

## ***Analysis of Vibrio vulnificus Infection Risk When Consuming Depurated Raw Oysters***

The Faculty of Oregon State University has made this article openly available.  
Please share how this access benefits you. Your story matters.

<b>Citation</b>	Deng, K., Wu, X., Fuentes, C., Su, Y. C., Welti-Chanes, J., Paredes-Sabja, D., & Torres, J. A. (2015). Analysis of Vibrio vulnificus Infection Risk When Consuming Depurated Raw Oysters. <i>Journal of Food Protection</i> , 78(6), 1113-1118. doi:10.4315/0362-028X.JFP-14-421
<b>DOI</b>	10.4315/0362-028X.JFP-14-421
<b>Publisher</b>	International Association for Food Protection
<b>Version</b>	Accepted Manuscript
<b>Terms of Use</b>	<a href="http://cdss.library.oregonstate.edu/sa-termsfuse">http://cdss.library.oregonstate.edu/sa-termsfuse</a>

1 Running title: Risk analysis of oyster depuration

2

3 **Analysis of *Vibrio vulnificus* infection risk when consuming depurated raw oysters**

4

5 **Kai Deng<sup>1,2</sup>, Xulei Wu<sup>3</sup>, Claudio Fuentes<sup>4</sup>, Yi-Cheng Su<sup>3</sup>, Jorge Welti-Chanes<sup>5</sup>,**

6 **Daniel Paredes-Sabja<sup>2,6\*</sup>, and J. Antonio Torres<sup>1\*\*</sup>**

7

8 (1) Food Process Engineering Group, Department of Food Science & Technology, Oregon State  
9 University, Corvallis, OR 97331, USA; (2) Laboratorio de Mecanismos de Patogénesis Bacteriana,  
10 Departamento de Ciencias Biológicas, Facultad de Ciencias Biológicas, Universidad Andrés Bello,  
11 Santiago, Chile; (3) Seafood Research and Education Center, Oregon State University, Astoria, OR  
12 97103, USA; and (4) Department of Statistics, Oregon State University, Corvallis, OR 97331, USA;  
13 (5) Escuela de Ingenierías y Ciencias, Tecnológico de Monterrey, Av. Eugenio Garza Sada 2501 Sur,  
14 Col. Tecnológico, 64849, Monterrey, NL; (6) Department of Biomedical Sciences, College of  
15 Veterinary Medicine, Oregon State University, Corvallis, OR, 97331

16

17

18

19

20

21

22

23

24 **Keywords:** Oyster, *Vibrio vulnificus*, dose-response, depuration, Monte Carlo, risk analysis

25

26 \* Corresponding author, Tel: +56 (2) 2770-3225; Fax: +56 (2) 2698-0414

27 Email: Daniel.Paredes.Sabja@gmail.com

28 \*\* Corresponding author, Tel: +1 (541) 737-4757; Fax: +1 (541) 737-1877

29 Email: J\_Antonio.Torres@oregonstate.edu

30 **ABSTRACT**

31 A beta Poisson dose-response model for *Vibrio vulnificus* food poisoning cases leading to  
32 septicemia was used when evaluating the effect of 15°C depuration on the estimated risk of raw  
33 oyster consumption. Statistical variability sources included *V. vulnificus* load at harvest, time and  
34 temperature during harvest and transportation to processing plants, decimal reductions (SV)  
35 observed during experimental circulation depuration treatments, refrigerated storage time before  
36 consumption, oyster size, and number of oysters per consumption event. Although reaching non-  
37 detectable *V. vulnificus* levels (<30 MPN/g) throughout the year and a 3.52 SV were estimated not  
38 possible at 95% confidence, depuration for 1, 2, 3, and 4 d would reduce the warm (Jun-Sep)  
39 season risk from 2,669 cases to 558, 93, 38, and 47 cases per 100 million consumption events,  
40 respectively. At 95% confidence, 47 and 16 h depuration would reduce the warm and transition  
41 (Apr-May, Oct-Nov) season risk, respectively, to 100 cases per 100 million consumption events  
42 assumed to be an acceptable risk, while 1 case per 100 million events would be the risk when  
43 consuming untreated raw oysters in the cold (Dec-Mar) season.

44

45 Pathogens frequently present in oysters include *Vibrio* species and noroviruses (6, 17).  
46 Among *Vibrio* species, 11 can cause human disease including *V. parahaemolyticus*, *V. vulnificus* and  
47 *V. cholerae* causing severe illnesses (19). During warm seasons, these halophilic Gram-negative  
48 bacteria can reach high numbers in oyster harvesting areas with moderate salinity (20). The CDC  
49 reports over 400 *Vibrio* illnesses each year including about 90 due to *V. vulnificus* occurring mostly  
50 during warm-weather months (5, 25). Diseases caused by *V. vulnificus* are among the most severe  
51 food-borne infections and have the highest case-fatality rate in the USA (23). A number of dose-  
52 response models have been used to predict the probability of illness when consumers are exposed  
53 to a given pathogen dose (12, 27). Since human dose-response studies cannot be conducted,  
54 modelling of the *V. vulnificus* dose-response relationship is based on estimates of dose exposure  
55 per serving, number of servings in the susceptible population and the number of oyster-associated  
56 cases of *V. vulnificus* septicemia cases reported to the CDC (1). The frequently used Beta-Poisson  
57 model (18) has been used to estimate the number of *V. vulnificus* cases likely to occur when  
58 consuming raw oysters harvested in the Gulf of Mexico (1) and was the model used in this study.

59 Depuration consists of placing live oysters in circulating seawater tanks. During treatment,  
60 the oysters' pumping activity expels *V. vulnificus* and other contaminants from their gills and  
61 intestinal tract (9, 22). To allow its reuse, seawater is filtered and then disinfected by UV, ozone or  
62 chlorine (21). A 15°C treatment temperature has been recommended for the depuration of oysters

63 in circulating seawater (9). A 44 h depuration time reducing *V. vulnificus* to non-detectable counts  
64 determined using a 3-tube most probable number (MPN) procedure and 1:10 dilution, i.e., less  
65 than 30 MPN/g is prescribed in the National Shellfish Sanitation Program (see p. 140, 3).  
66 Increasingly high consumer expectations of quality and safety make it necessary to develop  
67 depuration treatments that are effective for every raw oyster production lot. The design should  
68 consider the variability of production and handling factors including harvest, transportation, post-  
69 harvest processing, storage and other risk factors. This is not possible using deterministic  
70 algorithms or experimental test runs in processing plants. In this study, a Monte Carlo procedure  
71 was applied to estimate the risk of consuming raw oysters treated by depuration. Several recent  
72 reports describe its application to evaluate the uncertainty of food safety, quality and shelf-life  
73 estimations (10, 11, 26, 29, 31, 32). In this study, procedures were developed to estimate the  
74 number of septicemia infection cases per 100 million oyster consumption events, and to  
75 determine depuration times that would reduce this risk to an acceptable level defined as 100  
76 cases (31). This study included risk factors beginning at harvest and ending when raw oysters are  
77 consumed. The procedures used in this study estimated whether process objectives are met with a  
78 confidence set at 95%, while considering the statistical variability of these multiple factors.

79 **MATERIALS AND METHODS**

80 Statistical distributions can describe: (i) *V. vulnificus* load at harvest as a function of season;  
81 (ii) time and temperature during transportation from harvest site to processing plants; (iii) kinetic  
82 *V. vulnificus* growth parameters during harvest and transportation; (iv) depuration parameters for  
83 models developed using published circulation depuration laboratory data; (v) oyster size;  
84 (vi) refrigerated storage time before consumption; (vii) *V. vulnificus* die-off during refrigerated  
85 storage; and, (9) oysters consumed per serving.

86 Motes et al. (24) quantified the *V. vulnificus* ( $\log(\text{No, MPN/g oyster})$ ) in oysters  
87 collected from northern Gulf and Atlantic Coast sites including sites implicated in major *V.*  
88 *vulnificus* infection outbreaks yielding values of  $3.22 \pm 0.60$ ,  $2.01 \pm 1.12$ , and  $-0.29 \pm 0.51$  for  
89 the warm (Jun-Sep), transition (Apr-May, Oct-Nov) and cold (Dec-Mar) season,  
90 respectively. The USFDA Gulf Coast Seafood Laboratory described the time for the  
91 unrefrigerated oyster harvest and transportation time to processing plants using beta-  
92 PERT distributions (Table 1) (2). Data for Louisiana, Alabama, Texas and Florida (1000  
93 values for each state ) were generated using the Excel Add-in OpenPERT.xlam  
94 (<https://code.google.com/p/openpert/downloads/list>). These values were then grouped  
95 into cold, warm and transition season and used with the same Excel Add-in to find the *min*,  
96 *max* and *ml* values for each season in the four states. During harvest and transportation,  
97 oysters are exposed to air temperatures slightly higher than seawater. Since this difference

98 is small (1.6-3.3°C) (see p. 34, 1), oysters were assumed to be at air temperature with  
 99 normal distribution values of 13.1±4.3, 23.3±4.1, 27.2±2.0 and 16.4±5.5°C for Winter,  
 100 Spring, Summer and Fall, respectively, reported in the National Oceanic and Atmospheric  
 101 Administration/National Data Buoy Center database (1). As before, 1000 generated  
 102 temperature values were grouped to calculate the warm, transition and cold season  
 103 temperature values used in this study. Next, the *V. vulnificus* load for oysters arriving at the  
 104 processing plant ( $\text{Log}(N_1, \text{CFU/g})$ ) was estimated using Eq. (1) where  $\mu_m(T)$  represents the  
 105 growth rate at a random temperature  $T(^{\circ}\text{C})$  obtained from the seasonal air temperature  
 106 distribution,  $t(h)$  the unrefrigerated oyster handling time at this temperature  $T$ , and  $A$   
 107 ( $= \text{Log}(N_{1,max}, \text{CFU/g oyster}) = 6$ ) the maximum *V. vulnificus* counts possible in raw oysters  
 108 (1, 7). The temperature dependence of the growth rate  $\mu_m(T)$  in Eq. (1) was described by  
 109 Eq. (2) where  $T$  is the seasonal air temperature,  $k (= 0.011 \text{ Log}(\text{CFU})/(\text{h } ^{\circ}\text{C}))$  is the *V.*  
 110 *vulnificus* growth rate above  $T_0$ , and  $T_0 (= 13^{\circ}\text{C})$  a threshold temperature below which *V.*  
 111 *vulnificus* does not grow (see p. 32, 1, 13).

$$\text{Log}\left(N_1, \frac{\text{CFU}}{g_{\text{oyster}}}\right) = \min\left(\text{Log}\left(N_0, \frac{\text{CFU}}{g_{\text{oyster}}}\right) + \mu_m(T) \cdot t, A\right) \quad (1)$$

$$\mu_m(T) = \max(0, k(T - T_0)) \quad (2)$$

112 Data on the *V. vulnificus* load reduction by depuration at 15°C obtained by Chae et al. (9) by

113 sampling inoculated oysters every 24 h (Table 2a) was used to generate 1000 random pathogen  
 114 load values at 0, 24, 48, 72 and 96 h. Quadratic models fitted to these values were used to estimate  
 115 the *V. vulnificus* load after a 0 to 96 h depuration time ( $t_{depuration}$ ). The difference between the initial  
 116 *V. vulnificus* load and that after depuration was used to estimate decimal reduction (*SV*) values  
 117 achieved during that time for each of the 1,000 randomly generated datasets. *SV* values were then  
 118 used to estimate the microbial load after depuration ( $Log(N_2, MPN/g\ oyster)$ ) (Eq. 3).

$$Log(N_2, MPN / g_{oyster}) = Log(N_1, MPN / g_{oyster}) - SV_{depuration} \quad (3)$$

119 Cooling time to refrigeration temperature from depuration at 15°C was assumed short and  
 120 ignored in this study. Under refrigeration, the *V. vulnificus* die-off rate is 0.041 log CFU/day and  
 121 the refrigerated time before consumption follows a beta-PERT distribution with *min*, *ml* and *max*  
 122 values of 1, 6 and 21 d (14). The lognormal distribution for oyster meat weight reported by the  
 123 Interstate Shellfish Sanitation Conference (ISSC)/FDA is  $Log(W, g/oyster) = 1.18 \pm 0.15$  (1, 2). A  
 124 published survey of metropolitan areas within 100 miles of Cedar Key in Florida encompassing 5  
 125 million residents generated data for 306 oyster consumption events (15). Random sampling of this  
 126 non-parametric distribution (numbers (frequency) = 1, 2, 3(9x), 4(10x), 5(14x), 6(61x), 7, 8(11x),  
 127 10(15x), 12(95x), 13, 15(5x), 17, 18(8x), 20(8x), 24(37x), 25(5x), 30(3x), 36(7x), 40(3x), 45(2x),  
 128 48(4x), 50(3x), 60) was used in this study.

129 The shape and scale parameter  $\alpha$  ( $= 9.3 \times 10^{-6}$ ) and  $\beta$  ( $= 1.1 \times 10^5$ ) of the Beta distribution dose-



130 response model (Eq. 4), frequently used to estimate the *V. vulnificus* septicemia risk probability  
131 ( $P_{ill}$ ) for a population ingesting a given pathogen dose ( $D$ ) per serving published by the World  
132 Health Organization (WHO, 1), incorporate pathogenicity heterogeneity, i.e., not every ingested  
133 microorganism survives to cause an infection.

$$P_{ill} = 1 - \left(1 + \frac{D}{\beta}\right)^{-\alpha} \quad (4)$$

134 Safety risk was expressed as the number of septicemia cases in 100 million oyster  
135 consumption events. Estimations were divided into 8 steps repeated 1,000 times to obtain  
136 the number of infection cases with 95% confidence (31): (1) *V. vulnificus* oyster load at  
137 harvest ( $\text{Log}(N_0, \text{CFU/g})$ ); (2) *V. vulnificus* load for oysters arriving to processing plants  
138 ( $\text{Log}(N_1, \text{CFU/g})$ ); (3)  $SV$  value reached after a given depuration time ( $SV_{depuration}$ );  
139 (4) *V. vulnificus* load after depuration ( $\text{Log}(N_2, \text{CFU/g})$ ); (5) *V. vulnificus* load at  
140 consumption ( $\text{Log}(N_3, \text{CFU/g})$ ); (6) *V. vulnificus* dose per serving ( $D, \text{CFU/serving}$ ); (7)  
141 infection probability ( $P_{ill}$ ); and, (8) number of infection cases per 100 million consumption  
142 events ( $N, \text{cases}$ ). In each step, parameter values were randomly generated using their  
143 normal, lognormal or beta-PERT distribution or by random sampling (number of oysters  
144 per consumption event). In the depuration step, 1,000 sets of *V. vulnificus* load randomly  
145 generated using their lognormal distribution were fitted into quadratic expressions for  $SV$   
146 values after depuration times between 0 and 96 h. *V. vulnificus* load as a function of

147 depuration time, season, and handling step was analyzed by ANOVA tests. As in previous  
148 studies (30), a health risk of 100 cases per 100 million oyster consumption events at 95%  
149 confidence was used to recommend a depuration time.

## 150 RESULTS

151 The estimated *V. vulnificus* growth rate during harvest and transportation to processing  
152 plants was  $0.159 \pm 0.022$ ,  $0.074 \pm 0.047$ , and  $0.022 \pm 0.030$  *Log (MPN/g oyster)/h* in the warm,  
153 transition, and cold season, respectively, and combining these values with random harvest and  
154 transportation time values, the estimated pathogen load increase was  $1.21 \pm 0.28$ ,  $0.57 \pm 0.38$ , and  
155  $0.23 \pm 0.32$  *Log (MPN/g oyster)*, respectively. Combining the latter values with the harvest load  
156 variability yielded pathogen load values before depuration (*Log N<sub>1</sub>*) of  $4.41 \pm 0.28$ ,  $2.59 \pm 0.38$   
157 and  $-0.09 \pm 0.32$  *Log (MPN/g oyster)*, respectively.

158 SV values achieved by 24, 48, 72 and 96h depuration at 15°C using quadratic models  
159 (average  $R^2 = 0.95$ ) were  $1.55 \pm 0.17$ ,  $2.56 \pm 0.24$ ,  $3.03 \pm 0.24$ , and  $2.95 \pm 0.24$  *Log (MPN/g oyster)*,  
160 respectively, showing that after 72 h the pathogen load reduction is not significant ( $P\text{-value} > 0.05$ ).  
161 Microbial load values after depuration are summarized in Table 2b.

162 *V. vulnificus* die-off during refrigerated handling time before consumption was estimated as  
163  $0.31 \pm 0.15$  *Log (MPN/g oyster)* yielding pathogen loads for untreated oysters at consumption

164 of  $-0.38 \pm 0.6$  and  $4.14 \pm 0.7 \text{ Log (MPN/g oyster)}$  in cold and warm seasons, respectively, and a  
165 significantly lower value for the transition season ( $2.26 \pm 1.2 \text{ Log (MPN/g oyster)}$ ) when compared  
166 to the warm season (P value < 0.05). The season effect is reflected also in *V. vulnificus* loads for  
167 raw oyster treated by depuration for up to 96 h (Table 2c). The pathogen dose consumed per  
168 oyster serving (*D*) reflects the variability in the number of oysters consumed per serving, oyster  
169 weight, and pathogen load after depuration. In the case of untreated oysters, the dose of  $(10.0 \pm 3.1)$   
170  $\times 10^6 \text{ MPN/serving}$  in the warm season was significantly higher (P value < 0.05) than the  $(9.7 \pm 0.4)$   
171  $\times 10^5$  and  $283.1 \pm 61.7 \text{ MPN/serving}$  in the transition and cold season, respectively. Dose values  
172 after depuration for 24 to 96 h are shown in Table 2d.

173 At 95% confidence, the estimated number of infection cases in 100 million consumption  
174 events for untreated oysters in the warm season was 2669 cases, while 24 and 48h depuration  
175 would reduce it to 558 and 93 cases, respectively (Table 3). The depuration time required to reach  
176 the safety target used in this study (100 cases per 100 million consumption events) was estimated  
177 to be 47h. The discrepancy between the estimated 93 and 100 cases after 48 and 47h depuration  
178 time, respectively, reflects the non-deterministic procedure used in this study. In the transition  
179 season, the number of predicted cases would be 491 for untreated oysters, reaching the 100 cases  
180 safety-target after 16h depuration, meaning that the 44h recommended depuration (see p. 126, 3)  
181 could be considered over-processing. Only 35 cases would be observed after depuration for 24h,

182 while a 48h depuration is likely to reduce the number of infection cases to 4. The number of cases  
183 at 95% confidence that could be expected after the recommended 44h depuration time would be  
184 135, 8 and 0, for oysters harvested in the warm, transition and cold season, respectively (Table 3).

## 185 **DISCUSSION**

186 The U.S. FDA requires that “the dealer must demonstrate that the process reduces the level  
187 of *Vibrio vulnificus* and/or *Vibrio parahaemolyticus* ... to non-detectable (<30 MPN/gram) and that  
188 the process achieves a minimum 3.52 log reduction (see p. 140, 3).” Furthermore, *V. vulnificus* and  
189 *V. parahaemolyticus* levels must be determined following the sampling protocol (4) and microbial  
190 enumeration (see p. 345, 3) described by the National Shellfish Sanitation Program (NSSP). The  
191 USFDA/CFSAN and the ISSC state that treated oysters meeting these specified endpoint and  
192 decimal reduction levels can be labeled as "Processed to reduce *Vibrio vulnificus* to non-detectable  
193 levels" (see p. 187, 3). The 3.52 log reduction in these regulations is based on assuming extremely  
194 high *V. vulnificus* or *V. parahaemolyticus* loads observed sometimes in the Gulf Coast during  
195 summer months (100,000 MPN/g) being lowered by processing to reach non-detectable levels  
196 (<30 MPN/g) (see p. 172, 3). Although artificial seawater depuration tests showed that reduction  
197 in *V. vulnificus* counts leveled off after 48 h and that 96 h depuration cannot achieve 3.52 log  
198 reductions for (9), use of this experimental data showed that depuration would achieve large

199 reductions in the *V. vulnificus* infection risk reaching values below 100 cases per 100 million  
200 consumption events (Table 3).

201         The probability that depuration would reduce the *V. vulnificus* load to the non-detectable  
202 endpoint (<30 MPN/g) was also determined (Table 4). A 95% confidence level is typically used in  
203 process design calculations (26, 33). Oysters harvested in the cold season would meet this  
204 requirement at 95% confidence in less than 1h while in the transition and warm season it cannot  
205 be met even after 96h depuration. If pathogen loads after depuration ( $N_2$ ) are lower than the non-  
206 detectable 30 MPN/g, the estimated infection risk at 95% confidence per 100 million consumption  
207 events estimated using Eq. (4) would be less than 27 cases per year (calculations not shown).

208         An analysis of the effectiveness of the recommended depuration time (44 h) showed that  
209 only 23% and 50% of oysters harvested in the warm and transition seasons, respectively, would  
210 reach the non-detectable *V. vulnificus* level (i.e., below 30 MPN/g), while 100% would reach it in  
211 the cold season (Table 4). At 95% confidence, the use of the recommended 44 h depuration time  
212 would result in 135, 8, and 0 cases in the warm, transition, and cold season, respectively (Table 3).  
213 Thus, in the summer season it would be too short while for other seasons it would be too long. The  
214 microbial risk analysis of HPP-treated oysters completed by Serment-Moreno et al. (31) showed  
215 that 4 cases of *V. vulnificus* infection cases (95% confidence) per 100 million consumption events  
216 could be expected from the consumption of oysters harvested in the warm season if treated at 250

217 MPa for 2 min at 1°C with no cases expected in other seasons.

218           The number of oysters consumed per serving was obtained from central and north-central  
219 Florida surveys (15) where the number consumed per oyster serving could be higher than in  
220 inland states. Population inference errors may also occur when consumers recall consumption  
221 events (15) and consumers eating oysters more frequently may tolerate higher pathogen loads  
222 (28). Moreover, *V. vulnificus* in oysters were assumed to correspond to equally virulent strains  
223 (16). Although samples below detectable level were assigned a half-way value between 0 and the  
224 detection value, this can be ignored since the number of oyster samples below detection level was  
225 reported to be small (i.e., 2 of 24 samples, 24).

226           This study used reported experimental data (8) focusing on *V. vulnificus* oyster depuration.  
227 The same procedure could be applied to reduce the risks of other pathogens by depuration, or to  
228 analyze other oyster treatment technologies by modifying only the SV estimation. The procedure  
229 would be most effective when using data for individual processors or at least individual harvest  
230 regions. This would reduce statistical variability and lower the treatment intensity recommended  
231 (depuration time/temperature, pressure level/holding time, etc.). The positive effect of reducing  
232 statistical variability has been previously shown (10, 29).

233           In conclusion, the *V. vulnificus* infection risk associated with raw oyster consumption was  
234 quantified to estimate a recommended depuration time. The 44h depuration set independently of

235 oyster harvest season, non-detectable endpoint (<30 MPN/g), and 3.52 *V. vulnificus* decimal  
236 reduction, were analyzed using a Monte Carlo protocol. The analysis included the variability of the  
237 seasonal oyster pathogen load at the point of harvest, the time and local temperature during  
238 oyster harvest and transportation to oyster processing facilities, time that oysters are kept  
239 refrigerated after processing and before consumption, and the size and number of oysters  
240 consumed per serving. For untreated oysters, the *V. vulnificus* infection risk at 95% confidence is  
241 exceedingly low in the cold season but unacceptably high in the warm and transition season. An  
242 acceptable risk, defined in this study as 100 cases per 100 million consumption events, could be  
243 achieved with 95% confidence by oyster depuration for 47 and 16 h in the warm and transition  
244 season, respectively.

## 245 **ACKNOWLEDGMENTS**

246 The authors thank critical reviews by PhD Candidates M.C. Rosas González and L.E. García  
247 Amézquita. Financial support from the Tecnológico de Monterrey (Research Chair Funds CAT-200)  
248 for the Emerging Technologies and Nutrigenomic Research Groups, Fondo Nacional de Ciencia y  
249 Tecnología de Chile (FONDECYT Grant 1110569), Research Office of Universidad Andres Bello (DI-  
250 275-13/R 2013), Fondo de Fomento al Desarrollo Científico y Tecnológico (FONDEF) CA13I10077,  
251 and Formula Grants no. 2011-31200-06041 and 2012-31200-06041 from the USDA National  
252 Institute of Food and Agriculture is gratefully acknowledged.

## References

- 254 1. Anonymous. 2005. Risk assessment of *Vibrio vulnificus* in raw oysters. Interpretative  
255 summary and technical report. World Health Organization, Food and Agriculture Organization of  
256 the United Nations, Rome, Italy.
- 257 2. Anonymous. 2005. *Vibrio parahaemolyticus* risk assessment – Appendix 5: Details of the  
258 data analysis for exposure assessment. Center for Food Safety and Applied Nutrition, Food and  
259 Drug Administration, U.S. Department of Health and Human Services, Washington, DC.
- 260 3. Anonymous. 2011. Guide for the control of molluscan shellfish. U.S. Food and Drug  
261 Administration, Washington, DC.
- 262 4. Anonymous. 2011. Section IV. Guidance Documents, Chapter IV. Naturally Occurring  
263 Pathogens. In, National Shellfish Sanitation Program (NSSP) Guide for the Control of Molluscan  
264 Shellfish: 2011 Revision Interstate Shellfish Sanitation Conference and the Center for Food Safety  
265 and Applied Nutrition from the US Food and Drug Administration, College Park, MD.
- 266 5. Anonymous. 2012. Cholera and other *Vibrio* illness surveillance overview. Centers for  
267 Disease Control and Prevention, CDC, U.S. Department of Health and Human Services, Atlanta,  
268 Georgia.
- 269 6. Boxman, I. L. A., J. J. H. C. Tilburg, N. A. J. M. te Loeke, H. Vennema, K. Jonker, E. de Boer, and  
270 M. Koopmans. 2006. Detection of noroviruses in shellfish in the Netherlands. *Int J Food Microbiol.*  
271 108:391-396.
- 272 7. Buchanan, R. L., W. G. Damert, R. C. Whiting, and M. Van Schothorst. 1997. Use of  
273 epidemiologic and food survey data to estimate a purposefully conservative dose-response  
274 relationship for *Listeria monocytogenes* levels and incidence of listeriosis. *J Food Prot.* 60:918-922.
- 275 8. Chae, M. 2007. Low-temperature post-harvest processing for reducing *Vibrio*  
276 *parahaemolyticus* and *Vibrio vulnificus* in raw oysters In, Food Science and Technology vol. Master  
277 of Science. Oregon State University, Corvallis.
- 278 9. Chae, M., D. Cheney, and Y.-C. Su. 2009. Temperature effects on the depuration of *Vibrio*  
279 *parahaemolyticus* and *Vibrio vulnificus* from the American oyster (*Crassostrea virginica*). *J Food Sci.*  
280 74:M62-M66.
- 281 10. Chotyakul, N., C. Pérez-Lamela, and J. A. Torres. 2011. Effect of model parameter variability  
282 on the uncertainty of refrigerated microbial shelf-life estimates. *J Food Proc Eng.* 35:829-839.



- 283 11. Chotyakul, N., G. Velazquez, and J. A. Torres. 2011. Assessment of the uncertainty in thermal  
284 food processing decisions based on microbial safety objectives. *J Food Eng.* 102:247-256.
- 285 12. Christensen, E. R., and C.-Y. Chen. 1985. A General noninteractive multiple toxicity model  
286 including Probit, Logit, and Weibull transformations 41:711-725.
- 287 13. Cook, D. W. 1994. Effect of time and temperature on multiplication of *Vibrio vulnificus* in  
288 postharvest Gulf Coast shellstock oysters. *Appl Environ Microbiol.* 60:3483-3484.
- 289 14. Cook, D. W., P. O. O'Leary, J. C. Hunsucker, E. M. Sloan, J. C. Bowers, R. J. Blodgett, and A. de  
290 Paola. 2002. *Vibrio vulnificus* and *Vibrio parahaemolyticus* in U.S. retail shell oysters: A national  
291 survey from June 1998 to July 1999. *J Food Prot.* 65:79-87.
- 292 15. Degner, R. L., and C. Petrone. 1994. Consumer and restaurant manager reaction to  
293 depurated osysters and clams. Institute of Food and Agricultural Sciences, University of Florida,  
294 Gainesville, FL.
- 295 16. DePaola, A., J. L. Nordstrom, A. Dalsgaard, A. Forslund, J. Oliver, T. Bates, K. L. Bourdage, and  
296 P. A. Gulig. 2003. Analysis of *Vibrio vulnificus* from market oysters and septicemia cases for  
297 virulence markers *Appl Environ Microbiol.* 69:4006-4011.
- 298 17. Formiga-Cruz, M., G. Tofino-Quesada, S. Bofill-Mas, D. N. Lees, K. Henshilwood, A. K. Allard,  
299 A. C. Conden-Hansson, B. E. Hernroth, A. Vantarakis, A. Tsibouxi, M. Papapetropoulou, M. D.  
300 Furones, and R. Girones. 2002. Distribution of human virus contamination in shellfish from  
301 different growing areas in Greece, Spain, Sweden, and the United Kingdom. *Appl Environ Microbiol.*  
302 68:5990-5998.
- 303 18. Holcomb, D. L., M. A. Smith, G. O. Ware, Y. C. Hung, R. E. Brackett, and M. P. Doyle. 1999.  
304 Comparison of six dose - response models for use with food - borne pathogens. *Risk Anal.*  
305 19:1091-1100.
- 306 19. Janda, J., C. Powers, R. Bryant, and S. Abbott. 1988. Current perspectives on the  
307 epidemiology and pathogenesis of clinically significant *Vibrio* spp. *Clin Microbiol Rev.* 1:245-267.
- 308 20. Kaspar, C. W., and M. L. Tamplin. 1993. Effects of temperature and salinity on the survival of  
309 *Vibrio vulnificus* in seawater and shellfish. *Appl Environ Microbiol.* 59 2425-2429.
- 310 21. Lee, R., A. Lovatelli, and L. Ababouch. 2008. Bivalve depuration: Fundamental and practical  
311 aspects. Food and Agriculture Organization of the United Nations, Rome, Italy.
- 312 22. Lewis, M., S. Rikard, and C. R. Arias. 2010. Evaluation of a flow-through depuration system  
313 to eliminate the human pathogen *Vibrio Vulnificus* from oysters. *J Aquacult Res Dev.* 1:2.

- 314 23. Mead, P. S., L. Slutsker, P. M. Griffin, and R. V. Tauxe. 1999. Food-related illness and death in  
315 the United States. *Emerg Infect Dis.* 5:607-625.
- 316 24. Motes, M. L., A. de Paola, D. W. Cook, J. E. Veazey, J. C. Hunsucker, W. E. Garthright, R. J.  
317 Blodgett, and S. J. Chirtel. 1998. Influence of water temperature and salinity in *Vibrio vulnificus* in  
318 Northern Gulf and Atlantic Coast oysters. *Appl Environ Microbiol.* 64:1459-1465.
- 319 25. Nishibuchi, M., and A. DePaola. 2005. *Vibrio species.* p. 251–271. In P.M. Fratamico, A.K.  
320 Bhunia, and J.L. Smith (ed.), Foodborne pathogens: Microbiology and molecular biology Caister  
321 Academic Press, Norfolk, U K.
- 322 26. Rieu, E., K. Duhem, E. Vindel, M. Sanaa, and A. Theobald. 2007. Food safety objectives  
323 should integrate the variability of the concentration of pathogen. *Risk Anal.* 27:373-386.
- 324 27. Ritz, C. 2010. Toward a unified approach to dose-response modeling in ecotoxicology.  
325 *Environ Toxicol Chem.* 29:220-229.
- 326 28. Saavedra, J. M., A. Abi-Hanna, N. Moore, and R. H. Yolken. 2004. Long-term consumption of  
327 infant formulas containing live probiotic bacteria: tolerance and safety. *Am J Clin Nutr.* 79:261-267.
- 328 29. Salgado, D., J. A. Torres, J. Welti-Chanes, and G. Velazquez. 2011. Effect of input data  
329 variability on estimations of the equivalent constant temperature time for microbial inactivation  
330 by HTST and retort thermal processing. *J Food Sci.* 76:E495-E502.
- 331 30. Serment-Moreno, V., G. Barbosa-Cánovas, J. A. Torres, and J. Welti-Chanes. 2014. High  
332 pressure processing: Kinetic Models for Microbial and Enzyme Inactivation. *Food Eng Rev.* 6:56-  
333 88.
- 334 31. Serment-Moreno, V., K. Deng, Y.-C. Su, X. Wu, C. Fuentes, J. A. Torres, and J. Welti Chanes.  
335 2015. Monte Carlo analysis of the product handling and high-pressure treatment effects on the  
336 *Vibrio vulnificus* risk to raw oysters consumers. *J Food Eng.* 144:86–92.
- 337 32. Serment-Moreno, V., H. Mújica-Paz, J. A. Torres, and J. Welti-Chanes. 2012. Aplicación del  
338 método de Monte Carlo para simular la inactivación de pectinmetilesterasa (PME) en jugo de  
339 naranja con procesos combinados de altas presiones hidrostáticas (APH) y temperatura. *Rev Mex*  
340 *Ing Quim.* 11:363-372.
- 341 33. Smout, C., A. van Loey, M. E. Hendrickx, and J. Beirlant. 2000. Statistical variability of heat  
342 penetration parameters in relation to process design. *J Food Sci.* 65:685-693.

343  
344

**Table 1.** Harvest and transportation time (hours) to processing plants in the U.S. Gulf Coast

Location		Winter (Jan-Mar)	Spring (Apr-Jun)	Summer (Jul-Sept)	Fall (Oct-Dec)
Louisiana	<i>max</i> =	13	11	11	13
	<i>min</i> =	7	5	5	7
	<i>ml</i> =	12	9	9	12
Alabama, Texas, and Florida	<i>max</i> =	11	10	10	10
	<i>min</i> =	2	3	3	3
	<i>ml</i> =	8	7	7	7

345

**Table 2.** *Vibrio vulnificus* load and consumption dose of untreated and depurated oyster

Sample	Untreated*	Depuration time			
		24 h	48 h	72 h	96 h
a) Reduction during depuration test at 15°C**					
Inoculated	5.52 ± 0.16	3.49 ± 0.23	2.76 ± 0.19	2.67 ± 0.29	2.23 ± 0.20
b) Load after depuration for $t = 0$ (untreated) to 96 h, $\text{Log}(N_{2,t} \text{ MPN/g oyster})$					
Warm	4.41±0.28 <sup>1</sup>	2.89±0.7	1.87±0.7	1.41±0.7	1.48±0.7
Transition	2.59±0.38 <sup>1</sup>	0.95±1.2	-0.07±1.2	-0.54±1.3	-0.46±1.2
Cold	-0.09±0.32 <sup>1</sup>	-1.63±0.6	-2.65±0.6	-3.11±0.6	-3.03±0.62
c) Load after depuration and refrigerated storage, $\text{Log}(N_3, \text{ MPN/g oyster})$					
Warm	4.14±0.7	2.59±0.7	1.58±0.7	1.12±0.7	1.19±0.7
Transition	2.26±1.2	0.71±1.2	-0.31 ±1.2	-0.77±1.2	-0.69±1.2
Cold	-0.38±0.6	-1.95±0.6	-2.96±0.6	-3.43±0.6	-3.35±0.6
d) Dose when consuming raw oysters, $D, \text{ MPN/serving}$					
Warm	10.0±3.1x10 <sup>6</sup>	3.0±0.29x10 <sup>5</sup>	3.1±0.53x10 <sup>4</sup>	1.1±0.43x10 <sup>4</sup>	1.4± 0.16x10 <sup>4</sup>
Transition	9.7±0.4x10 <sup>5</sup>	3.3±0.04x10 <sup>4</sup>	3.6± 0.07 x10 <sup>3</sup>	1.1±0.06 x10 <sup>3</sup>	1.3±0.02 x10 <sup>3</sup>
Cold	283.1±61.7	8.0±0.57	0.83±0.11	0.29±0.09	0.37±0.03

(\*) Untreated load corresponds to the Log N1 value obtained in the previous calculation step

(\*\*) Data obtained from Chae et al. (9)

346

347

**Table 3.** Probable number of septicemia infection cases by consuming raw oysters treated by depuration for t = 0 (untreated) to 96 h and recommended depuration time ( $t_{depuration}$ )<sup>1</sup>

Season	Untreated	Depuration time					$t_{depuration}$ (h)
		24 h	48 h	72 h	96 h	44 h <sup>2</sup>	
Warm	2669	558	93	38	47	135	47
Transition	491	35	4	1	2	8	16
Cold	1	0	0	0	0	0	0 <sup>3</sup>

(1) depuration time to reach N = 100 cases per 10<sup>8</sup> consumption events

(2) # of cases at 95% confidence expected after the recommended 44h depuration time

(3) depuration treatment is not recommended for cold season

**Table 4.** Probability that the *V. vulnificus* load in raw oysters is reduced to <30 MPN/g by depuration for  $t = 0$  to 96 h and recommended depuration time ( $t_{30MPN/g}$ ) to reach this load with 95% confidence

Season	Untreated	Depuration time					$t_{30MPN/g}$
		24 h	48 h	72 h	96 h	44 h <sup>1</sup>	
Warm	0%	3%	31%	55%	50%	23%	>96 h
Transition	17%	66%	88%	95%	94%	50%	72 h
Cold	99%	100%	100%	100%	100%	100%	0 h

(1) Currently recommended depuration time