

A dynamic *in silico* model of rumen microbial fermentation and methane production

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The rumen microbiota: key player of the animal

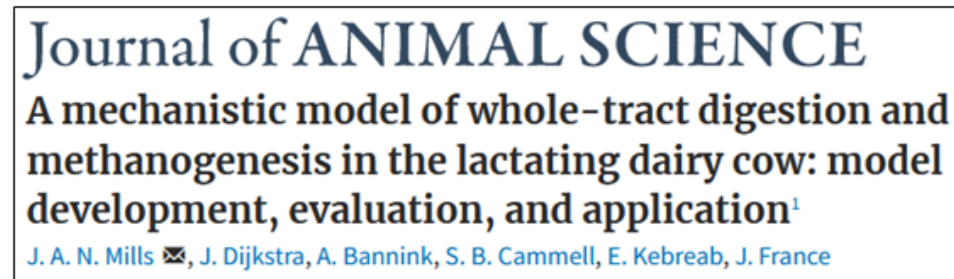
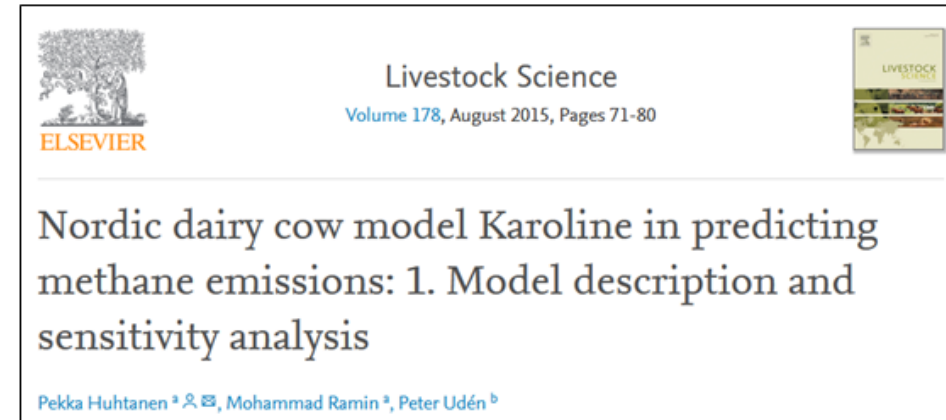
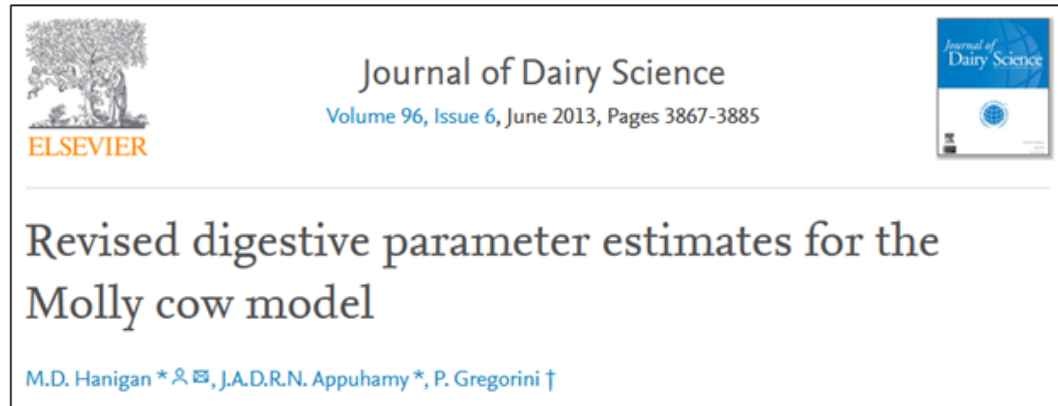


- Feed efficiency
- Ruminant production
- Health and well-being
- Environmental footprint

Our goal: to develop mathematical models

- Enhance understanding of rumen function
- Provide virtual platform to simulate the dynamics of rumen fermentation and methane production
 - Potential use in the design of nutritional strategies to improve animal health, production, and reduce methane production

Dynamic rumen models: longstanding research area



Rumen modelling: room for improvement

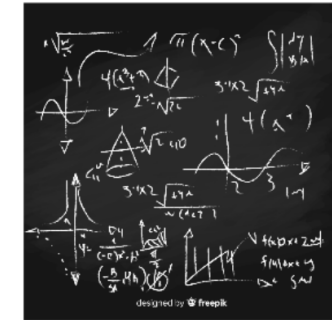
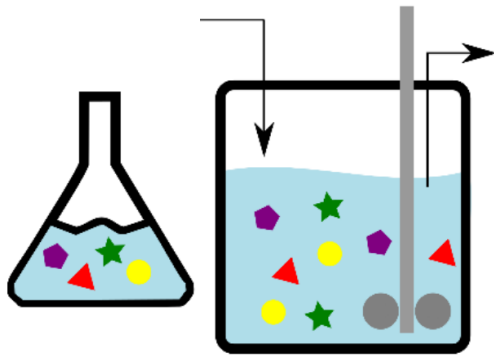
Aspects for improvement ([Ellis et al., 2008, J. Agric. Sci.](#); [Janssen 2010, AFST](#))

- Better representation of the microbiota
- Incorporation of hydrogen dynamics
- Thermodynamic control

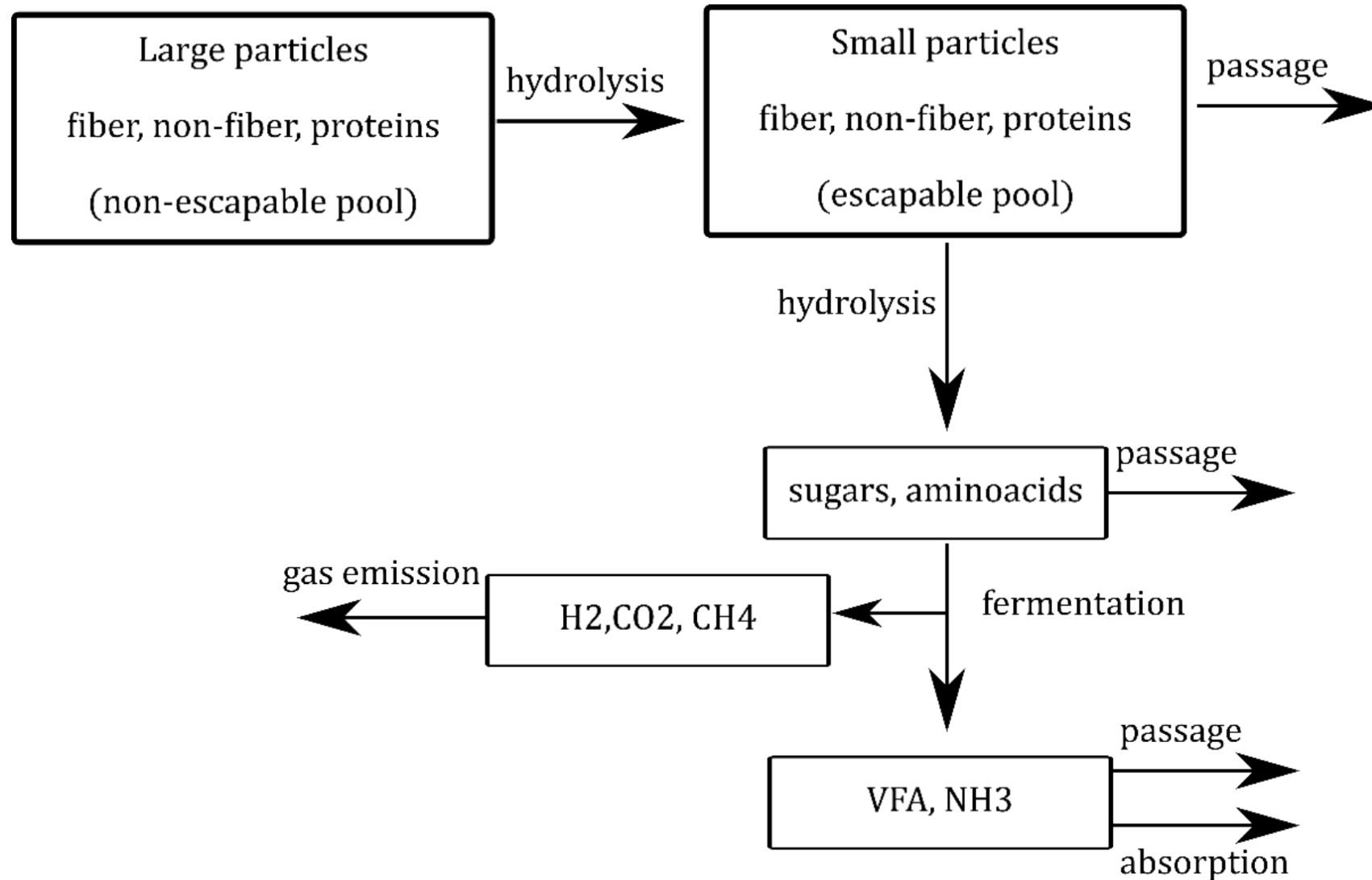
Dynamic component has been seldom considered in model evaluations and in model exploitation

- Assumption of steady-state
- An exception ([van Lingen et al., 2019, JTB](#))

Our modelling strategy: integrating *in vitro*, *in vivo*, *in silico*

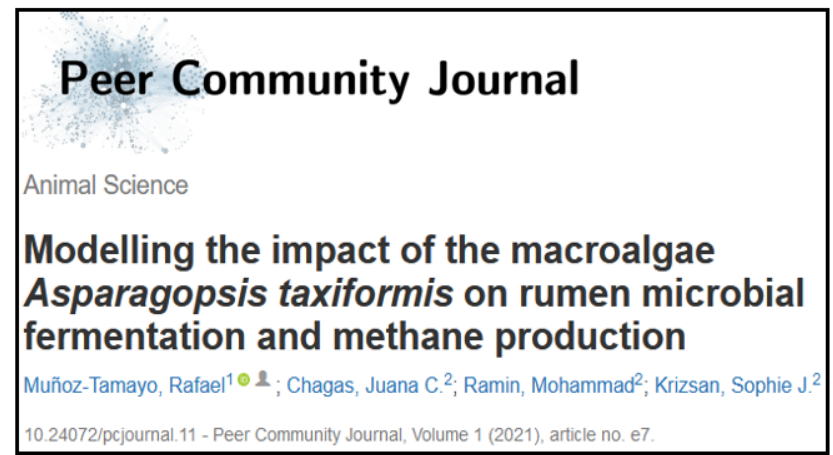
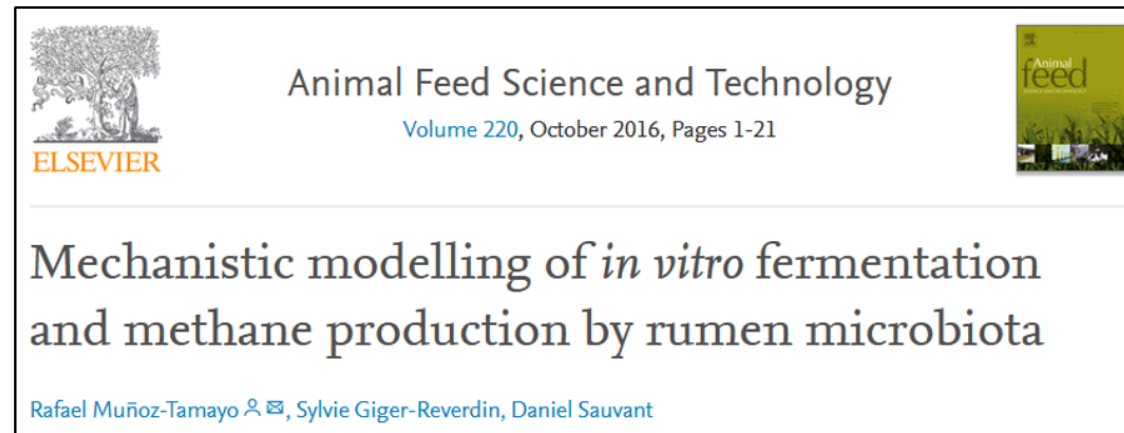
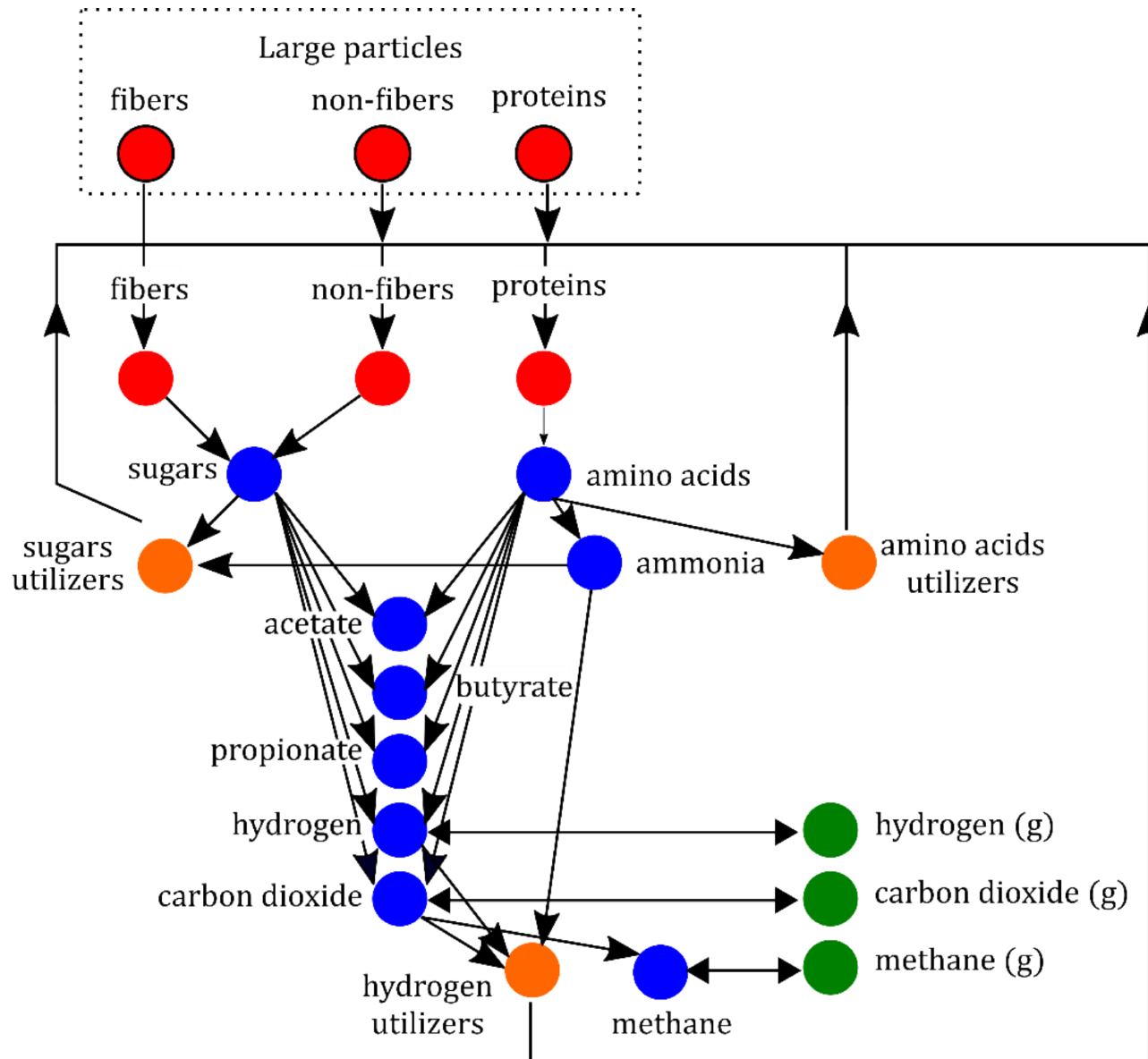


Our model in figures: mass fluxes





Our model in figures: fermentation pathway



Our model: some equations

- Mass balance equations: ordinary differential equations

$$\frac{dx}{dt} = \mathbf{S} \cdot \boldsymbol{\rho}(\mathbf{x}, \mathbf{p}) - \mathbf{g}(\mathbf{x}, \mathbf{p})$$

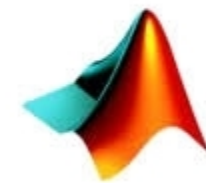
$\mathbf{x}$: vector of metabolites concentrations

$\mathbf{S}$: stoichiometry matrix

$\boldsymbol{\rho}(\mathbf{x}, \mathbf{p})$: vector of reaction rates

$\mathbf{g}(\mathbf{x}, \mathbf{p})$: function vector of transport phenomena

- 21 state variables (compartments)
- 37 parameters
 - Physicochemical and fermentation parameters (yield factors, Monod parameters of the kinetic rates): [Muñoz-Tamayo et al., \(2016\), AFST](#)
 - absorption rate constants: initial guess from [Dieho et al., \(2016\), JDS](#)
 - microbial biomass data: [Ahvenjärvi et al. \(2002\), JAS](#); (2018) JDS
 - passage rates: [Bayat et al., \(2011\), AFST](#)



MATLAB®

Implementation

Model calibration (parameter estimation)

- Experimental data from 4 Nordic-Red dairy cows rumen fistula (2 days)
 - Dry matter intake: 3 h
 - Rumen sampling for VFA (acetate, butyrate, propionate) determination: 3 h
 - Methane in respiration chambers: 17 min
- Maximum likelihood estimation
 - 5 parameters:
 - absorption rate constants of VFAs
 - Hydrolysis rate constants of NSC and protein
 - Check for identifiability



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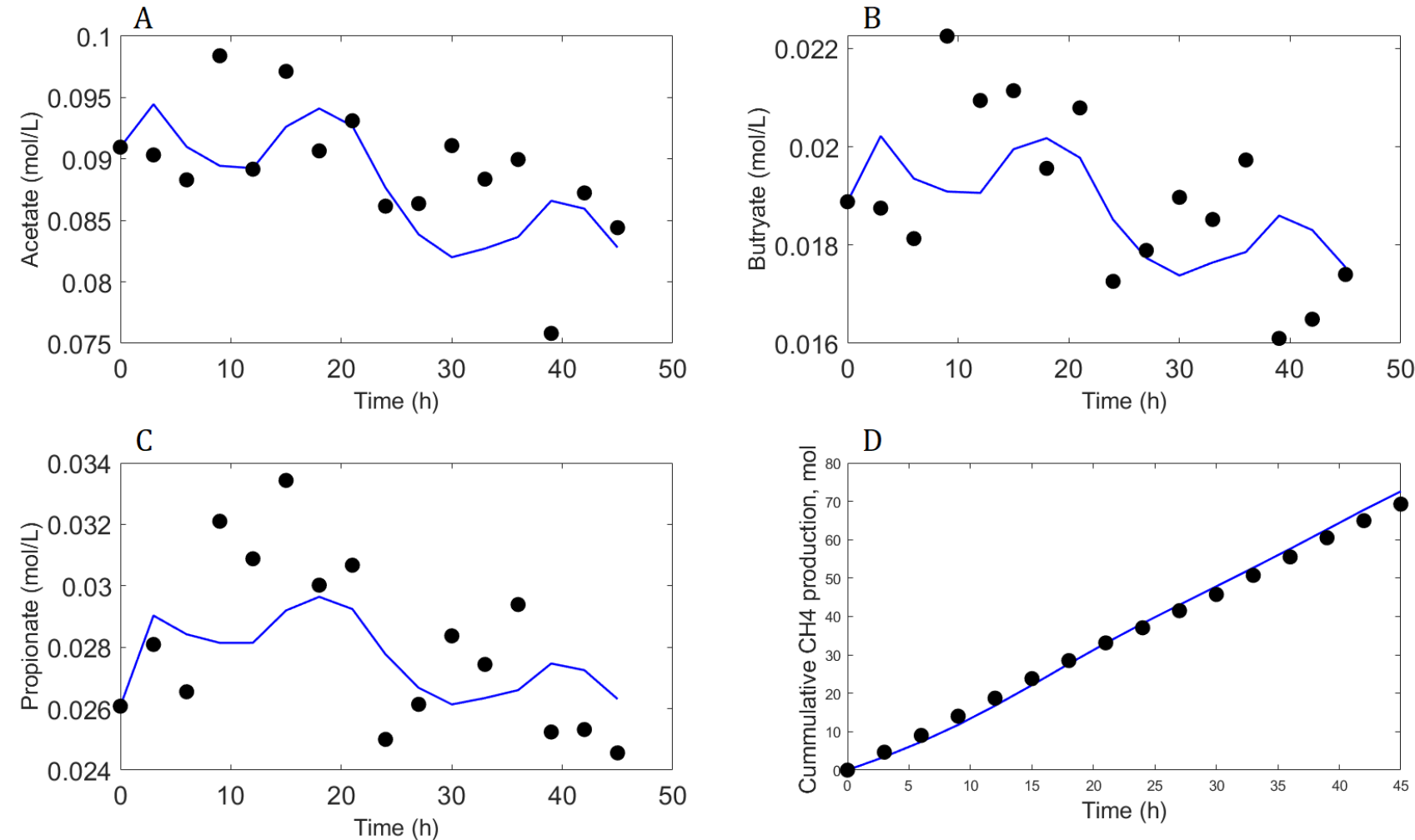
IDEAS: a Parameter Identification Toolbox with Symbolic Analysis of Uncertainty and its Application to Biological Modelling

R. Muñoz-Tamayo ^{*}, B. Laroche ^{*}, M. Leclerc ^{*}, E. Walter ^{*}

<https://doi.org/10.3182/20090706-3-FR-2004.00211>

Example results: one cow

Experimental data (●) , — predicted variables



Capability of the model to represent the dynamic pattern of fermentation

	CVRMSE
Acetate	9%
Butyrate	11%
Propionate	13%
CH ₄	21%
Cummulative CH ₄	5%

Take home message

- Our mathematical model represents satisfactorily the rumen fermentation and CH₄ production by dairy cows
- Ongoing work
 - Include mechanistic description of pH ([Muñoz-Tamayo et al., 2016, AFST](#))
 - Include hydrogen thermodynamic control ([Muñoz-Tamayo et al., 2021, PCJ](#))
 - Make the code Open Source and available for the scientific community
 - User interface: R or Python

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- Cows were offered total mixed ration at 7:00, 13:00 and 19:00 h consisting of grass silage (timothy-meadow fescue sward) provided at 45:55 forage to concentrate ratio on a dry matter basis.

TMR = 58 kg/dDMI = 26 kg/d

- NDF = 39%, NSC = 36% , PRO = 15%
- Acclimatisation period in chambers: 1 day