



Modulagem de contaminação cruzada: teoria e aplicação

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Publication date:
2014

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Citation (APA):

Møller, C. O. D. A. (Author). (2014). Modulagem de contaminação cruzada: teoria e aplicação. Sound/Visual production (digital)

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Modulagem de contaminação cruzada: teoria e aplicação

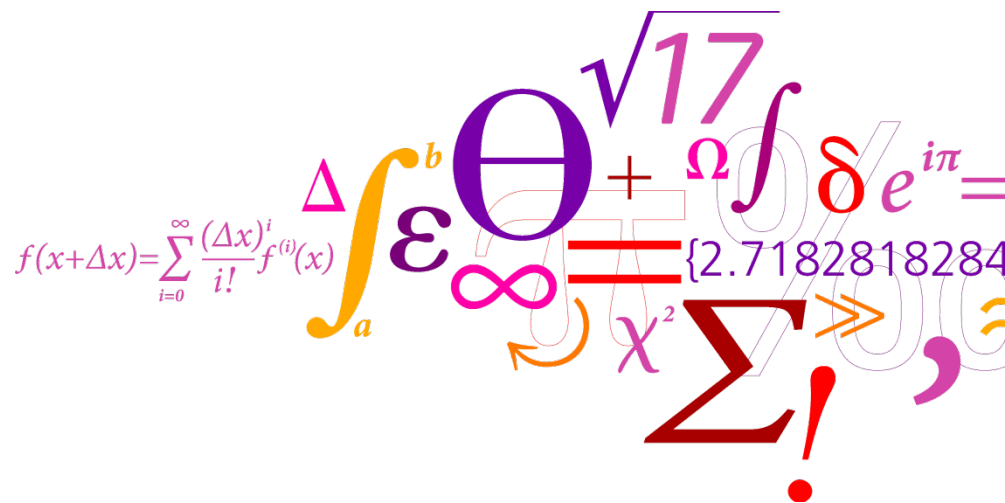
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Outline:

- Modelagem de contaminação cruzada
 - ✓ Introdução
 - ✓ Objetivos do estudo
 - ✓ Resumo do trabalho desenvolvido
 - ✓ Processo para construir o modelo
 - ✓ Resultados
 - ✓ Desafios e perspectivas
- Projeto Brasil-Dinamarca
- Aplicação

Modelling transfer of *Salmonella*

Typhimurium DT104 during grinding of pork

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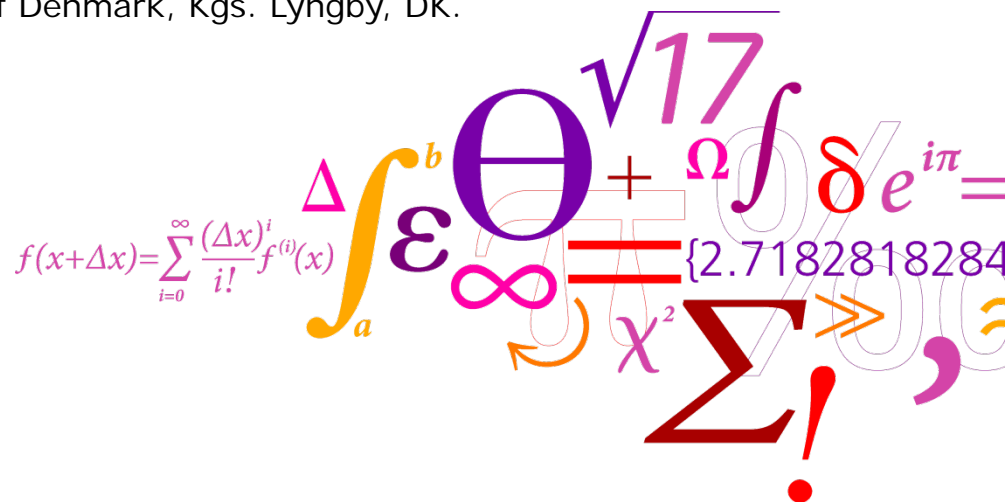
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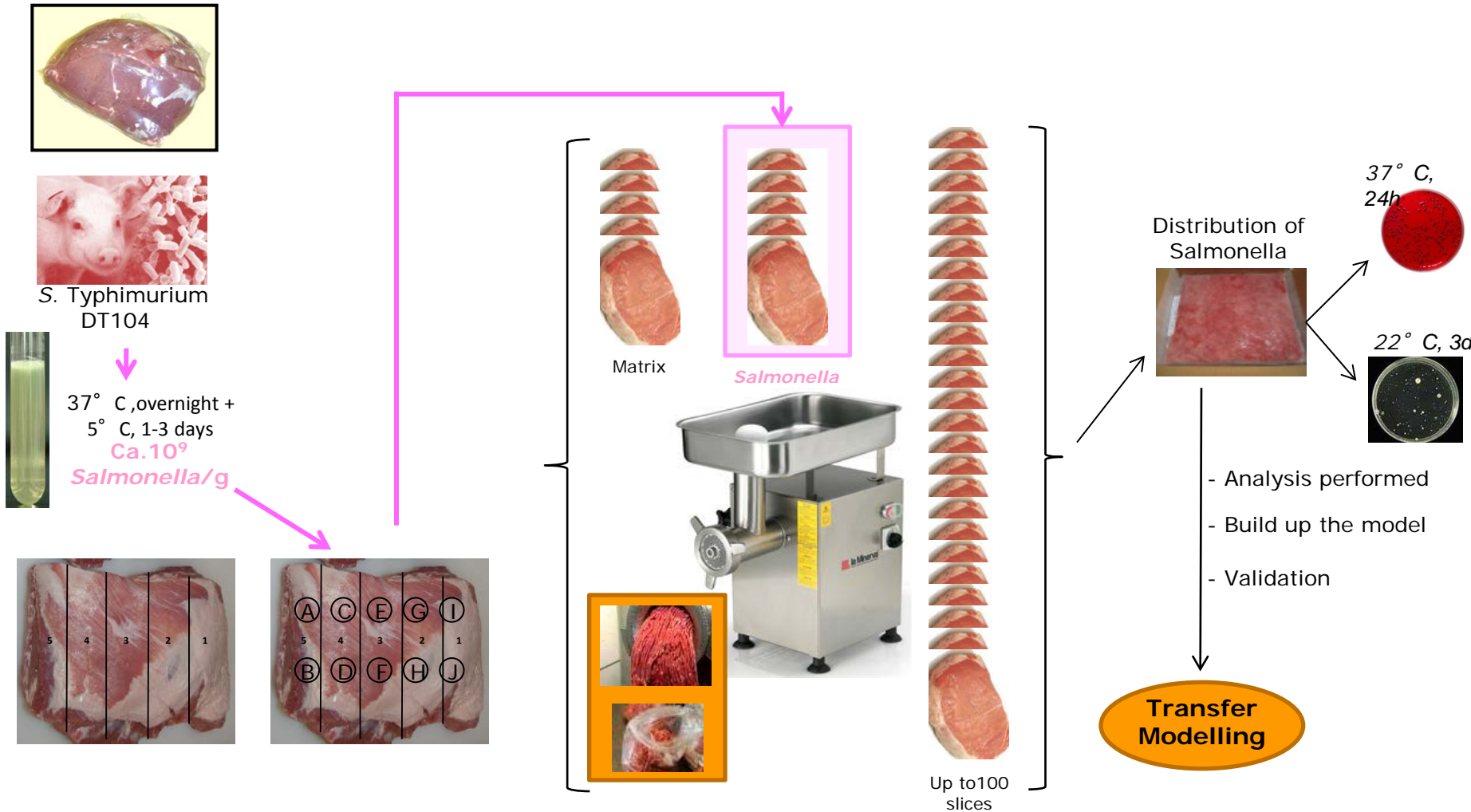


Introduction

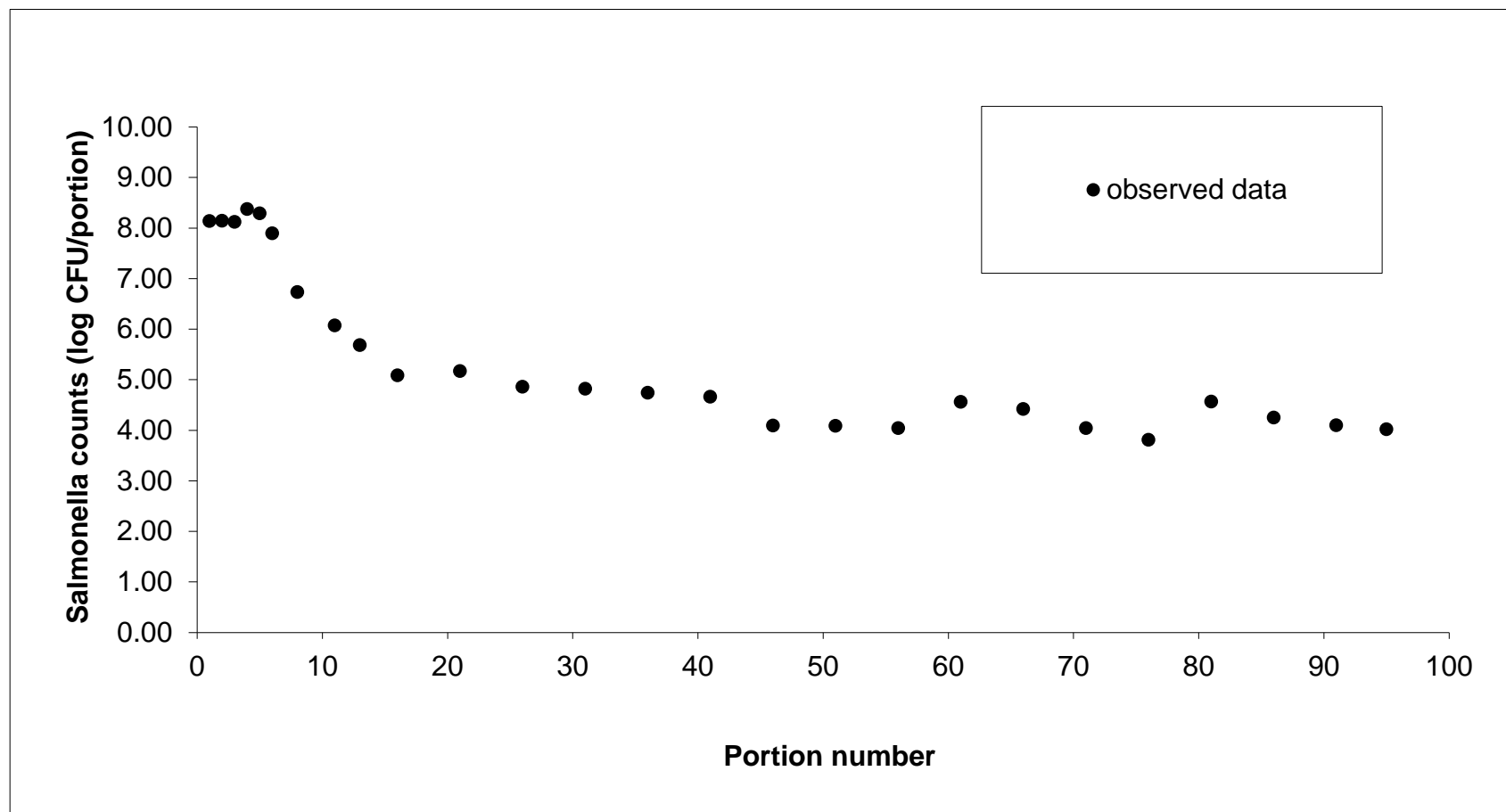
- *Salmonella* is a critical pathogen (CDC, 2011; EFSA, 2010).
- Pork still is an important source of salmonellosis (EFSA, 2010; van Hoek *et al.*, 2012; Wegener *et al.*, 2003).
- Ground meat is frequently associated with outbreaks of salmonellosis (Stock and Stolle, 2001).
- Up to 70% of foodborne illnesses are estimated to be linked to catered food (Filion and Powell, 2011; Hensen *et al.*, 2006; Jones *et al.*, 2004; Lee and Middleton, 2003).
- In Denmark, 61 of 86 reported outbreaks in 2011 were associated with outside-the-home settings (anonymous, 2012).
- To model the distribution of pathogens during the processing operation are of major relevance to risk analysts (Flores, 2006).

Objective

The aim of this study was to develop a model able to predict cross contamination of *Salmonella* in pork grinding.

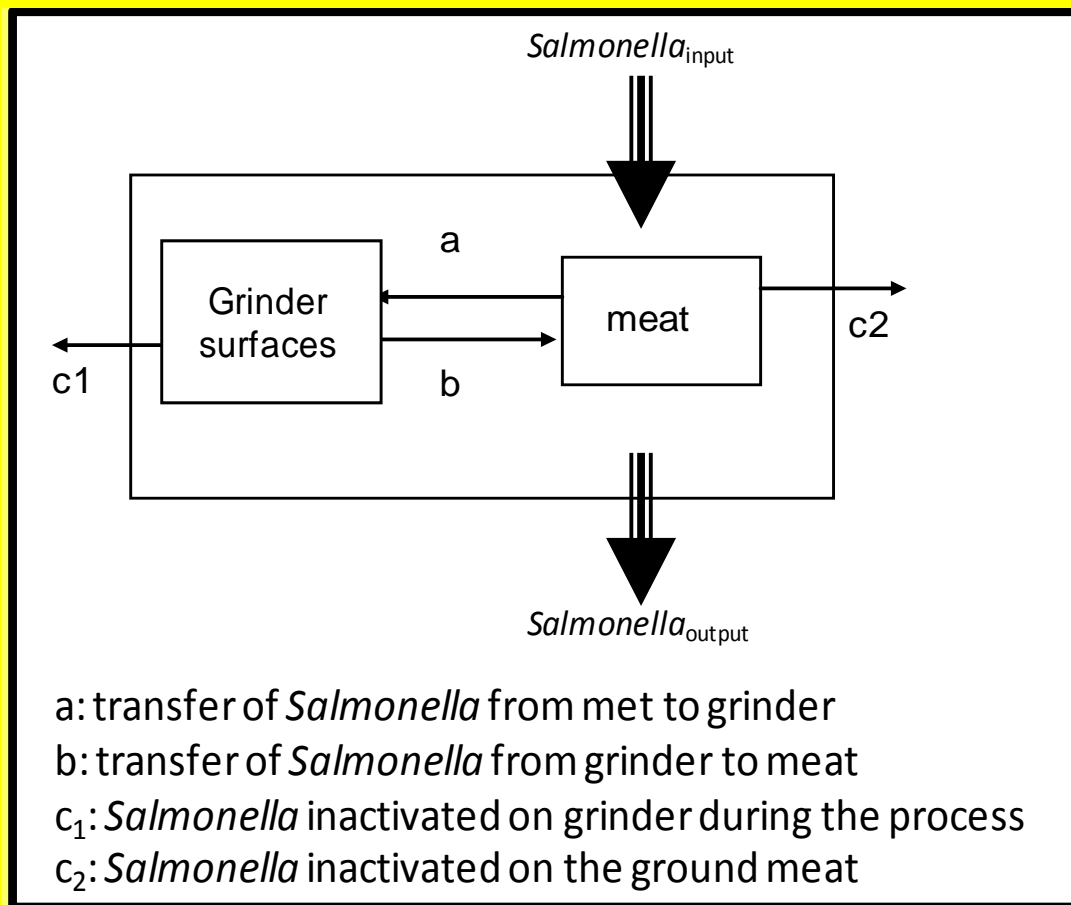


Describing the transfer rates of *Salmonella* during pork grinding

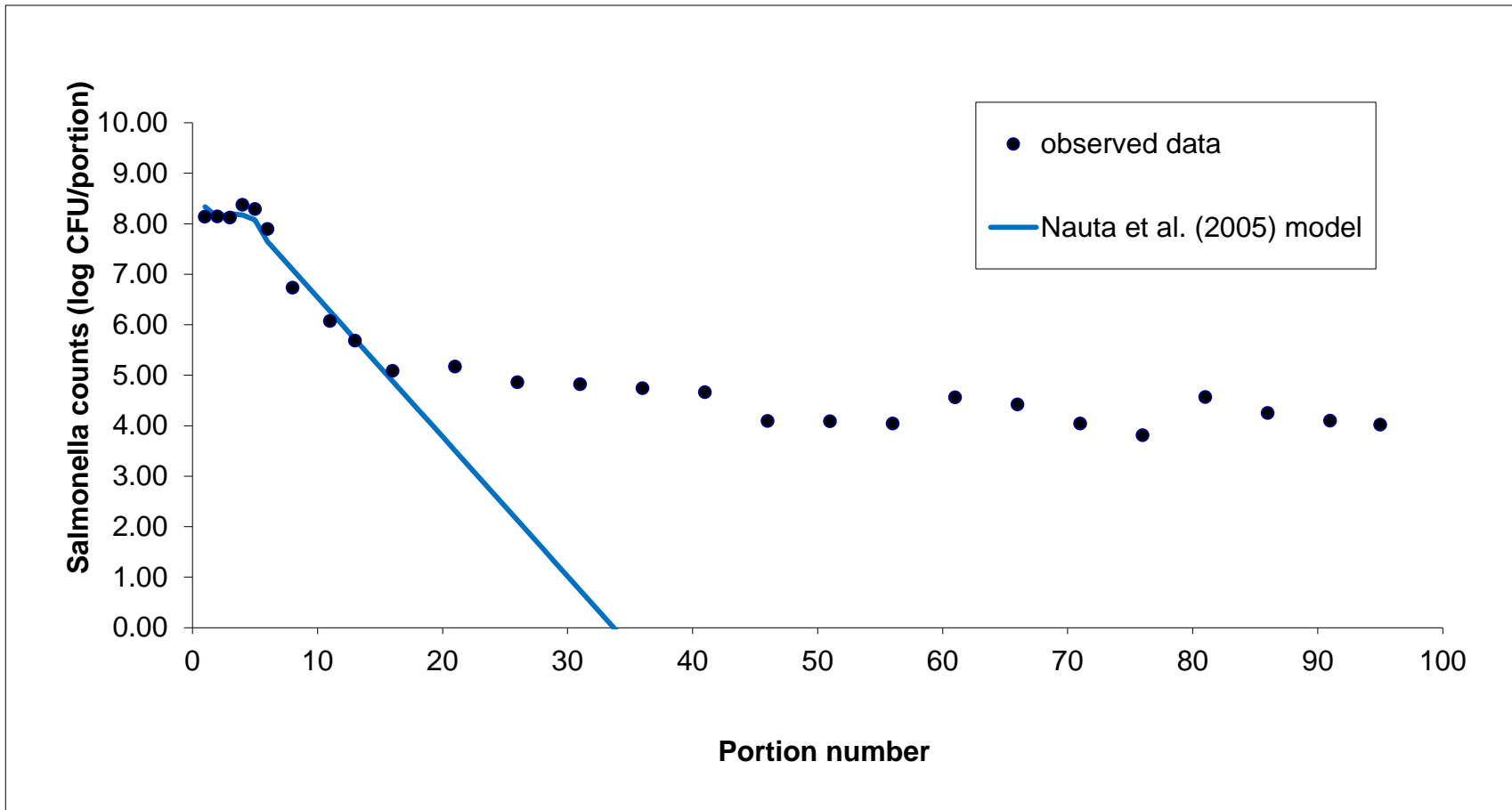


Modelling cross-contamination during pork grinding

Nauta et al. (2005) Model



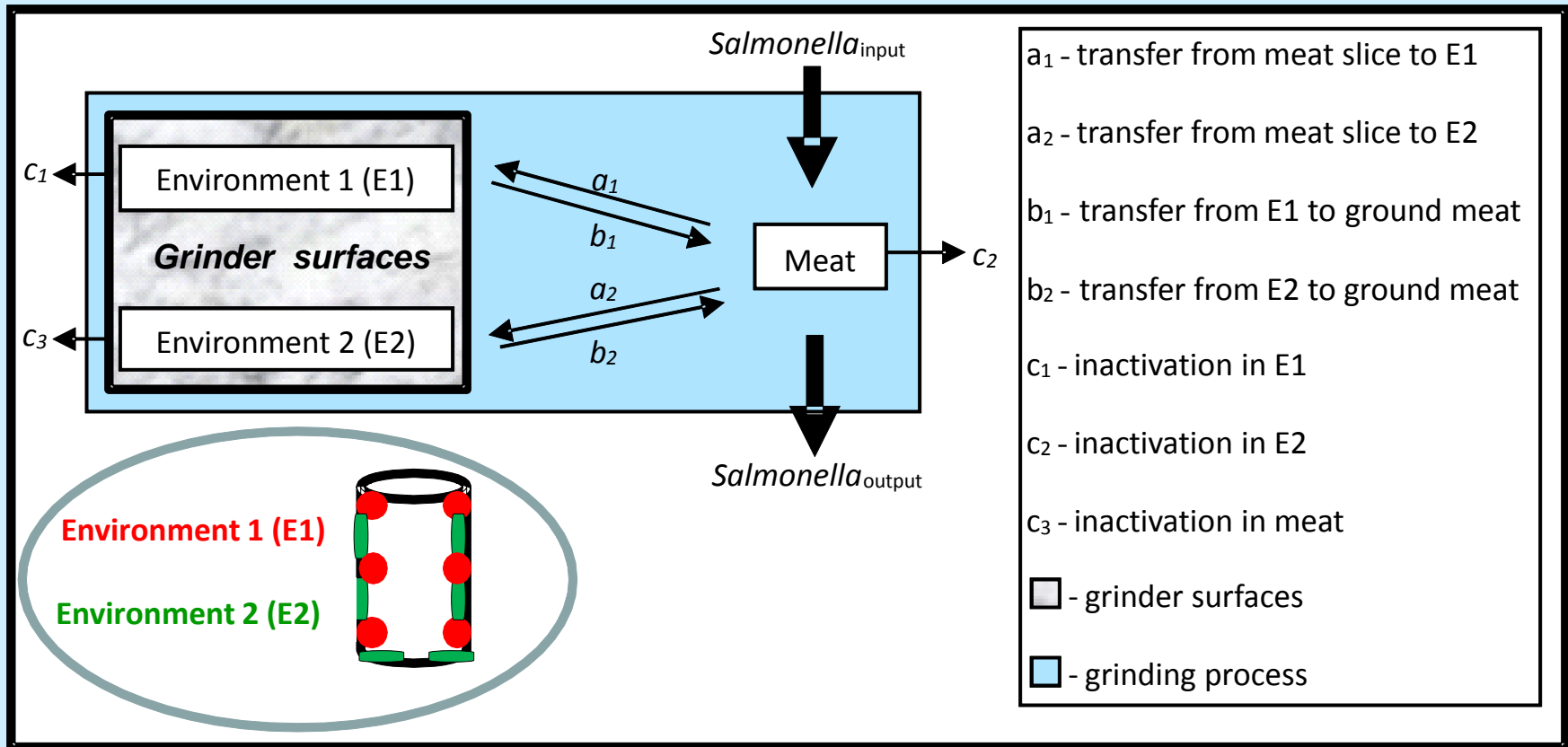
Describing the transfer rates of *Salmonella* during pork grinding



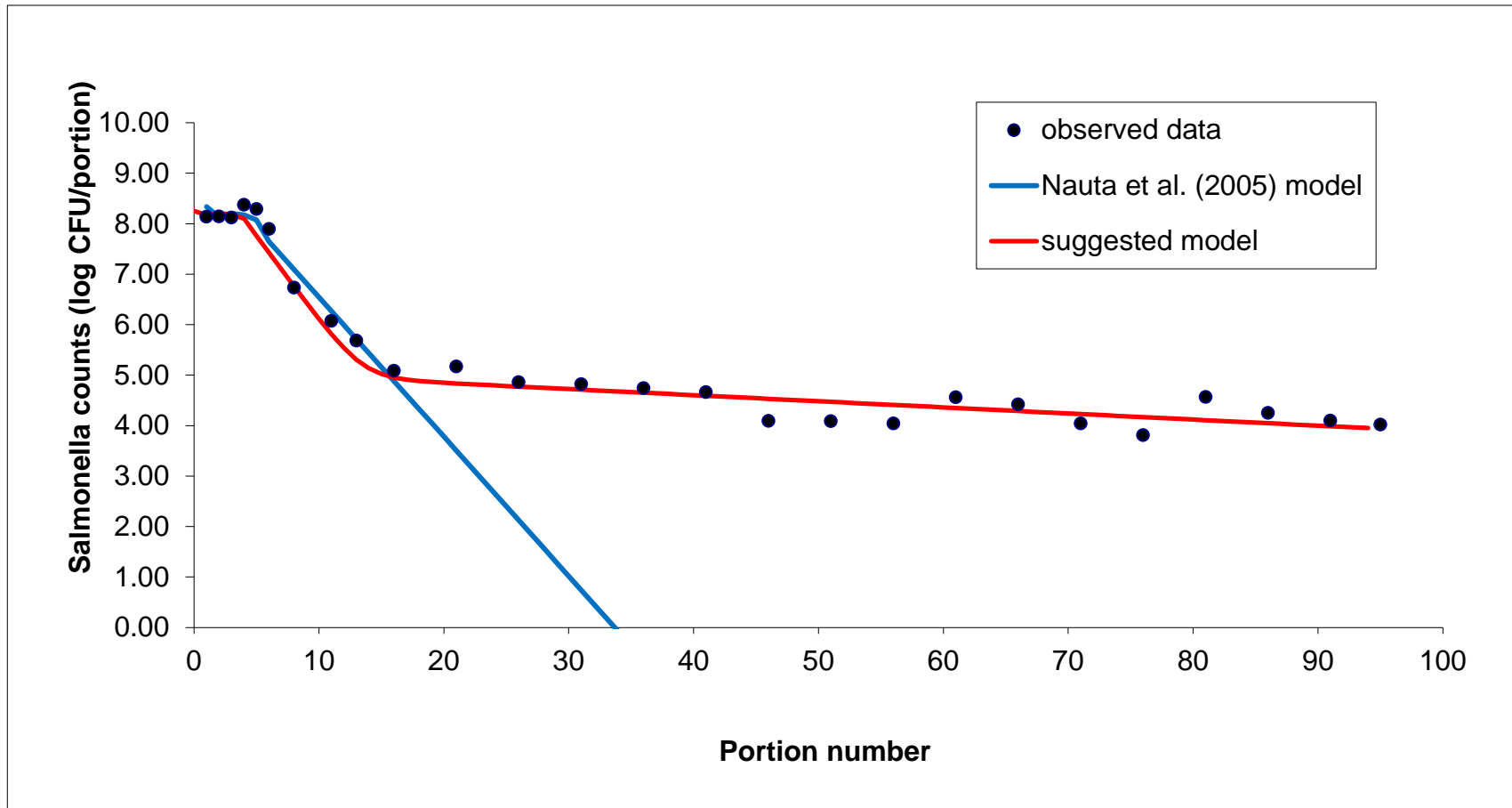
Transfer rates of *Salmonella* DT104 based on cell count data fitted to the suggested model

Modelling cross-contamination during pork grinding

Suggested Model

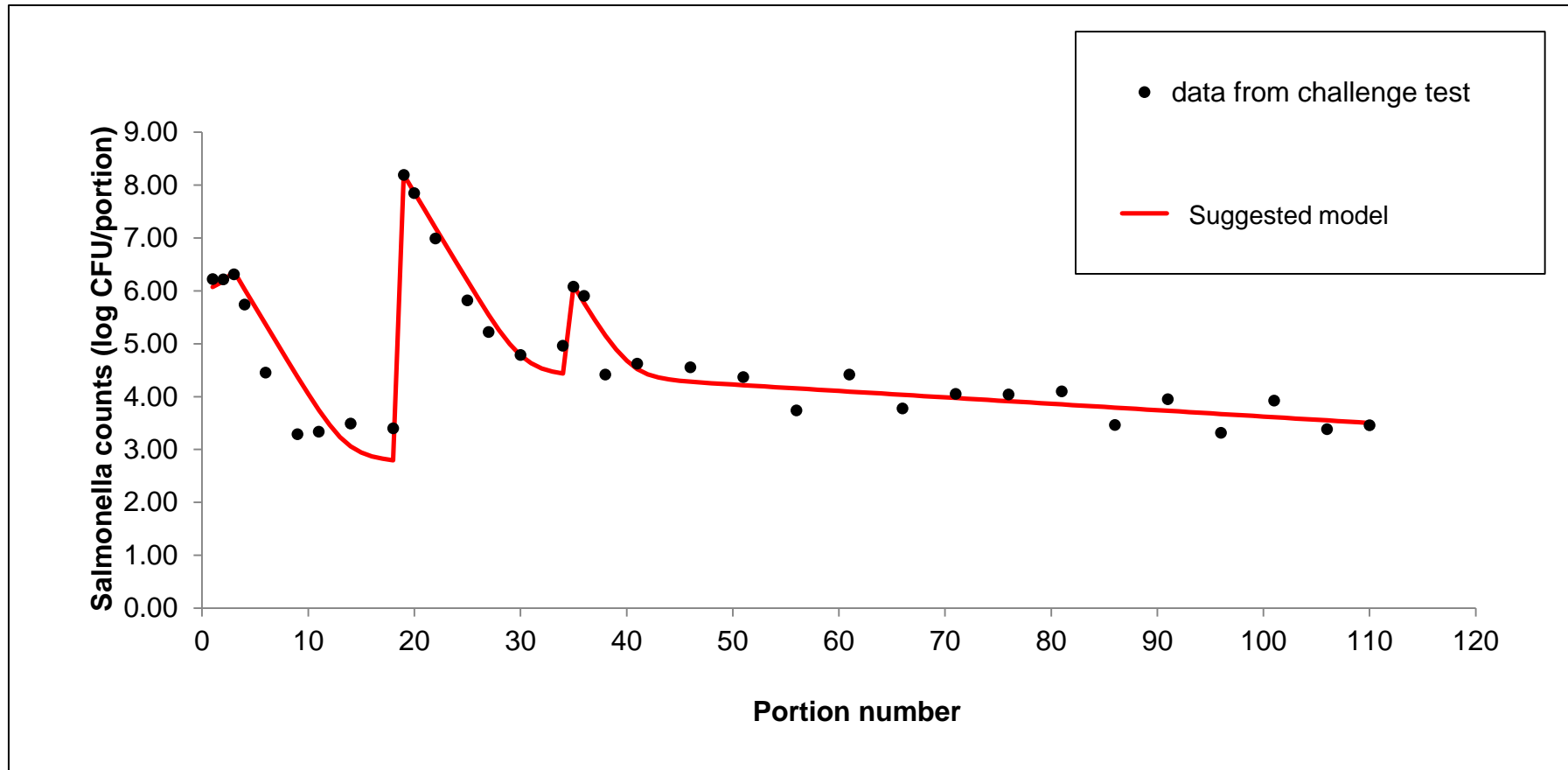


Describing the transfer rates of *Salmonella* during pork grinding



Transfer rates of *Salmonella* DT104 based on cell count data fitted to the suggested model

Challenges in cross-contamination during pork grinding



Describing the transfer rates of *different pathogens* during slicing in other matrices rather than pork using literature data:

- when applying the data published by Vorst *et al.* (2006), simulating cross contamination of *L. monocytogens* during turkey slicing, $R^2 = 0.86$, was found.
- when the data presented by Aarnisalo *et al.* (2007), regarding transfer of *L. monocytogens* during slicing of gravad salmon, was used $R^2 = 0.74$ was obtained.
- and for the data published by Sheen and Hwang (2010) related to cross contamination of *E. coli* O157:H7 during ham slicing, R^2 was 0.78.

ORIGINAL ARTICLE

Modelling transfer of *Salmonella* Typhimurium DT104 during simulation of grinding of pork

C.O.A. Møller¹, M.J. Nauta¹, B.B. Christensen², P. Dalgaard³ and T.B. Hansen¹

- Tail phenomenon
 - ✓ Food processors
 - Control measures
 - Cleaning and sanitization
- Observed transfer successfully modelled
- Model can describe different processes
- Tool to support risk assessors

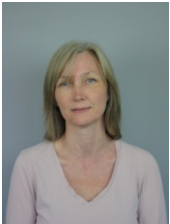
- **To be investigated:**
 - ✓ Food matrices
 - ✓ Pathogens
 - ✓ Inoculum levels
 - ✓ Processings



Acknowledgements

✓ This research project was financed by the Technical University of Denmark through the FoodDTU programme

✓ Constructive advice and critical comments were given by:



Tina Beck Hansen



Maarten Nauta



Paw Dalgaard



Bjarke Bak Christensen



✓ Skillful technical assistance was provided by colleagues from the Division of Food Microbiology at the National Food Institute.



Rikke Krag



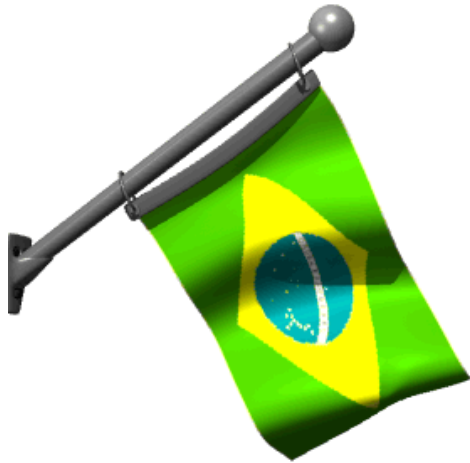
Mette Kemp



Kate Vibefeldt



Louise Vignæs



Meat-Cross-Con

MEAT SAFETY: An innovative modelling approach to evaluate microbial pathogen transfer and cross contamination from farm to fork

Acknowledgements



Thank
You!

Let's start the practical application of the cross contamination model?



Thanks