm<sup>2</sup>) of the mandibular region. The same steps during the chilling and cold storage of carcasses were  
226 applied as for the sternal region. Carcasses containing less than 0 log<sub>10</sub> CFU/100 cm<sup>2</sup> after blast  
227 chilling (C<sub>mci</sub>) were considered negative. The number of pathogenic *Y. enterocolitica* positive head  
228 meat cuts per batch (n<sub>phb</sub>) was calculated similar to the pork bellies, assuming a weight of an  
229 individual cheek of 75 g (W<sub>hm</sub>), and a prevalence of 28.9% (P<sub>initial,m</sub>). The number of cfu per head meat  
230 cut was simulated for each positive cut separately, starting each time from C<sub>initial,m</sub>. The numbers of  
231 pathogenic *Y. enterocolitica* on positive head meat cuts were added to the numbers on pork bellies  
232 to determine the total number of pathogenic *Y. enterocolitica* per batch of minced meat (N<sub>mb</sub>).

233 The addition of tonsillar tissue (scenarios B4-B6) was simulated using a prevalence (P<sub>initial,t</sub>) of  
234 pathogenic *Y. enterocolitica* in pig tonsils during slaughter of 44.3% and an initial concentration  
235 (C<sub>initial,t</sub>) with a minimum of 1.00 log<sub>10</sub> CFU/g, most likely of 4.00 log<sub>10</sub> CFU/g and a maximum of 5.91  
236 log<sub>10</sub> CFU/g (Van Damme et al., 2015). Inactivation and growth during carcass chilling and cold  
237 storage was included as described before. Numbers were modelled for each individual positive tonsil  
238 and were added to the total number of pathogenic *Y. enterocolitica* from pork bellies to calculate the  
239 total number of pathogenic *Y. enterocolitica* per batch of minced meat (N<sub>mb</sub>). As alternative scenarios,  
240 we evaluated the addition of one piece of tonsillar tissue of 1 g (scenario B4), one piece of tonsillar  
241 tissue of 10 g (scenario B5), and 10 pieces of tonsillar tissue (of 10 different pigs) of 1 g each  
242 (scenario B6).

243 Besides a batch weight of 900 kg in the baseline scenario, the effect of smaller and larger minced  
244 meat batches were simulated by changing W<sub>b</sub> to 140 kg and 1500 kg in the alternative scenarios B7  
245 and B8, respectively.

### 246 2.3.3. Consumer storage practices

247 Alternative scenarios for consumer storage (C1-C9) were evaluated by replacing Temp<sub>cg</sub> or Time<sub>cg</sub> by  
248 different fixed values (4°C, 7°C, 10°C and 15°C for Temp<sub>cg</sub> and 0, 1, 2, 3, and 4 days for Time<sub>cg</sub>). The  
249 effect of consumer storage scenarios was evaluated for both MAP packaging and storage under  
250 ambient atmosphere. Storage under ambient atmosphere was simulated by changing the formulas  
251 for μ<sub>max</sub> both at retail and consumer level (Table 5). The formula was created using ComBase data as  
252 described before, but omitting the parameter "CO<sub>2</sub>".

253 For simulation of MAP packages that are consumed at the use-by date (scenarios C10 and C11), a  
254 shelf-life of 9 days was assumed based on company information. Storage of minced meat at  
255 consumer level until the use-by date (scenario C10) was simulated setting the storage time at  
256 consumer level at 7 days. For simulation of MAP packages that are sold and consumed/prepared at

257 the use-by date (scenario C11), the storage time at retail ( $\text{Time}_{\text{rg}}$ ) was set at 9 days and storage time  
258 at consumer level ( $\text{Time}_{\text{cg}}$ ) was set at 0 days.

#### 259 2.3.4. Uncertainty analysis

260 Uncertainty analyses were performed by estimating the prevalence and proportion of packages  
261 containing more than 3  $\log_{10}$  CFU by changing one parameter value in the model to a value that  
262 represents the low or high end of the uncertainty interval around the value chosen in the baseline  
263 model. The parameter values that were changed are shown in Table 6.

264 The uncertainty regarding the initial concentration on carcasses ( $C_{\text{initial}}$ ) was evaluated by changing  
265 the mean or standard deviation with  $\pm 0.5 \log_{10}$  (U1-U4). For the prevalence ( $P_{\text{initial}}$ ), the upper (U5)  
266 and lower limit (U6) of the 95% confidence interval for the prevalence at the sternal region were  
267 used (Van Damme et al. 2015). A different value for the reduction during blast chilling ( $I_{\text{cc}}$ ) was based  
268 on the 7% cell inactivation that was observed by El-Zawahry and Grecz (1981) when freezing  
269 pathogenic *Y. enterocolitica* in broth at  $-18^{\circ}\text{C}$  for one hour (U7). A larger reduction during blast  
270 chilling (U8) was simulated using the  $-0.8 \log$  reduction of *Y. enterocolitica* that was observed by King  
271 et al. (2012) when applying a water wash before freezing pig organs. Scenario U9 assumed no growth  
272 of pathogenic *Y. enterocolitica* during carcass cold storage, which was based on the results of Greer  
273 and Dilts (1995), who found no growth of pathogenic *Y. enterocolitica* O:4,32 during storage at  $4^{\circ}\text{C}$   
274 for over ten days after artificial inoculation of lean pork tissue. As Greer and Dilts (1995) observed  
275 immediate growth of *Y. enterocolitica* O:4,32 on pork fat at  $4^{\circ}\text{C}$ , a lag phase of 0 hours was assumed  
276 in scenario U10. The doubling time in scenario U10 was based on ComBase results assuming a  
277 temperature of  $4^{\circ}\text{C}$ , pH of 6.5 (Greer and Dilts, 1995), and  $A_w$  of 0.990 (van Netten et al., 1997). The  
278 percentage of pathogenic *Y. enterocolitica* that remain on a belly cut after derinding was set at 25%  
279 and 75% to represent less and more removal during cutting and removal (U11 and U12). The lower  
280 and upper limits of the uncertainty about the weight of a batch of minced meat ( $W_b$ ), the proportion  
281 of bellies that is used (%bellies), the weight of a belly cut ( $W_{\text{bdr}}$ ), the temperature ( $\text{Temp}_{\text{rg}}$ ) and the  
282 time during storage at retail ( $\text{Time}_{\text{rg}}$ ) were considered reasonable by the authors (U13-U22). The  
283 uncertainty regarding the growth of pathogenic *Y. enterocolitica* in minced meat was studied by  
284 reducing the maximum growth rate by half (U23).

### 285 **3. Results**

286 Using the baseline scenario, the prevalence of pathogenic *Y. enterocolitica* in 0.5-kg minced meat  
287 packages was estimated at 15.4% ( $\geq 1$  CFU/package). Only a small percentage of packages (1.4%, i.e.  
288 9.2% of the contaminated packages) contained more than  $10^3$  pathogenic *Y. enterocolitica* at the end

289 of storage. The distribution of pathogenic *Y. enterocolitica* in positive minced meat packages at the  
290 end of storage (just before consumption or preparation) in the baseline scenario is shown in Figure 2.

### 291 3.1. Initial contamination of carcasses before chilling

292 The effect of initial carcass contamination on pathogenic *Y. enterocolitica* contaminated minced meat  
293 packages was evaluated varying the initial prevalence and concentration of pathogenic *Y.*  
294 *enterocolitica* on carcasses ( $P_{\text{initial}}$  and  $C_{\text{initial}}$ ) to represent minced meat that is produced using  
295 carcasses from slaughterhouses with either low or high contamination with pathogenic *Y.*  
296 *enterocolitica*. Lowering the prevalence of pathogenic *Y. enterocolitica* on carcasses from 16.39% to  
297 7.5% reduced the proportion of highly contaminated meat packages by half (Figure 3). A similar  
298 reduction was seen when the average initial concentration on pork carcasses is reduced by  $0.5 \log_{10}$   
299 CFU/cm<sup>2</sup>. The combined effect of reducing the prevalence and the concentration resulted in the  
300 highest effect, with a more than 5-fold decrease in the number of highly contaminated packages  
301 before consumption. A similar but opposite effect was seen for a higher prevalence and/or higher  
302 concentration (Figure 3).

### 303 3.2. Effect of decontamination

304 The results of different scenarios to evaluate the effect of decontamination methods for carcasses at  
305 slaughterhouse level are shown in Figure 4. The use of solely conventional chilling resulted in twice  
306 as many pathogenic *Y. enterocolitica* contaminated minced meat packages compared to when it's  
307 combined with blast chilling, during which the carcass surface is frozen. Steam condensation had a  
308 larger effect on the final outcome estimates as it would reduce the number of contaminated and  
309 highly contaminated pathogenic *Y. enterocolitica* packages 95 to 158 times. The use of 2% lactic acid  
310 sprays would also reduce the proportion of pathogenic *Y. enterocolitica* contaminated minced meat  
311 packages, resulting in a larger effect in combination with blast chilling than with conventional air  
312 chilling. Using carcasses that are chilled for 68 h resulted in more than 10 times as many pathogenic *Y.*  
313 *enterocolitica* contaminated 0.5-kg minced packages compared to minced meat that is produced  
314 using 20h-chilled carcasses (Figure 4).

### 315 3.3. Addition of head meat and tonsillar tissue

316 The additional use of 1% to 50% head meat for the production of minced meat increased the  
317 proportion of pathogenic *Y. enterocolitica* positive minced meat packages 2 to 6 times compared to  
318 the baseline scenario that only assumed pork bellies as a source of pathogenic *Y. enterocolitica*  
319 contamination (Figure 5). The impact of adding head meat was larger for highly contaminated  
320 packages than for the prevalence of pathogenic *Y. enterocolitica* positive minced meat packages. The

321 use of 10% head meat in minced meat resulted in almost 20 times as many highly contaminated  
322 minced meat packages at time of consumption (Figure 5).

323 The addition of 1 g tonsillar tissue to a 900-kg minced meat batch resulted in a 7-fold increase of the  
324 number of minced meat packages containing  $>3$  log pathogenic *Y. enterocolitica* at time of  
325 consumption (Figure 5 and Figure 2). The addition of one tonsil of 10 g resulted in a similar but  
326 slightly higher increase. The addition of 1-g tonsil pieces of 10 different pigs resulted in over 35 times  
327 as many highly contaminated minced meat packages at time of consumption (Figure 5).

328 Changing the batch size ( $W_b$ ) from 900 kg to 140 kg or 1500 kg had very little effect on the endpoint  
329 estimates (data not shown).

#### 330 3.4. Consumer storage

331 When storage of minced meat at consumer level would always be at 4°C, the proportion of highly  
332 contaminated packages would be reduced with more than a 1000-fold compared to the baseline  
333 scenario (Figure 6). If minced meat would always be consumed or prepared within one day after  
334 purchase, a reduction of the endpoint estimate was observed, whereas a constant storage time of  
335 two or more days increased the proportion of highly contaminated packages compared to the  
336 baseline scenario. For each of the scenarios, storage at ambient atmosphere resulted in a higher  
337 proportion of highly contaminated packages than storage in MAP (Figure 6). Storage of minced meat  
338 until the use-by date was simulated using a storage time at consumer level of 7 days or storage at  
339 retail for 9 days (to simulate purchase and consumption at the end of shelf life). Both scenarios  
340 estimated that nearly all pathogenic *Y. enterocolitica* positive packages after packaging (15%) would  
341 contain  $> 10^3$  pathogenic *Y. enterocolitica* at the end of the 9-day storage period. The endpoint  
342 estimate was higher when packages were stored until the use-by date in MAP, as compared to  
343 storage at ambient atmosphere for two days or less (Figure 6).

#### 344 3.5. Uncertainty

345 The results for the uncertainty analyses are shown in Figure 7. A reduced growth rate during storage  
346 at retail and consumer level had the highest impact on the proportion of highly contaminated minced  
347 meat packages. Uncertainty regarding the standard deviation of pathogenic *Y. enterocolitica*  
348 numbers on carcasses before chilling ( $C_{initial}$ ), reduction during blast chilling, and growth during  
349 carcass cold storage had a large effect on both endpoint estimates. For all variables that were  
350 evaluated, the uncertainty had a larger effect on the proportion of highly contaminated packages  
351 than on the prevalence of pathogenic *Y. enterocolitica* in minced meat packages. The uncertainty

352 during minced meat production regarding the exact weight of a minced meat batch, the proportion  
353 of bellies and the weight of a pork belly had only a minor effect on the endpoint estimates.

#### 354 4. Discussion

##### 355 4.1. Modelling approach

356 The consumption of raw minced pork has been shown to be the main risk factor for yersiniosis  
357 infections in Germany (Rosner et al., 2012) and the knowledge of consumers regarding the correct  
358 handling of raw minced meat seems to be limited (Bremer et al., 2005). Therefore, the effect of  
359 different control measures during the production of minced meat on pathogenic *Y. enterocolitica*  
360 contaminated and highly contaminated minced meat packages were evaluated in this study. The  
361 modelling approach used was based on the Modular Process Risk Model approach (Nauta, 2008) that  
362 has frequently been applied to model the transmission of microbial pathogens through food chains  
363 for quantitative microbiological risk assessment (e.g. Nauta et al., 2007; Daelman et al., 2013; Møller  
364 et al., 2015). A full risk assessment, ending at an estimation of the risk of illness, was not feasible as  
365 only few reports are available estimating the numbers of *Yersinia* spp. in food products that are  
366 related to yersiniosis cases (Pärn et al., 2015; Todd et al., 2008) and, to our knowledge, no dose-  
367 response model is available for pathogenic *Y. enterocolitica*. Moreover, due to a lack on consumption  
368 data of raw minced pork and uncertainty about preparation styles, it was decided to end the analysis  
369 at the end of storage, just before consumption of raw minced pork or preparation. Using a similar  
370 approach as Nauta et al. (2003), and acknowledging that all microbial dose response models show an  
371 increasing probability of illness with an increasing dose, it was assumed that every contaminated  
372 package may pose a health risk and that the risk of yersiniosis is higher for highly contaminated  
373 packages. The choice of the critical level  $10^3$  was arbitrary, balancing the need for a high level with  
374 the need for a level that occurs regularly, as to get robust results with a feasible number of model  
375 iterations. When comparing two scenarios, it is assumed that the relative proportion of highly  
376 contaminated packages can be considered a reasonable surrogate for the relative risk as applied  
377 elsewhere (e.g. Møller et al., 2015).

##### 378 4.2. Uncertainties of the model and relevant data gaps

379 The present model used pathogenic *Y. enterocolitica* numbers that are found on the sternal region of  
380 carcasses as input variables to represent contamination of the belly area, and assumed that pork  
381 bellies were the sole source of contamination of minced meat. Laukkanen-Ninios et al. (2014b)  
382 quantified plasmid-carrying *Yersinia* in meat cuts in Finland that were intended to be used in minced  
383 meat and found *Yersinia* in 39% of pork cuts, varying between 0.1 and 1.6 MPN/g (average 0.41  
384 MPN/g) using nested PCR. Nevertheless, as pathogenic *Y. enterocolitica* were isolated from one pork

385 cut only (0.6%) (Laukkanen-Ninios et al., 2014b), the contamination level of meat cuts for the  
386 production of minced meat seems very low. Nevertheless, since contamination from shoulder cuts  
387 and cross contamination between belly cuts were not included in the present model, the  
388 contamination of meat cuts with pathogenic *Y. enterocolitica* before grinding is probably  
389 underestimated. Moreover, the uncertainty analysis showed that the standard deviation of the initial  
390 concentration on carcasses had a large effect on the final prevalence of contaminated packages and  
391 especially for the proportion of highly contaminated packages. This importance of the standard  
392 deviation of concentrations has been found previously (Duarte et al., 2016). Clearly, more accurate  
393 estimations on the numbers of pathogenic *Y. enterocolitica* on bellies and other pork cuts that are  
394 used for minced meat production, including the variation between carcasses and slaughterhouses,  
395 could improve the estimations of the model.

396 The level of growth and inactivation of *Y. enterocolitica* has been shown to differ according to the  
397 tissue. As such, Greer and Dilts (1995) observed immediate growth of pathogenic *Y. enterocolitica* at  
398 4°C after artificial inoculation of fat tissue whereas no growth was observed on lean tissue for several  
399 days after inoculation. The authors also found that pathogenic *Y. enterocolitica* on lean tissue were  
400 more resistant to lactic acid than those on fat tissue (Greer and Dilts, 1995). Moreover, larger  
401 reductions of *Y. enterocolitica* have been observed on pig skin compared to muscle tissue when  
402 evaluating steam-ultrasound decontamination (Morild et al., 2011) or water spraying followed by  
403 steam decontamination (Smulders et al., 2012). Nevertheless, the effect of lactic acid treatment has  
404 been shown to vary between studies. As such, van Netten et al. (1997) found a 4.7 log immediate  
405 death of *Y. enterocolitica* serotype O:3 on pork skin after dipping in 2% lactic acid at 37°C for 120s.  
406 Such reductions would reduce the proportion of highly contaminated packages with more than a  
407 1000-fold (data not shown), though this is likely an overestimation of the reduction as such  
408 conditions may not be accomplished under field conditions. Besides the immediate effect of lactic  
409 acid, the present model assumed a reduced growth of *Y. enterocolitica* during carcass cold storage  
410 after the application of 2% lactic acid, which are based on data using pork skin (van Netten et al.,  
411 1997). Nevertheless, Greer and Dilts (1995) observed a persistent reduction of *Y. enterocolitica* in the  
412 next seven days following a 3% lactic acid treatment of pig lean and fat tissue stored at 4°C.  
413 Therefore, studies quantifying the immediate and long-term effect of lactic acid on carcasses under  
414 field conditions are necessary to improve the predictions for lactic acid decontamination. As the  
415 attachment, inactivation, and growth of pathogenic *Y. enterocolitica* may differ according to the  
416 surface type (Greer and Dilts, 1995; Morild et al., 2011), the inclusion of these differences would be a  
417 more realistic approach to model *Y. enterocolitica* on carcasses, but this would considerably increase  
418 the complexity of the model. Moreover, this would require comprehensive data on the distribution,



419 growth and inactivation of the pathogens on each of the different tissues on carcasses, which are  
420 currently not available. Nevertheless, as the level of growth and inactivation of *Y. enterocolitica*  
421 during cold storage may have a large influence on the outcome variables, more accurate studies on  
422 the level of reduction of pathogenic *Y. enterocolitica* on carcasses under different chilling and cold  
423 storage conditions - including the biological and strain variation - should be performed to obtain  
424 more accurate endpoint estimates.

425 Data regarding the growth of pathogenic *Y. enterocolitica* on pork, and minced meat in particular, are  
426 limited. Therefore, the growth rate represented a large uncertainty in the present model. Kleinlein  
427 and Untermann (1990) observed growth of pathogenic *Y. enterocolitica* in minced beef stored in  
428 MAP (20% CO<sub>2</sub>, 80% O<sub>2</sub>), especially at temperatures of 10°C or higher, whereas Strotmann et al.  
429 (2008) observed a reduction of *Y. enterocolitica* bioserotype 4/O:3 during storage at 2°C, regardless  
430 of the CO<sub>2</sub> concentration. After 13 days of storage of pig cheeks at 6°C in 30% CO<sub>2</sub> and 70% O<sub>2</sub>,  
431 Fredriksson-Ahomaa et al. (2012) observed *Y. enterocolitica* bioserotype 4/O:3 in numbers varying  
432 between 2.3 and 5.4 log CFU/g. Due to the different factors affecting growth and the large impact it  
433 has on prevalence and concentrations found in packages after consumer storage, more studies are  
434 needed regarding the growth of the pathogen in minced meat at different temperatures, including  
435 the variation between strains and varying meat characteristics.

#### 436 4.3. Interventions to control pathogenic *Y. enterocolitica*

437 The prevalence of pathogenic *Y. enterocolitica* on carcasses was set at 16.4% for the baseline model,  
438 though the proportion of carcasses that are pathogenic *Y. enterocolitica* positive at the sternal region  
439 have been shown to vary between slaughterhouses from 7.5 to 37.5% (Van Damme et al., 2015).  
440 Comparing minced meat that is produced from carcasses originating from “good” slaughterhouses  
441 (that produce carcasses with a low prevalence and low concentration) compared to “bad”  
442 slaughterhouses (that produce carcasses with a high prevalence and a high concentration), results in  
443 a more than 30-fold increase in the proportion of highly contaminated *Y. enterocolitica* minced meat  
444 packages. This finding demonstrates the utility of risk differentiation of slaughterhouses (EFSA, 2011)  
445 to control pathogenic *Y. enterocolitica* transmission via minced meat. As the combined effect of  
446 reducing the prevalence and concentration of pathogenic *Y. enterocolitica* on carcasses resulted in  
447 the greatest reduction of highly contaminated minced meat packages, measures to decrease both  
448 the number of positive carcasses and the concentration of pathogenic *Y. enterocolitica* on carcasses  
449 would result in the largest benefit. Many different physical and chemical decontamination  
450 treatments have been described to reduce bacterial contamination on pig carcasses (Loretz et al.,  
451 2011). Besides the effect of (blast) chilling as the most conventional way to reduce bacterial  
452 contamination on carcasses, the effect of steam decontamination and lactic acid decontamination

453 were simulated to represent commonly used physical and chemical decontamination procedures of  
454 pig carcasses. Although blast chilling before conventional chilling has been shown to result in a larger  
455 reduction than conventional air chilling alone for different pathogens (Loretz et al., 2011), blast  
456 chilling has been shown not to reduce pathogenic *Y. enterocolitica* recovery from carcasses  
457 (Nesbakken et al., 2008). The effect of blast chilling on the outcome estimate also seemed rather  
458 limited in the present model. The use of decontamination procedures on carcasses before chilling  
459 was estimated to result in higher reductions of the proportion of highly contaminated minced meat  
460 packages, and would thus likely reduce the public health risk.

461 The baseline model assumed pork bellies as the only source of pathogenic *Y. enterocolitica*  
462 contamination during the production of minced meat. Meat cuts originating from other parts of the  
463 carcass may be contaminated in higher levels and numbers, which would increase the numbers of  
464 pathogenic *Y. enterocolitica* in a minced meat batch and the resulting minced meat packages. Pork  
465 cheeks and tongues have been shown to be highly contaminated with pathogenic *Y. enterocolitica*  
466 (Laukkanen-Ninios et al., 2014b; Messelhauser et al., 2011). As such, the addition of different levels  
467 of head meat for the production of minced meat was simulated using qualitative and quantitative  
468 data from the mandibular region on pig carcasses before cooling as input data to represent meat  
469 from pork cheeks and the throat region. The use of head meat for the production of minced meat  
470 increased the proportion of pathogenic *Y. enterocolitica* positive minced meat packages with  
471 increasing amounts of head meat and had a larger effect on highly contaminated minced meat  
472 packages. The addition of just 10% head meat in minced meat resulted in almost 20 as many highly  
473 contaminated minced meat packages at time of consumption. The addition of pork cheeks and other  
474 potentially highly contaminated meat cuts (such as throat meat) should thus be avoided for the  
475 production of minced meat that is potentially consumed raw.

476 Tonsils have been shown to be highly contaminated with human pathogenic *Yersinia* spp. (Bonardi et  
477 al., 2016; Van Damme et al., 2010) and represent an important risk for carcass contamination. Tonsils  
478 should be removed hygienically after *post mortem* inspection according to EU regulation (EC) No.  
479 853/2004, though parts may remain in the head and result in contamination further down the pork  
480 production line (Fredriksson-Ahomaa et al., 2004). The addition of minimal amounts of tonsillar  
481 tissue in minced meat resulted in a large effect in the proportion of highly contaminated minced  
482 meat packages before consumption, so special care should be taken to remove all remaining tonsillar  
483 tissue from the carcass.

484 Minced meat produced on Monday resulted in a higher proportion of highly contaminated packages  
485 than minced meat produced on Tuesday to Friday. Industrially produced minced meat is usually

486 made from carcasses that are slaughtered the previous day, though carcasses from pigs that are  
487 slaughtered on Friday are stored during the weekend for processing on Monday, resulting in a longer  
488 cold storage. After storage of pork bellies during 4 and 8 days at 4°C, van Netten et al. (1997)  
489 observed more than 1 and 4 log<sub>10</sub> increase of cold and acid adapted *Y. enterocolitica* serotype O:3.  
490 Therefore, minced meat that is produced from carcasses that have been stored for several days may  
491 represent a larger risk for public health than freshly slaughtered pig carcasses. This implies that the  
492 shelf life for minced meat may be adapted depending on the cold storage time of carcasses to reduce  
493 the proportion of minced meat packages that are (highly) contaminated with pathogenic *Y.*  
494 *enterocolitica*.

495 Consumer practices were shown to have a large effect on the proportion of minced meat packages  
496 with high numbers of pathogenic *Y. enterocolitica* at time of consumption. When all consumers  
497 would store minced meat at 4°C, a 1000-fold reduction in the number of highly contaminated  
498 packages could be expected. A similar reduction was seen if consumers would consume the minced  
499 meat at the day of purchase. Storage of minced meat in ambient atmosphere leads to higher  
500 maximum growth rates for *Y. enterocolitica* compared to packaging with 30% CO<sub>2</sub>, resulting in higher  
501 estimates of highly contaminated packages at the end of storage. Nevertheless, the storage time at  
502 ambient atmosphere is presumably shorter compared to minced meat stored under MAP conditions  
503 due to the shorter shelf life (Strotmann et al., 2008). Limbo et al. (2010) calculated that the mean  
504 shelf life of MAP minced beef was 9 days at the recommended storage temperature of about 4°C.  
505 The proportion of highly contaminated packages in the present study was higher when all MAP  
506 would be stored until the use-before date compared to the storage of packages at ambient  
507 atmosphere for two days or less. Although MAP is introduced to reduce bacterial growth and prolong  
508 shelf-life of products, the longer shelf-life could potentially increase the risk of yersiniosis due to the  
509 growth of pathogenic *Y. enterocolitica* during prolonged storage at refrigerated conditions.

## 510 **5. Conclusions**

511 Meat producers should focus on reducing the number of pathogenic *Y. enterocolitica* contaminated  
512 minced meat packages, which can be achieved by using meat cuts that are less contaminated with  
513 pathogenic *Y. enterocolitica*. As such, belly cuts should be preferred over head meat. Moreover, meat  
514 produced from carcasses of slaughterhouses with lower contamination results in less pathogenic *Y.*  
515 *enterocolitica* contaminated minced meat packages. Finally, it's important that the tonsils are  
516 completely removed in the slaughterhouse as the (accidental) addition of minimal amounts of  
517 tonsillar tissue has a large effect on the proportion of highly contaminated minced meat packages.  
518 Nevertheless, the number of packages that contain high numbers of pathogenic *Y. enterocolitica*,  
519 which are expected to cause the highest risk of yersiniosis, is primarily influenced by consumer

520 storage practices. A reduced storage time (under one day) or a storage temperature (below 4°C)  
521 would largely reduce the proportion of packages containing high numbers of pathogenic *Y.*  
522 *enterocolitica*.

523

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644 **Figure captions**645 **Figure 1. Food pathway of the baseline model to describe *Y. enterocolitica* in minced meat**  
646 **produced by an industrial meat processing plant.**

647 The model starts with the contamination of carcasses in the slaughterhouse after evisceration and  
648 ends with a 0.5-package of minced pork just before consumption and/or preparation.

649 **Figure 2. Distributions of concentrations of *Y. enterocolitica* in 0.5-kg minced meat packages after**  
650 **storage at consumer level (based on 100 000 iterations) using (1) the baseline scenario that only**  
651 **assumed pork bellies as a source of contamination (dashed line) and (2) the alternative scenario in**  
652 **which 1 g of tonsillar tissue is added to a 900-kg minced batch (solid line).**

653 Concentrations of *Y. enterocolitica* are given for contaminated packages only; the areas under the  
654 curves reflect the prevalence of 15.4% in the baseline scenario and 37.9% in the alternative scenario.

655 **Figure 3. Effect of initial pig carcass contamination in slaughterhouses on *Y. enterocolitica***  
656 **contamination of minced meat packages just before consumption.**

657 The proportion of *Y. enterocolitica* positive minced meat packages of the alternative scenarios are  
658 expressed relative to the proportion of minced meat packages in the baseline model. Relative  
659 proportions are log transformed, so the baseline gets a value zero, and -1 and 1 represent a tenfold  
660 reduction and increase of the proportion, respectively. The baseline model used a prevalence ( $P_{\text{initial}}$ )  
661 of 16.4% and a mean concentration ( $C_{\text{initial}}$ ) of  $-2.565 \log_{10} Y. enterocolitica/\text{cm}^2$ . Alternative scenarios  
662 were simulated using a lower/higher prevalence ( $P_{\text{initial}}$  of 7.5% or 37.5%, respectively) and/or a  
663 lower/higher concentration (mean  $C_{\text{initial}}$  of 0.5  $\log_{10}$  lower or higher compared to the baseline value,  
664 respectively). The grey bars represent the outcome of the proportion of *Y. enterocolitica* positive 0.5-  
665 kg minced meat packages. The black bars represent the results for 0.5-kg minced meat packages that  
666 contain more than  $3 \log_{10} Y. enterocolitica$  at time of consumption or preparation.

667 **Figure 4. Effect of cooling and carcass decontamination steps on *Y. enterocolitica* contaminated**  
668 **minced meat packages.**

669 The proportion of *Y. enterocolitica* positive minced meat packages of the alternative scenarios are  
670 expressed relative to the proportion of minced meat packages in the baseline model. The baseline  
671 model assumed a 0.6 log reduction of *Y. enterocolitica* during blast chilling. The storage time of  
672 carcasses in the baseline model was 20h (for carcasses of pigs slaughtered on Monday-Thursday) or  
673 68h (for carcasses of pigs slaughtered on Friday).

674 The grey bars represent the outcome of the proportion of *Y. enterocolitica* positive 0.5-kg minced  
675 meat packages. The black bars represent the proportion of 0.5-kg minced meat packages that  
676 contain more than  $3 \log_{10} Y. enterocolitica$  at time of consumption or preparation.

677 **Figure 5. Evaluation of the addition of head meat and tonsillar tissue to a 900-kg batch of minced**  
678 **meat on *Y. enterocolitica* contaminated minced meat packages just before consumption.**

679 The proportion of *Y. enterocolitica* positive minced meat packages of the alternative scenarios are  
680 expressed relative to the proportion of minced meat packages in the baseline model. The baseline  
681 model only assumed pork bellies as a source of *Y. enterocolitica* contamination. The grey bars  
682 represent the outcome of the proportion of *Y. enterocolitica* positive 0.5-kg minced meat packages.  
683 The black bars represent the proportion of 0.5-kg minced meat packages that contain more than 3  
684  $\log_{10}$  *Y. enterocolitica* at time of consumption or preparation.

685 **Figure 6. Evaluation of consumer practices on *Y. enterocolitica* contaminated minced meat**  
686 **packages just before consumption.**

687 The proportion of highly contaminated ( $> 3 \log_{10}$ ) *Y. enterocolitica* 0.5-kg minced meat packages of  
688 the alternative scenarios are expressed relative to the proportion of highly contaminated 0.5-kg  
689 minced meat packages in the baseline model (= stored in modified atmosphere packages (MAP), 30%  
690 CO<sub>2</sub>). The black bars represent minced meat packages stored in MAP. The bars with diagonal stripes  
691 represent storage at ambient atmosphere. \* Storage until use-by date was only simulated for MAP  
692 minced meat.

693 **Figure 7. Results of the uncertainty analyses of the baseline model.**

694 The grey bars represent the outcome of the proportion of *Y. enterocolitica* positive 0.5-kg minced  
695 meat packages. The black bars represent the proportion of 0.5-kg minced meat packages that  
696 contain more than 3  $\log_{10}$  *Y. enterocolitica* at time of consumption or preparation. The relative  
697 proportion for U23 (reduced growth in minced meat) was truncated at -1.5.



**Table 1. Overview of the different steps, processes and units that were used in the risk assessment model for *Y. enterocolitica* in minced pork.**

Processing step	Basic process	Unit
1 Contamination of carcasses (after evisceration, before chilling)	Initial contamination	Carcass half – belly area (2000 cm <sup>2</sup> )
2 Chilling room	Inactivation Growth	Carcass half – belly area (2000 cm <sup>2</sup> )
3 Cutting and derinding	Removal	Belly cut (2000 cm <sup>2</sup> ; 7.5 kg)
4 Grinding and seasoning	Mixing	Batch of minced meat (900 kg)
5 Packaging	Partitioning	Minced meat package (0.5 kg)
6 Storage (meat processing plant and retail)	Growth	Minced meat package (0.5 kg)
7 Storage (consumer)	Growth	Minced meat package (0.5 kg)

**Table 2. Overview of variables and parameters in the baseline Modular Process Risk Model (MPRM) for human pathogenic *Y. enterocolitica* in minced meat.**

Module	Variable	Description	Unit	Value/distribution/equation	Source
Input (carcasses, sternal region, after evisceration)	$P_{\text{initial}}$	Prevalence of <i>Y. enterocolitica</i> on pig carcasses (sternal region) after evisceration	%	16.39	Van Damme et al. (2015)
	$C_{\text{initial}}$	Concentration of <i>Y. enterocolitica</i> on pig carcasses (sternal region) after evisceration (positive carcasses only)	$\text{Log}_{10}$ CFU/cm <sup>2</sup>	~ Normal(-2.565; 0.736) truncated at a minimum value of -1.85	Calculated based on data from Van Damme et al. (2015)
Inactivation during carcass chilling	$I_{\text{cc}}$	Inactivation	$\text{Log}_{10}$ reduction	-0.6	King et al. (2012)
	$C_{\text{cci}}$	Concentration on pig carcasses after inactivation during chilling	$\text{Log}_{10}$ CFU/cm <sup>2</sup>	$= C_{\text{initial}} + I_{\text{cc}}$	Calculation
Growth during carcass cold storage	$\text{Time}_{\text{ccg}}$	Cold storage time of carcasses and all head meat and tonsils applied in the same batch	h	~ Discrete(20, 68), (4, 1)	Company info
	$\lambda_{\text{ccg}}$	Lag phase during carcass cold storage	h	24	Van Netten et al. (1997)
	$D_{\text{ccg}}$	Doubling time during cold storage	h	9.978	ComBase
	$N_{\text{ccg}}$	Number of <i>Y. enterocolitica</i> after growth during cold storage	CFU/cm <sup>2</sup>	$= 10^{C_{\text{cci}}} \times 2^{(\text{Time}_{\text{ccg}} - \lambda_{\text{ccg}})/D_{\text{ccg}}}$	Calculation
Cutting and derinding	$S_{\text{bc}}$	Surface of belly cut	cm <sup>2</sup>	2000	Assumption
	$N_{\text{bc}}$	Number of <i>Y. enterocolitica</i> per belly after cutting	CFU/belly	$= N_{\text{ccg}} \times S_{\text{bc}}$ (rounded to an integer value)	Calculation
	$R_{\text{bd}}$	Proportion of <i>Y. enterocolitica</i> that remain on the belly cut after derinding	%	50%	Assumption
	$N_{\text{bdr}}$	Number of <i>Y. enterocolitica</i> on belly cut after derinding	CFU/belly	~ Binomial( $N_{\text{bc}}$ , $R_{\text{bd}}$ )	Calculation
Mixing and grounding	$W_{\text{b}}$	Weight of a batch of minced meat	kg	900	Company information
	%bellies	Proportion of bellies per batch (w:w)	%	34	Company information
	$W_{\text{bc}}$	Weight of a belly cut	kg	7.5	Company information
	$n_{\text{bb}}$	Number of bellies per batch		$= \frac{W_{\text{b}} \times \% \text{bellies}}{W_{\text{bc}}}$	Calculation
	$n_{\text{pbb}}$	Number of positive bellies per batch		~ Binomial( $n_{\text{bb}}$ , $P_{\text{initial}}$ )	Calculation

	$N_{mb}$	Number of <i>Y. enterocolitica</i> in one minced meat batch	CFU	$= \sum_{i=1}^{n_{pbb}} N_{bdr,i}$	Calculation
Partitioning/packaging	$W_{mp}$	Weight per minced meat package	kg	0.5	Company information
	$N_{mp}$	Number of <i>Y. enterocolitica</i> in one minced meat package after packaging/partitioning	CFU	$\sim \text{Binomial}(N_{mb}, W_{mp}/W_b)$	Assumption
Storage at retail	$Temp_{rg}$	Temperature during storage in meat processing plant and at retail	°C	4	Assumption
	$Time_{rg}$	Time between packaging and selling at retail	h	48	Assumption
	$\mu_{max,rg}$	Maximum growth rate (MAP)	Log <sub>10</sub> CFU/h	$= 0.0003 \times Temp_{rg}^2 + 0.0005 \times Temp_{rg} + 0.0103$	ComBase
	$N_{rg}$	Number of <i>Y. enterocolitica</i> in one package of minced meat after storage at retail	CFU	$= N_{mp} \times 10^{\mu_{max,rg} \times Time_{rg}}$	Calculation
Storage at consumer level	$Temp_{cg}$	Temperature of home refrigerators	°C	$\sim \text{Pert}(25\% 5; 50\% 7; 75\% 9)$	Devriese et al. (2006)
	$Time_{cg}$	Time between purchase and consumption/preparation	days	$\sim \text{Pert}(0;1;4)$	Marklinder et al. (2004)
	$\mu_{max,cg}$	Maximum growth rate (MAP)	Log <sub>10</sub> CFU/h	$0.0003 \times Temp_{cg}^2 + 0.0005 \times Temp_{cg} + 0.0103$	ComBase
	$N_{cg}$	Number of <i>Y. enterocolitica</i> in one package of minced meat at the end of storage (just before consumption or preparation)	CFU/0.5-kg package	$= N_{rg} \times 10^{\mu_{max,cg} \times Time_{cg} \times 24}$	Calculation

**Table 3. Overview of variables and parameters to evaluate alternative scenarios at slaughterhouse level.**

Code	Description of the scenario	Variable	Alternative value/distribution/model	Source
A1	Lower initial prevalence on carcasses	$P_{\text{initial}}$	7.5	Van Damme et al. (2015)
A2	Lower initial concentration on carcasses	$C_{\text{initial}}$	$\sim \text{Normal}(-3.065; 0.736)$ truncated at a minimum value of -1.85	Based on data from Van Damme et al. (2015)
A3	Lower initial prevalence and concentration on carcasses	$P_{\text{initial}}$	7.5	Van Damme et al. (2015)
		$C_{\text{initial}}$	$\sim \text{Normal}(-3.065; 0.736)$ truncated at a minimum value of -1.85	Based on data from Van Damme et al. (2015)
A4	Higher initial prevalence on carcasses	$P_{\text{initial}}$	37.5	Van Damme et al. (2015)
A5	Higher initial concentration on carcasses	$C_{\text{initial}}$	$\sim \text{Normal}(-2.065; 0.736)$ truncated at a minimum value of -1.85	Based on data from Van Damme et al. (2015)
A6	Higher initial prevalence and concentration on carcasses	$P_{\text{initial}}$	37.5	Van Damme et al. (2015)
		$C_{\text{initial}}$	$\sim \text{Normal}(-2.065; 0.736)$ truncated at a minimum value of -1.85	Based on data from Van Damme et al. (2015)
A7	Only conventional air chilling (no blast chilling)	$I_{\text{cc}}$	-0.1	King et al. (2012)
A8	Steam condensation followed by conventional chilling	$I_{\text{cc}}$	$\sim \text{-Pert}(0.7, 2.2, 4) - 0.1$	Smulders et al. (2012) and King et al. (2012)
A9	Steam condensation followed by blast chilling and conventional chilling	$I_{\text{cc}}$	$\sim \text{-Pert}(0.7, 2.2, 4) - 0.6$	Smulders et al. (2012) and King et al. (2012)
A10	Lactic acid treatment followed by conventional chilling and cold storage	$I_{\text{cc}}$	-0.7	King et al. (2012)
		$\lambda_{\text{ccg}}$	48	van Netten et al. (1997)
		$D_{\text{ccg}}$	12.4	van Netten et al. (1997)
A11	Lactic acid treatment followed by blast chilling and conventional chilling and cold storage	$I_{\text{cc}}$	-1.6	King et al. (2012)
		$\lambda_{\text{ccg}}$	48	van Netten et al. (1997)
		$D_{\text{ccg}}$	12.4	van Netten et al. (1997)
A12	Minced meat produced using carcasses stored over weekend	$\text{Time}_{\text{ccg}}$	68h	Company information
A13	Minced meat produced using carcasses the day after slaughter	$\text{Time}_{\text{ccg}}$	20h	Company information

**Table 4. Overview of variables and parameters to evaluate alternative scenarios during grinding.**

Scenario	Variable	Description	Alternative value/distribution/model	Source
B1-B3: Addition of head meat	$P_{initial,m}^*$	Prevalence of <i>Y. enterocolitica</i> on pig carcasses (mandibular region) after evisceration	28.89%	Van Damme et al. (2015)
	$C_{initial,m}^*$	Concentration of <i>Y. enterocolitica</i> on pig carcasses (mandibular region) after evisceration (positive carcasses only)	$\sim$ Normal (-0.578; 1.256) truncated at a minimum of 0.15 (in $\log_{10}$ CFU/100 $\text{cm}^2$ )	Based on data from Van Damme et al. (2015)
	$C_{mci}^*$	Concentration on pig carcasses (mandibula) after inactivation during chilling	$= C_{initial,m} + I_{cc}$ (in $\log_{10}$ CFU/100 $\text{cm}^2$ )	Calculation
	$N_{mcg}^*$	Number of <i>Y. enterocolitica</i> after growth during cold storage	$= 10^{C_{mci}} \times 2^{(Time_{ccg} - \lambda_{ccg})/D_{ccg}}$ (in CFU/100 $\text{cm}^2$ )	Calculation
	%headmeat*	% of head meat in a batch of minced meat (w:w)	1% (B1), 10% (B2) or 50% (B3)	Assumption
	$W_{hm}^*$	Weight of a piece of head meat	0.075 kg	Company information
	$n_{hb}^*$	Number of head meat cuts per batch	$n_{hb} = \frac{W_b \times \%headmeat}{W_{hm}}$	Calculation
	$n_{phb}^*$	Number of positive head meat cuts per batch	$\sim$ Binomial ( $n_{hb}, P_{initial,m}$ )	Assumption
	$N_{mb}$	Number of <i>Y. enterocolitica</i> in one minced meat batch	$N_{mb} = \sum_{i=1}^{n_{phb}} N_{bdr,i} + \sum_{i=1}^{n_{phb}} N_{mb,i}$ (in CFU)	Calculation
B4-B6: Addition of tonsillar tissue	$P_{initial,t}^*$	Prevalence of <i>Y. enterocolitica</i> in pig tonsils at time of evisceration	44.33%	Van Damme et al. (2015)
	$C_{initial,t}^*$	Concentration of <i>Y. enterocolitica</i> in pig tonsils at time of evisceration	Pert(1.00;4.00;5.91) in $\log_{10}$ CFU/g	Based on data from Van Damme et al. (2015)
	$C_{tci}^*$	Concentration during chilling (after inactivation)	$= C_{initial,t} + I_{cc}$ (in $\log_{10}$ CFU/g)	Calculation
	$N_{tcg}^*$	Number of <i>Y. enterocolitica</i> after growth during cold storage	$N_{tci} \times 2^{(Time_{ccg} - \lambda_{ccg})/D_{ccg}}$ (in CFU/g)	Calculation
	$n_{tb}^*$	Number of tonsil pieces per batch	1 (B4 and B5) or 10 (B6)	Scenarios
	$W_t^*$	Weight of a tonsil piece	1g (B4 and B6) or 10 g (B5)	Scenarios
	$n_{ptb}^*$	Number of positive tonsil pieces per batch	$\sim$ Binomial ( $n_{tb}, P_{initial,t}$ )	Calculation
	$N_{mb}$	Number of <i>Y. enterocolitica</i> in one minced meat batch	$N_{mb} = \sum_{i=1}^{n_{ptb}} N_{bdr,i} + W_t \sum_{i=1}^{n_{ptb}} N_{tcg,i}$ (in CFU)	Calculation
B7: Smaller	$W_b$	Weight of a batch of minced meat	140 kg	Company information

batch of minced meat				
B8: Larger batch of minced meat	$W_b$	Weight of a batch of minced meat	1500 kg	Assumption

\* new variable

**Table 5. Overview of variables and parameters to evaluate alternative scenarios at consumer level.**

Code	Description	Parameter	Value	Source
C1-4	Consumer storage temperature of 4°C, 7°C, 10°C or 15°C	Temp <sub>cg</sub>	4°C (C1), 7°C (C2), 10°C (C3) or 15°C (C4)	Scenarios
C5-9	Consumer storage for 0, 1, 2, 3 or 4 days	Time <sub>cg</sub>	0 days (C5), 1 day (C6), 2 days (C7), 3 days (C8) or 4 days (C9)	Scenarios
C1-9 at ambient atmosphere	Storage at ambient atmosphere	μ <sub>max,rg</sub>	$0.0004 \times \text{Temp}_{rg}^2 + 0.0012 \times \text{Temp}_{rg} + 0.0174$ (in log <sub>10</sub> CFU/h)	ComBase
		μ <sub>max,cg</sub>	$0.0004 \times \text{Temp}_{cg}^2 + 0.0012 \times \text{Temp}_{cg} + 0.0174$ (in log <sub>10</sub> CFU/h)	ComBase
C10	Consumer storage until the use-by date	Time <sub>cg</sub>	7 days	Company info
C11	Purchase and consumption at use-by-date	Time <sub>rg</sub>	9 days	Company info
		Time <sub>cg</sub>	0 days	Assumption

**Table 6. Overview of the variables and parameters to evaluate uncertainty.**

Code	Variable	Alternative value/distribution	Source
U1	C <sub>initial</sub>	~ Normal(-2.065; 0.736) (in log <sub>10</sub> CFU/cm <sup>2</sup> )	Assumption
U2	C <sub>initial</sub>	~ Normal(-3.065; 0.736) (in log <sub>10</sub> CFU/cm <sup>2</sup> )	Assumption
U3	C <sub>initial</sub>	~ Normal(-2.565; 1.236) (in log <sub>10</sub> CFU/cm <sup>2</sup> )	Assumption
U4	C <sub>initial</sub>	~ Normal(-2.565; 0.236) (in log <sub>10</sub> CFU/cm <sup>2</sup> )	Assumption
U5	P <sub>initial</sub>	23.1%	Van Damme et al. (2015)
U6	P <sub>initial</sub>	13.3%	Van Damme et al. (2015)
U7	I <sub>cc</sub>	-0.03 log <sub>10</sub> reduction	El-Zawahry and Grecz (1981)
U8	I <sub>cc</sub>	-0.8 log <sub>10</sub> reduction	King et al. (2012)
U9	L <sub>ccg</sub>	77 h	Greer and Dilts (1995)
U10	L <sub>ccg</sub>	0 h	Greer and Dilts (1995)
	D <sub>ccg</sub>	10.36 h	ComBase
U11	R <sub>bd</sub>	25%	Assumption
U12	R <sub>bd</sub>	75%	Assumption
U13	W <sub>b</sub>	850 kg	Assumption
U14	W <sub>b</sub>	950 kg	Assumption
U15	%bellies	29%	Assumption
U16	%bellies	39%	Assumption
U17	W <sub>bdr</sub>	7 kg	Assumption
U18	W <sub>bdr</sub>	8 kg	Assumption
U19	Temp <sub>rg</sub>	2°C	Assumption
U20	Temp <sub>rg</sub>	6°C	Assumption
U21	Time <sub>rg</sub>	1d	Assumption
U22	Time <sub>rg</sub>	3d	Assumption
U23	μ <sub>max,rg</sub>	$\frac{0.0003 \times \text{Temp}_{rg}^2 + 0.0005 \times \text{Temp}_{rg} + 0.0103}{2}$	Assumption
	μ <sub>max,cg</sub>	$\frac{0.0003 \times \text{Temp}_{cg}^2 + 0.0005 \times \text{Temp}_{cg} + 0.0103}{2}$	Assumption



Figure 1

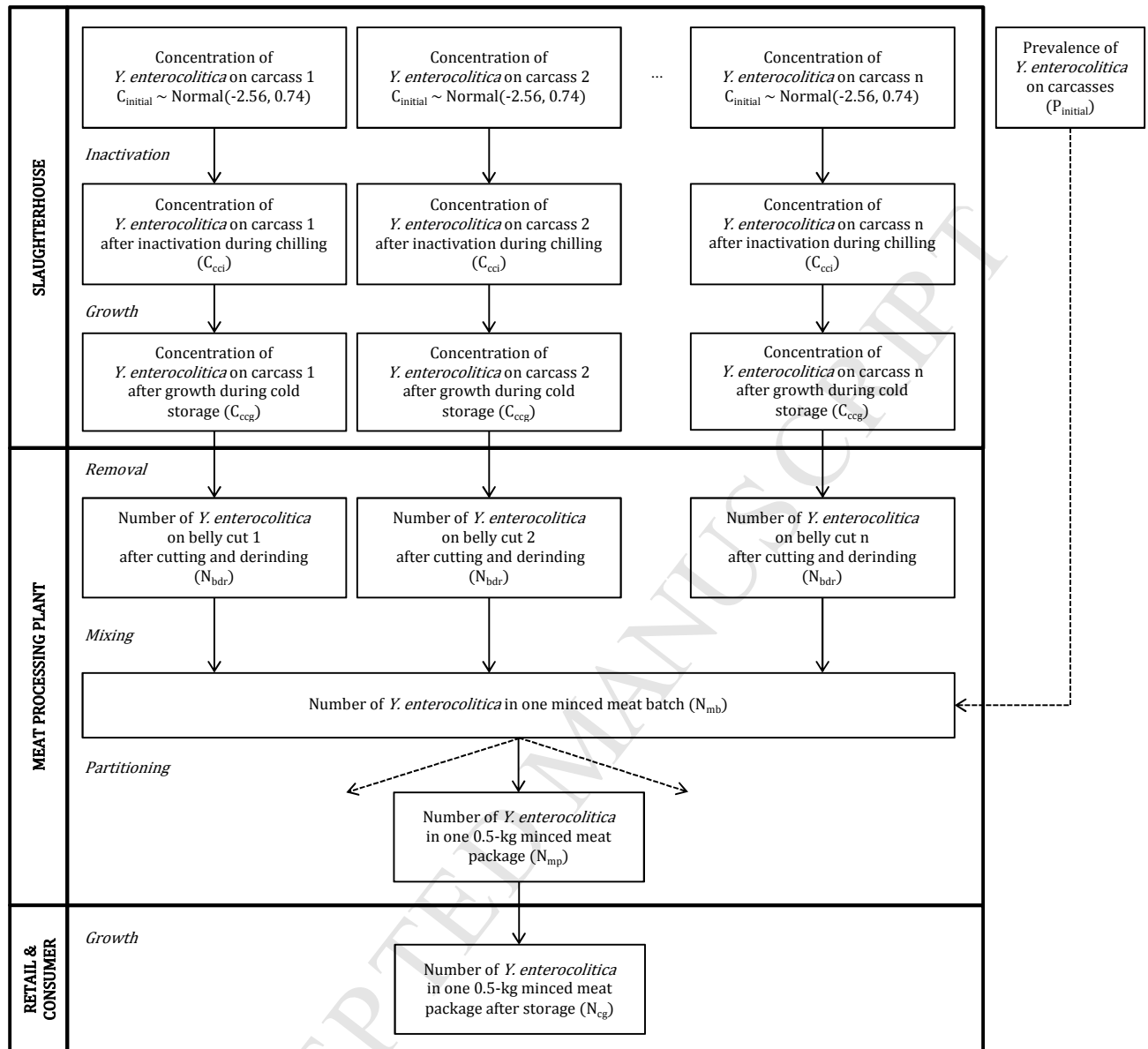


Figure 2

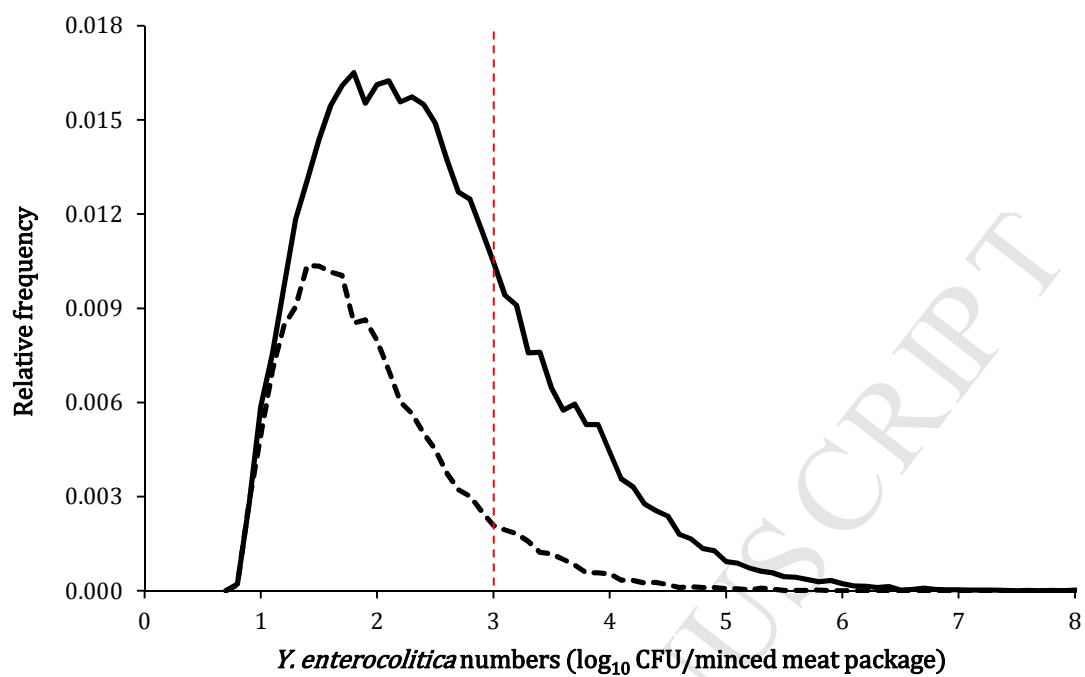
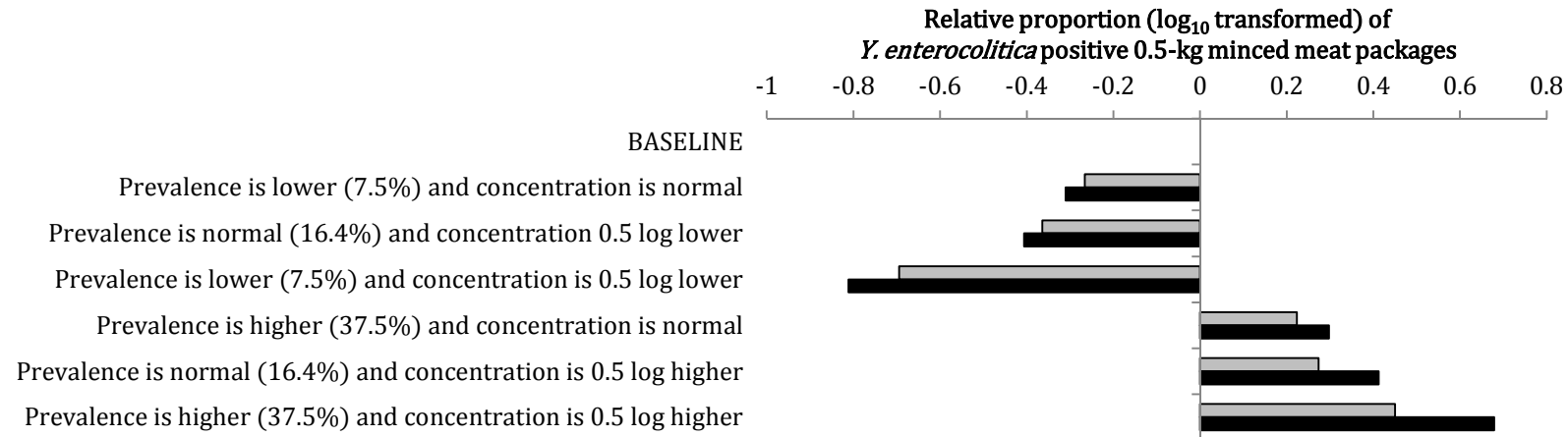


Figure 3



ACCEPTED MANUSCRIPT

Figure 4

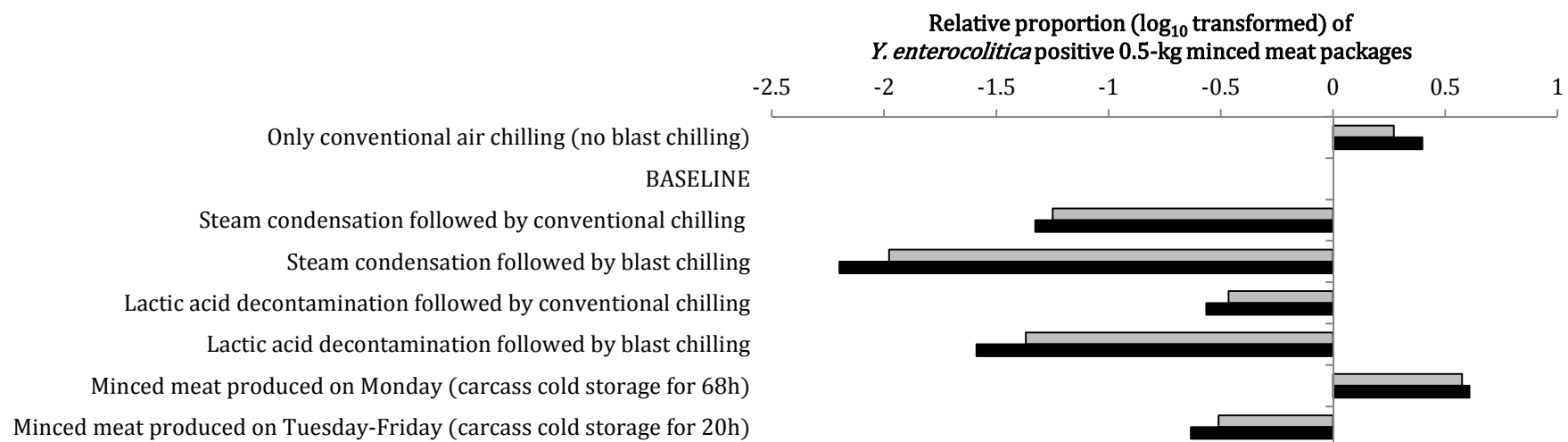


Figure 5

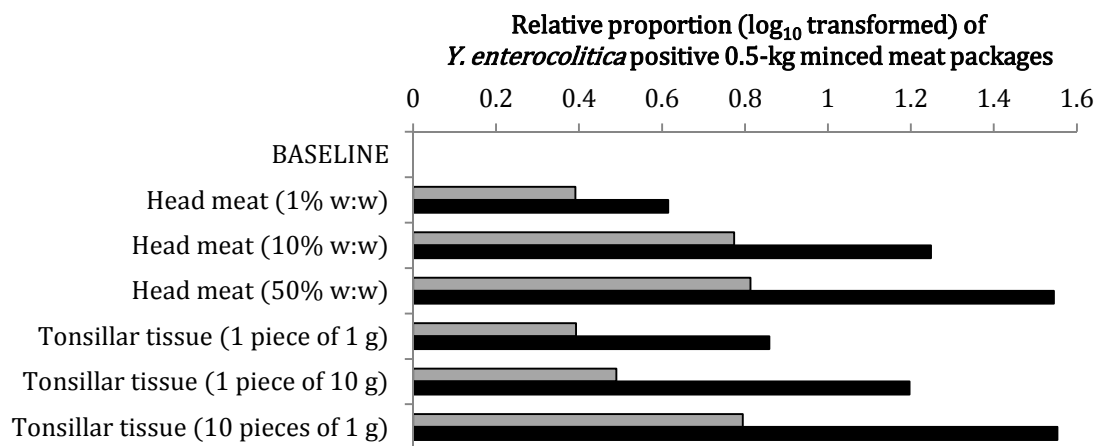


Figure 6

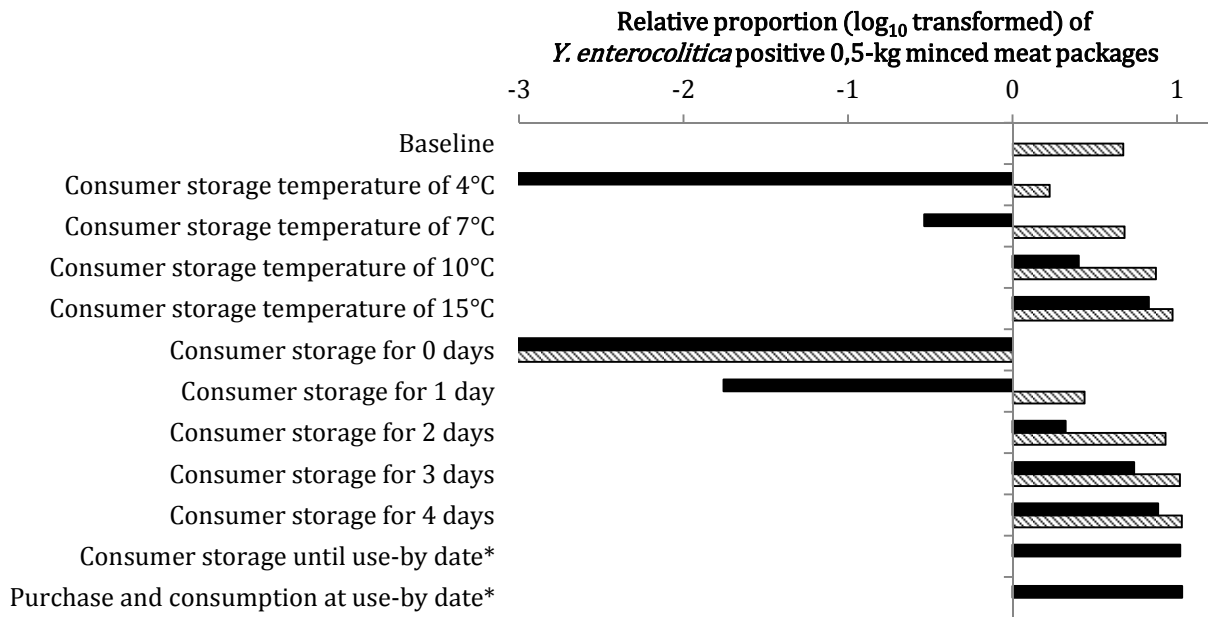
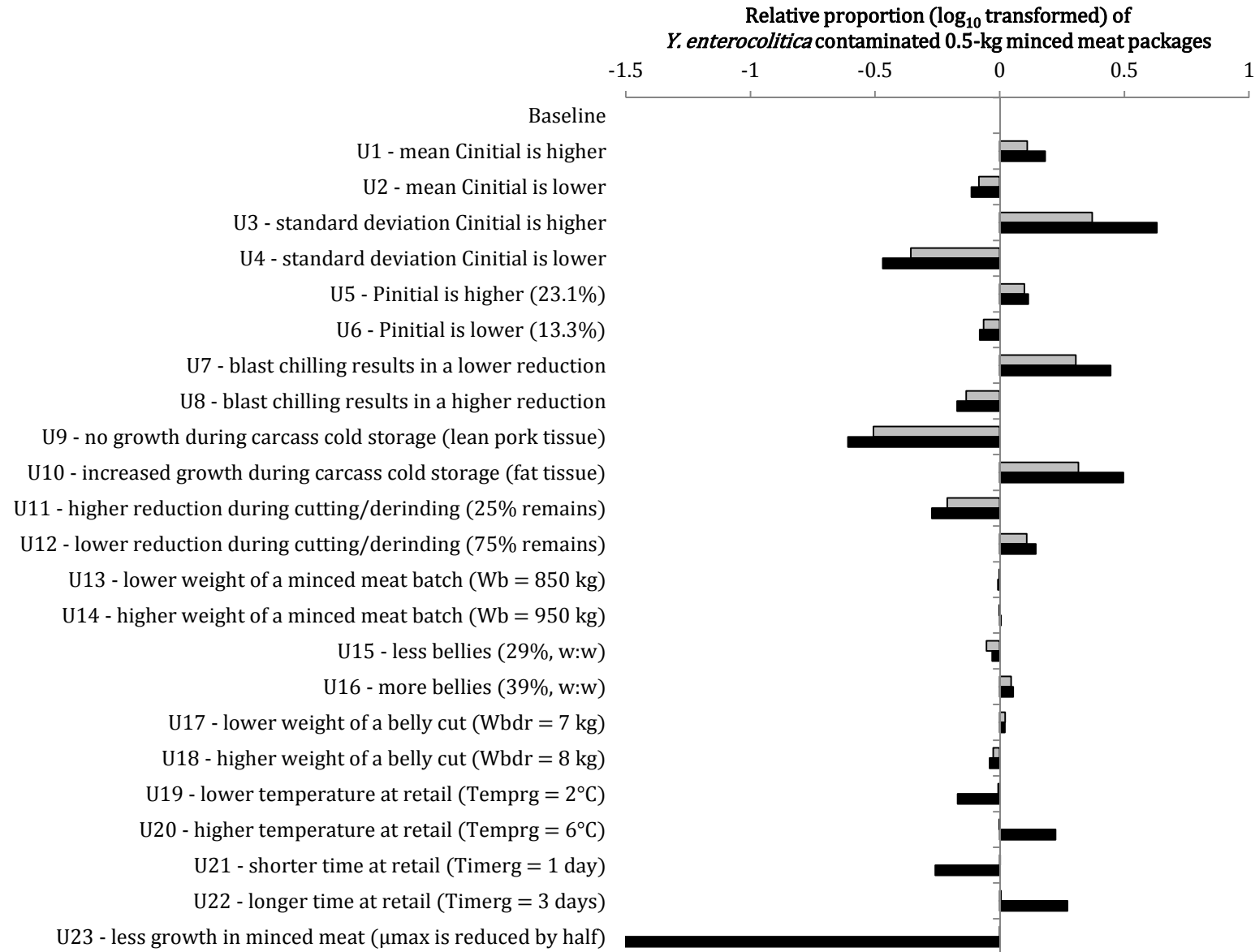


Figure 7



**Highlights**

- Contamination of minced meat with human pathogenic *Y. enterocolitica* was modelled.
- The endpoint of the assessment was the proportion of (highly) contaminated packages.
- Control of *Y. enterocolitica* contamination at slaughterhouse level is important.
- Pork bellies are preferred over head meat for the production of minced meat.
- Consumer practices strongly influence the number of highly contaminated packages.