

Absence of *L. monocytogenes* growth during raw milk cheesemaking: a modelling approach

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Background

Predictive
Microbiology

Domain
Modelling

Modelling
growth

Conclusions



Background - Motivation



- ❑ Lack of data on growth during cheesemaking
- ❑ Most predictions based on lab. media experiments under static conditions
- ❑ Dynamic conditions of food not accounted for
- ❑ Data showing differences between models in liquid and solid media
- ❑ Need for models in food

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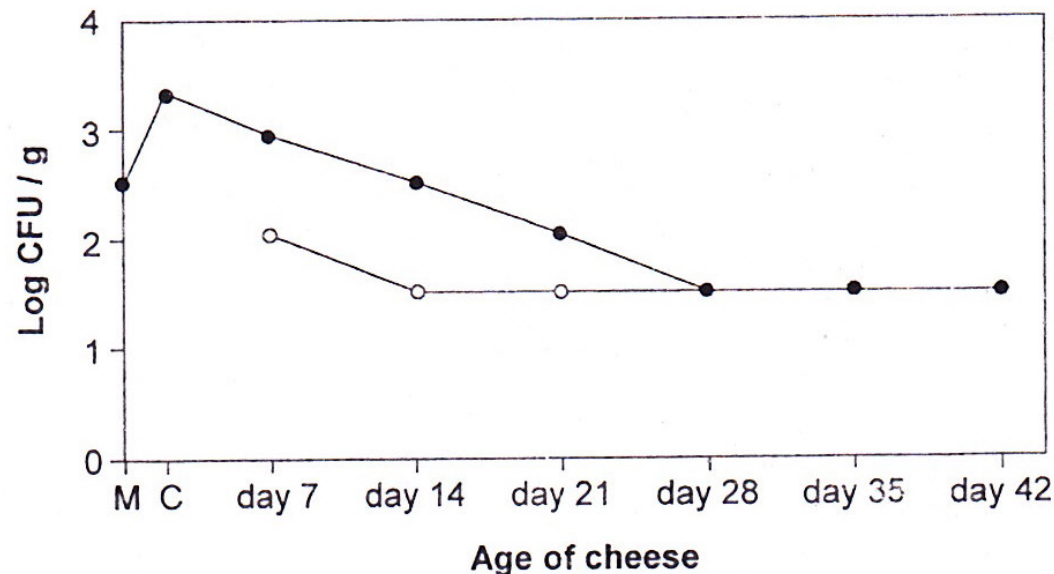
Conclusions



Literature data



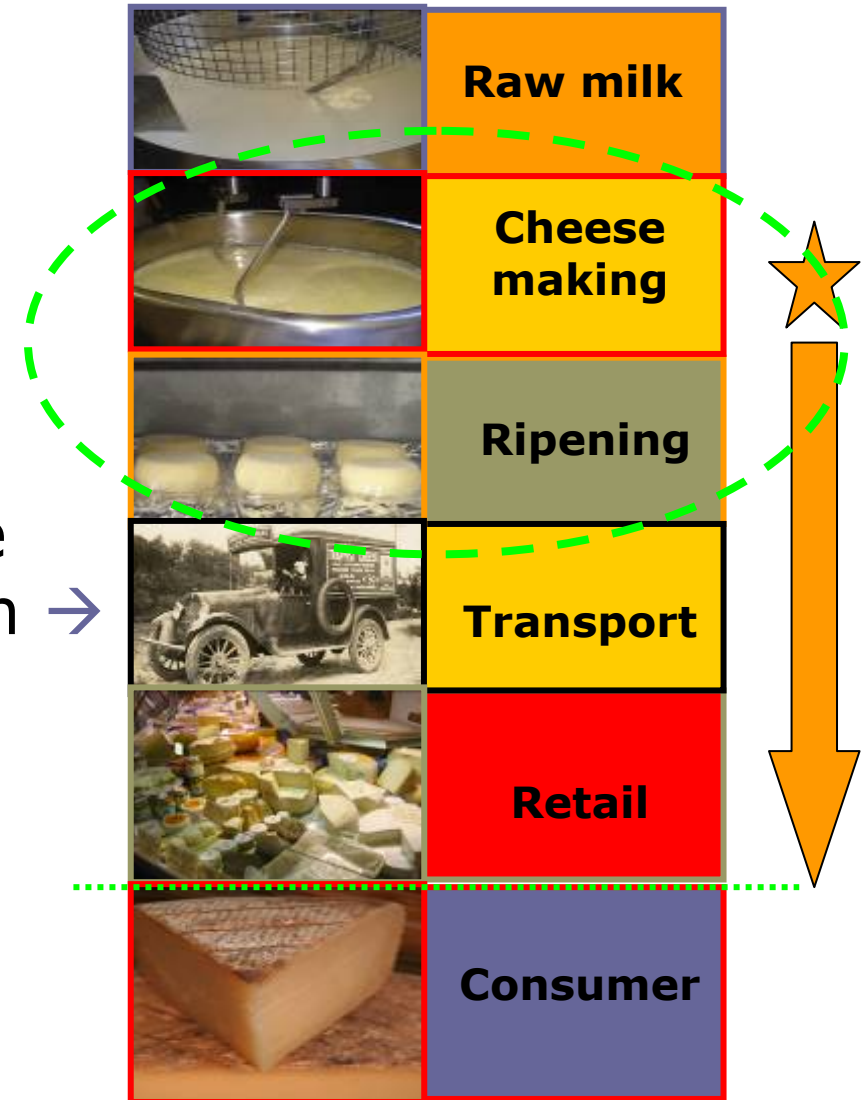
- Inaccurate data from literature – apparent growth
- We used cfu/g dry wt – compare liquid and solid directly



The bigger picture - Domain modelling



- BIG model comprised by several smaller models
- Food chain
- Each step of the production chain → modelled



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Pasteurised vs. Raw milk cheesemaking



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- During Cheesemaking and Ripening
- Laboratory scale cheesemaking
- Pasteurised and raw milk
- Milk spiked with 2 strains *L. monocytogenes* (low numbers)
- 4 Replicates
- Populations monitored through cheesemaking and ripening
- Physicochemical parameters measured

Materials and methods



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<i>Protocol</i>	<i>Sampling</i>
Milk heated to 30°C	Milk
Rennet addition – pH 6.55	Milk
Cutting of the gel	Curd
Cooked to 36°C	Curd
Moulded	Cheese
Brined	Cheese
Smeared	Core/Surface
Ripened	Core/Surface

Cheesemaking

Ripening

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Materials and methods



□ Cheesemaking

- *L. monocytogenes* counts:
 - Plate count enumeration
 - chromogenic agar ALOA
- pH, water activity, temperature, lactose



□ Ripening

- *L. monocytogenes* counts:
 - Plate count enumeration
 - chromogenic agar ALOA
- pH, water activity, temperature, lactose

SAMPLING OF CORE AND SURFACE



Modelling methodology



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- Qualitative interpretation of the data
 - **Comparison of the growth non growth model predictions with observed data**
 - **Extraction of the kinetics showing growth of *L. monocytogenes* during cheese-making and ripening**

- Quantitative interpretation of the data
 - **where growth occurred, adjustment of primary and secondary cardinal growth models with:**
Y = Numeration in CFU/g dry wt.,
Xi = Values of physicochemical parameters, cardinal values for *Listeria*

 - **Assessment of the optimal growth rate of *L. monocytogenes* for different phases**

- Novelty
 - **Dynamic physico-chemical conditions (SAS Model Procedure)**
 - **Dynamics of the matrix : liquid vs solid (CFU / gram of dry weight)**

Determination of the optimal growth rate



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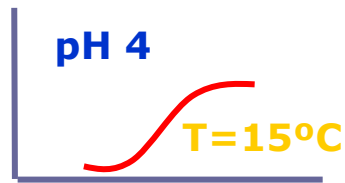
Modelling growth

Conclusions

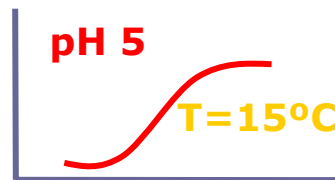


USUAL APPROACH

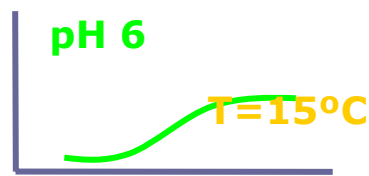
Cfu/g or ml



$\mu(4,15)$



$\mu(5,15)$



$\mu(6,15)$

t

Primary model

$$x(t) = \frac{X_{\max}}{1 + \left(\frac{X_{\max}}{X_0} - 1\right)e^{-\mu \cdot t}}$$

Secondary model

$$\mu_{\max} = \mu_{\text{opt}} CM_2(T) CM_1(\text{pH}) SR_1(\text{aw}) SR_2(\text{Lac}) \zeta(T, \text{pH}, \text{aw}, \text{Lac})$$

μ_{opt}

Determination of the optimal growth rate *in dynamic conditions*



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Modelling growth

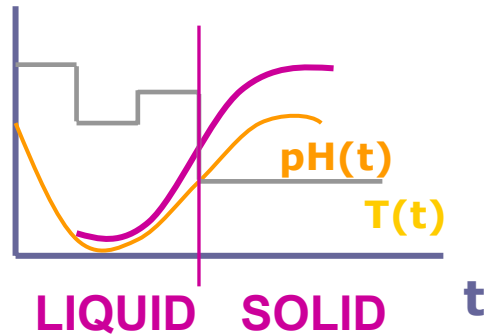
Conclusions



NOVEL APPROACH

cfu/gdw(t)

cfu/ml(t) cfu/g(t)



Primary model Secondary model

$$dx(t) = \frac{X_{\max}}{1 + \left(\frac{X_{\max}}{X_0} - 1\right)e^{-\mu \cdot t}} \mu_{\max} = \mu_{\text{opt}} CM_2(T) CM_1(\text{pH}) SR_1(\text{aw}) SR_2(\text{Lac}) \zeta(T, \text{pH}, \text{aw}, \text{Lac}) dt$$

Physico-chemical parameters

$$\text{pH}(t) = f(t)$$

$$T^\circ(t) = g(t)$$

$$\text{aw}(t) = h(t) \dots$$

μ_{opt}

Observed data

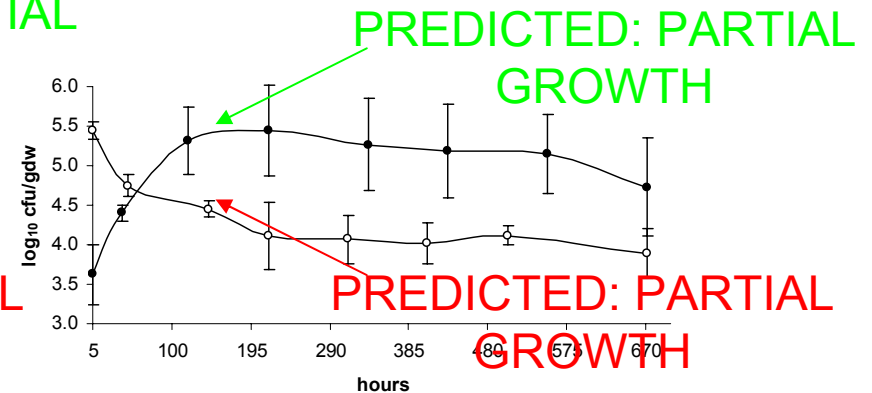
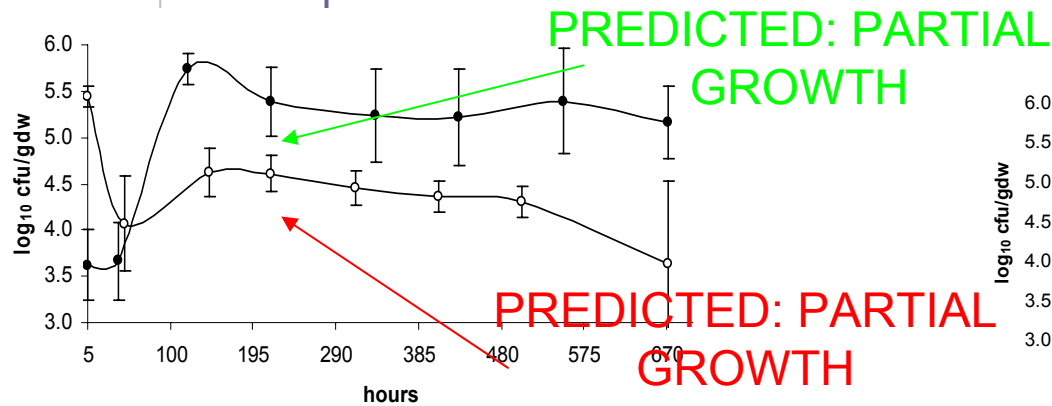
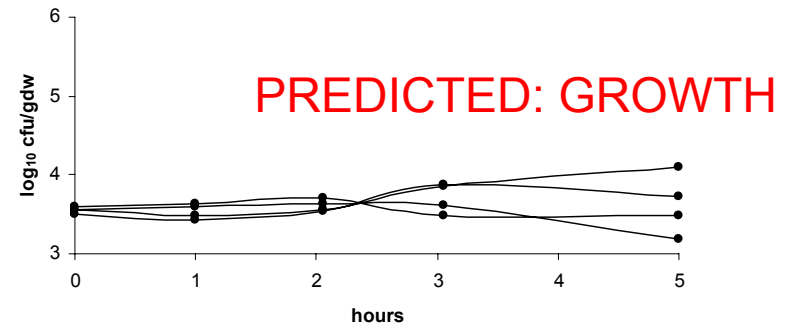
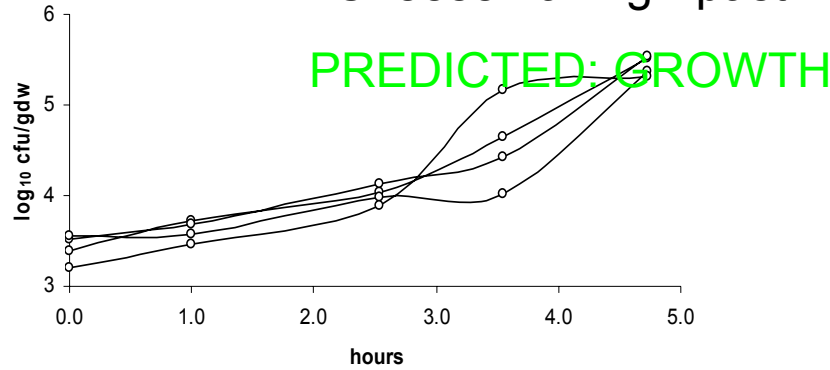
vs

growth/no growth predictions



Cheesemaking - past

Cheesemaking - raw



Ripening / Core

- = raw
- = pasteurised



Ripening / Rind

- = raw
- = pasteurised

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Predictive Microbiology

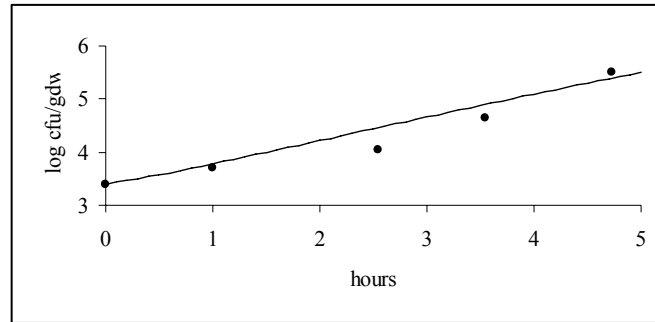
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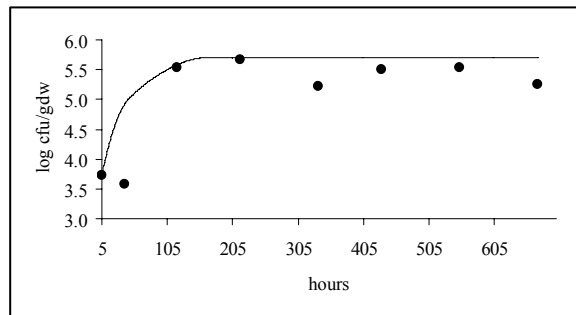


Optimal growth rates assessed



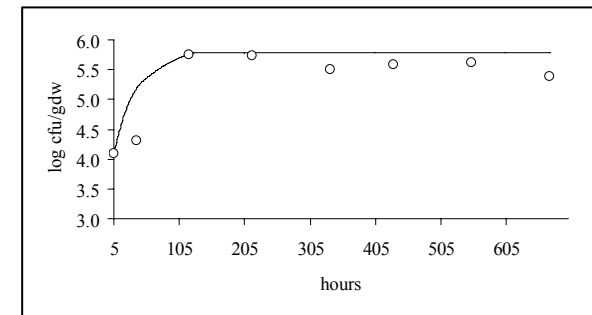
Cheese making / pasteurized milk

$\mu_{opt} \sim \text{Normal}(1.2; 0.01)$



Ripening / raw / core

$\mu_{opt} \sim \text{Normal}(0.18; 0.002)$



Ripening / raw / surface

$\mu_{opt} \sim \text{Normal}(0.16; 0.001)$

Summary



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□ Cheesemaking

- Growth in pasteurised milk – no growth in raw milk
 - Raw milk – natural flora inhibited starter and listeria – low lactic acid production

□ Ripening

- Growth in raw milk cheese – inactivation in pasteurised milk cheese
 - Raw milk – low lactic acid and increasing pH allowed growth

□ Models in food are a useful tool in growth situations

Future Perspectives



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- Predictive Modelling useful to understand, predict and describe behaviour in food
- Combase online tool → static environmental conditions
- More data in food systems is needed / Validation in foods
- Inactivation models in dynamic conditions need to be developed

Acknowledgments



- **The work was supported by the European Union funded Integrated Project BIOTRACER (contract 036272) under the 6th RTD Framework**



- **CNIEL (National Interprofessional Centre for the Economy of Dairy Products) Department of Scientific Research – Paris, France**

- **Alexandre Maffre, Fanny Tenenhaus-Aziza, Moez Sanaa**



- **TEAGASC Walsh Fellowship**



- **UCD, Prof. Francis Butler**



Thank You for your
attention!

