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GROWTH OF MIXED CULTURES OF *LISTERIA* AND *PSEUDOMONAS* SPP. IN BEEF MEAT AND THEIR PREDICTION USING POLYNOMIAL MODELS

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Background

Among the psychrotrophic Gram-negative bacteria that can multiply in meat products, dairy products and vegetables, only a few have an influence on the quality and the shelf-life of these products and can be responsible for considerable financial losses. The principal flora responsible for such spoilage during aerobic storage are the pseudomonads. *Pseudomonas* spp. grew faster than other strains present in meat at temperatures between 2 and 15°C. Several pathogenic bacteria also grow within this temperature range and may cause foodborne outbreaks: *Listeria monocytogenes* has been shown to be responsible for outbreaks of food poisoning in France and in many countries (Rocourt and Bille, 1997).

Objective

To predict the growth of both spoilage Pseudomonas spp. and pathogenic Listeria in meat products

Methods

A first set of experiments were performed in decontaminated beef meat that was inoculated with *Listeria* or *Pseudomonas* strains. Strains were grown alone or together. The second set of experiments was performed in naturally contaminated meat that was purchased at the supermarket, meats were available in oxygen-permeable bags (experiments B and D) or in vacuum packs (experiments A and C) (Figure 1). The dominant *Pseudomonas* spp. were grown alone or in the presence of inoculated *Listeria*. Meat was stored under aerobic conditions at low temperatures. Bacterial counts were performed on: Tryptone Soja Agar (Difco, OSI, Maurepas, France) for non selective growth; on Palcam Agar (Merck, Nogent-sur-Marne, France) for *Listeria*; on *Pseudomonas* Agar (Oxoïd, Unipath Ltd, Basingstocke, England) for *Pseudomonas*.

Growth parameters (A, logarithmic increase in the bacterial population, L, lag time, GT, generation time) were calculated by fitting the growth curves using a modified Gompertz equation (Zwietering et al., 1990). The *Listeria* model of Lebert et al. (1998) and the three *Pseudomonas* models of Robles (1999) were used as they took the growth variations into account and calculate a growth response interval.

Results and discussion

Three models for one rapid and one slow growing strain of *P. fragi* and one slow growing strain of *P. fluorescens* were developed in a meat broth and tested in naturally-contaminated meat. TSA and CFC counts were similar indicating that *Pseudomonas* spp. were dominant. Table 1 shows that experimental L were longer than predicted L but that the observed GT were between the GT predicted by *Pfr*162 and *Pfr*K1 model. The *Pseudomonas* models can be used to predict the growth of *Pseudomonas* spp. in meat.

In decontaminated meat inoculated with Listeria or Pseudomonas strains (Table 2), all the models provided satisfactory predictions. In mixed population experiments (Table 2), generation times for Listeria were similar when grown alone or with P. fragi, and were predicted satisfactory by the models. A comparison of the growth parameters of Pfr162 alone and with Lm14 showed that they were similar if account was taken of the confidence intervals. When co-inoculated, the growth of neither organism was either inhibited or stimulated. No interactions were observed between the strains, and their growth parameters were satisfactorily predicted by their respective models.

In naturally-contaminated meat inoculated with Listeria, Figure 1 shows that the experimental points for Pseudomonas were within the curves generated by the three models. In the four cases, observed GT were well predicted by the models. In experiments A and C, growth was inside the curves predicted by PfrK1 and Pfl58 models (slow models). In experiments B and D, growth was near the curves predicted by Pfr162 models (fast model). A and D experiments (meat stored in vacuum packs) were characterised by longer lag times than those observed in experiments B and D (meat stored in oxygen-permeable bags): these differences can be due to the presence of different bacterial flora on the meat because of the way of storage. In meat stored in oxygen-permeable film, the microflora was already aerobic and mainly composed of *Pseudomonas* spp., that explains a small L. With vacuum-packaged meat, it took longer for the *Pseudomonas* to outgrow the anaerobic flora present in the meat. No *Listeria* growth was observed until *Pseudomonas* reached the stationary phase, slight increases (1 to 2 log(CFU)) were then observed. Predictions of *Listeria* growth were calculated for an A of 4.1 as found in Table 2 and Lm14 model failed to predict the slow growth of *Listeria*.

Conclusions

In experiments with mixed populations, three situations were observed: (1) in decontaminated meat, L. monocytogenes inoculated alone grew well at 6°C and the growth was correctly predicted by the model; (2) in decontaminated meat that was inoculated with Listeria and Pseudomonas strains, L. monocytogenes grew well and was not affected by the presence of the Pseudomonas, growth of both organisms were correctly predicted by the models; (3) in naturally contaminated meat inoculated with L. innocua, the strain did not grow until the Pseudomonas had reached the stationary phase. The models satisfactorily predicted the growth of Pseudomonas strains.

In consequence, the Lm14 model can not be used for refrigerated meat stored aerobically as it indicated a too 'fail-safe' prediction: meat had already reached a spoilage state even though no increase of *Listeria* was observed. The *Pseudomonas* models accurately predicted the growth of naturally occurring *Pseudomonas* spp.

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Pertinent literature

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<u>Table 1:</u> Observed and predicted values (lag time (L) and generation time (GT)) of *Pseudomonas* spp. grown on the surface of naturally contaminated beef muscle at a relative humidity of air \approx 100% and pH 5.8. In small prints: lower and upper confidence intervals at 95%.

T	logN ₀	Α	L obs.	L pred. (h)			GT obs.	GT pred. (h)		
(°C)	in the second		(h)	Pfr162 model	PfrK1 model	Pf158 model	(h)	Pfr162 model	PfrK1 model	Pfl58 model
6	4.4	5.8	19 (13-26)	6.7 (3.9-11.6)	3.3 (0.8-13.5)	12 (5.8-24)	5.1 (4.5-5.6)	4.3 (3.5-5.4)	6.1 (4.5-8.2)	6.5 (4.5-9.3)
6	3.0	7.5	17 (12-22)	6.7 (3.9-11.6)	3.3 (0.8-13.5)	12 (5.8-24)	4.4 (4.1-4.8)	4.3 (3.5-5.4)	6.1 (4.5-8.2)	6.5 (4.5-9.3)
4	3.0	7.4	18 (12-24)	10.3 (5.5-19.3)	4.4 (0.9-22.3)	14 (6.4-32)	5.7 (5.2-6.2)	6.2 (4.8-7.9)	8.8 (6.3-12.4)	10 (6.7-15)

<u>Table 2</u>: Observed and predicted lag times and generation times (h) for single strain or mixed cultures grown on the surface of decontaminated meat at 4°C, relative humidity $\approx 100\%$. In small prints: lower and upper confidence intervals at 95%.

Experiments	Strains	Т	pH	logN ₀	Α	L obs.	L pred. (h)		GT obs.	GT pred. (h)	
	100000000000000000000000000000000000000	(°C)				(h)	Pfr162 model	Lm14 model	(h)	Pfr162 model	Lm14 model
Listeria +	Pfr162	4	5.9	4.0	6.4	8 (4-12)	10 (6-16)	with the first standard and	5.2 (4.8-5.6)	6.1 (5.0-7.3)	tell Sin ante alm
Pseudomonas	LiCLIP	4	5.9	3.1	4.1	69 (41–97)		67 (10-446)	28 (23-33)		26 (14-49)
Listeria	<i>Lm</i> 14	4	6.3	4.1	4.8	36 (10-62)		56 (10-301)	33 (29-36)		28 (13-39)
Pseudomonas	<i>Pfr</i> 162	4	5.7	4.0	6.2	26 (20-31)	11 (5-20)		5.2 (4.5-5.8)	6.3 (4.8-8.2)	





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