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# The structure of the food matrix at different length scales affects the mechanisms of digestion and the nutrient bioavailability

Didier Dupont

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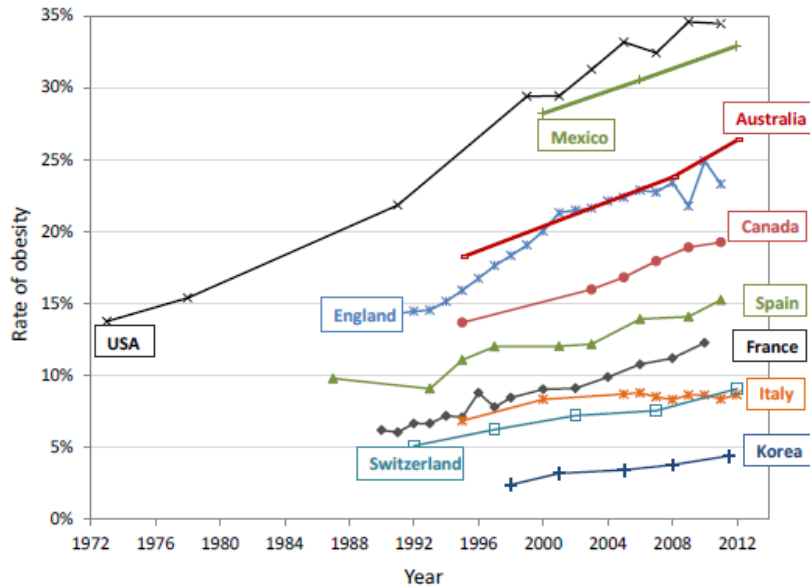
# The structure of the food matrix at different length scales affects the mechanisms of digestion and the nutrient bioavailability



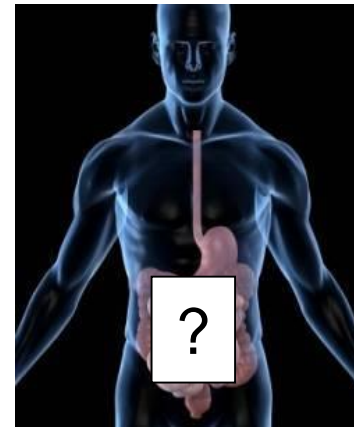
Dr Didier DUPONT, INRAE, STLO, Rennes, France



# Food and human health: the key role of digestion



Diet-related diseases ↑  
Prevent these pathologies rather than  
cure them



**Gut = interface between food and human body**

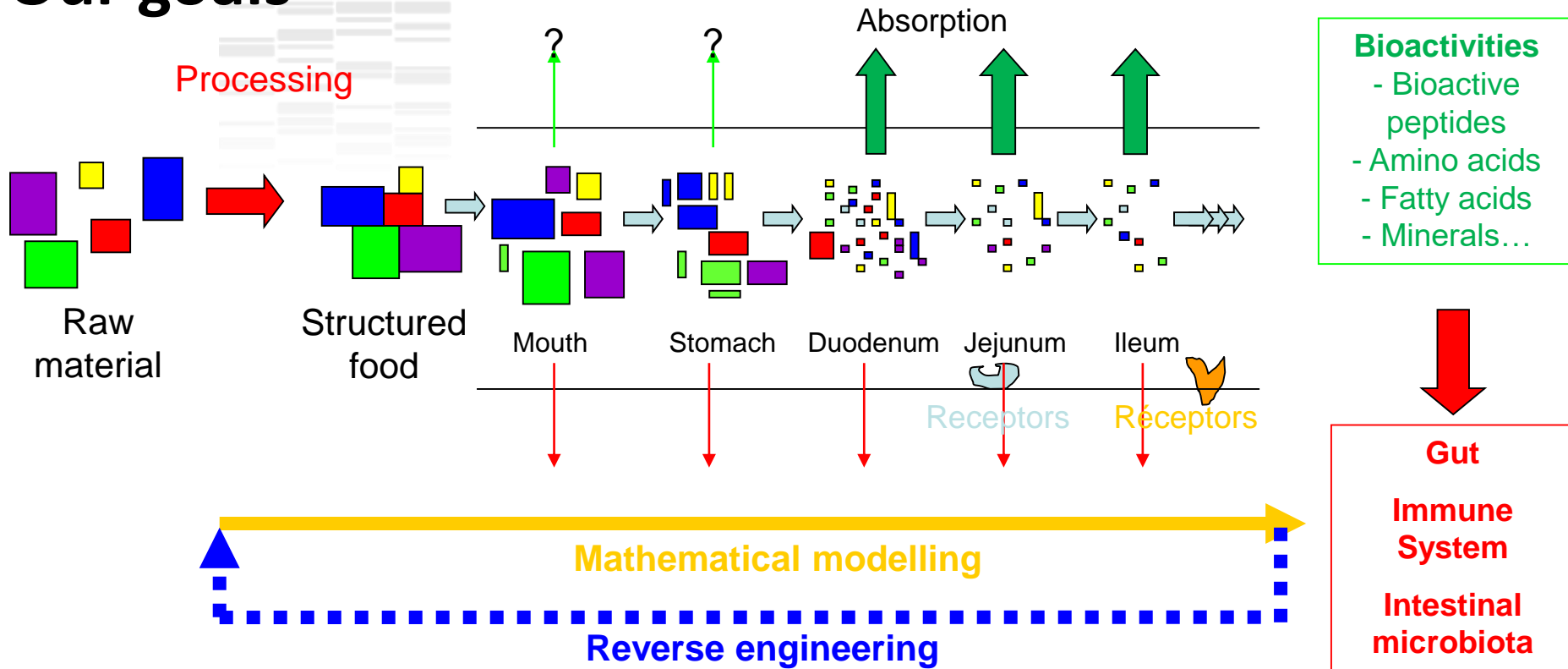
**Digestion releases food components that can have a beneficial or a deleterious effect on human health**

... but the mechanisms of food disintegration in the gastrointestinal tract remain unclear and the digestive process has been considered as a black box so far

**By increasing our knowledge on food digestion, we will increase our knowledge on the effect of food on human health**

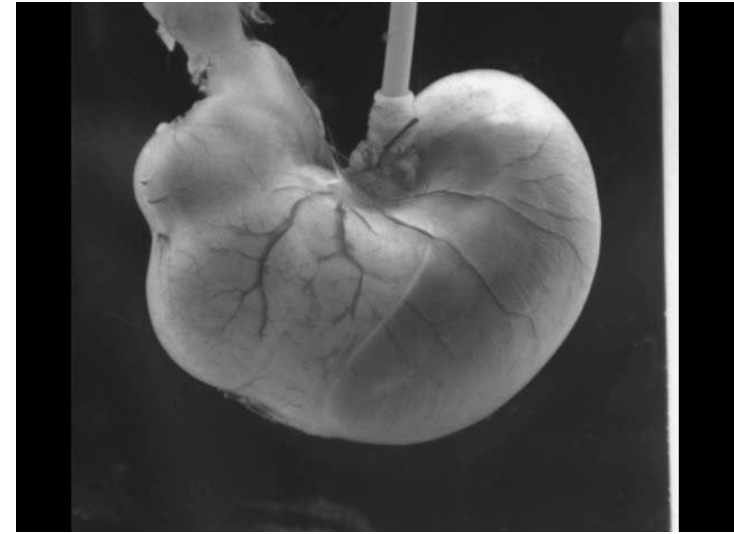
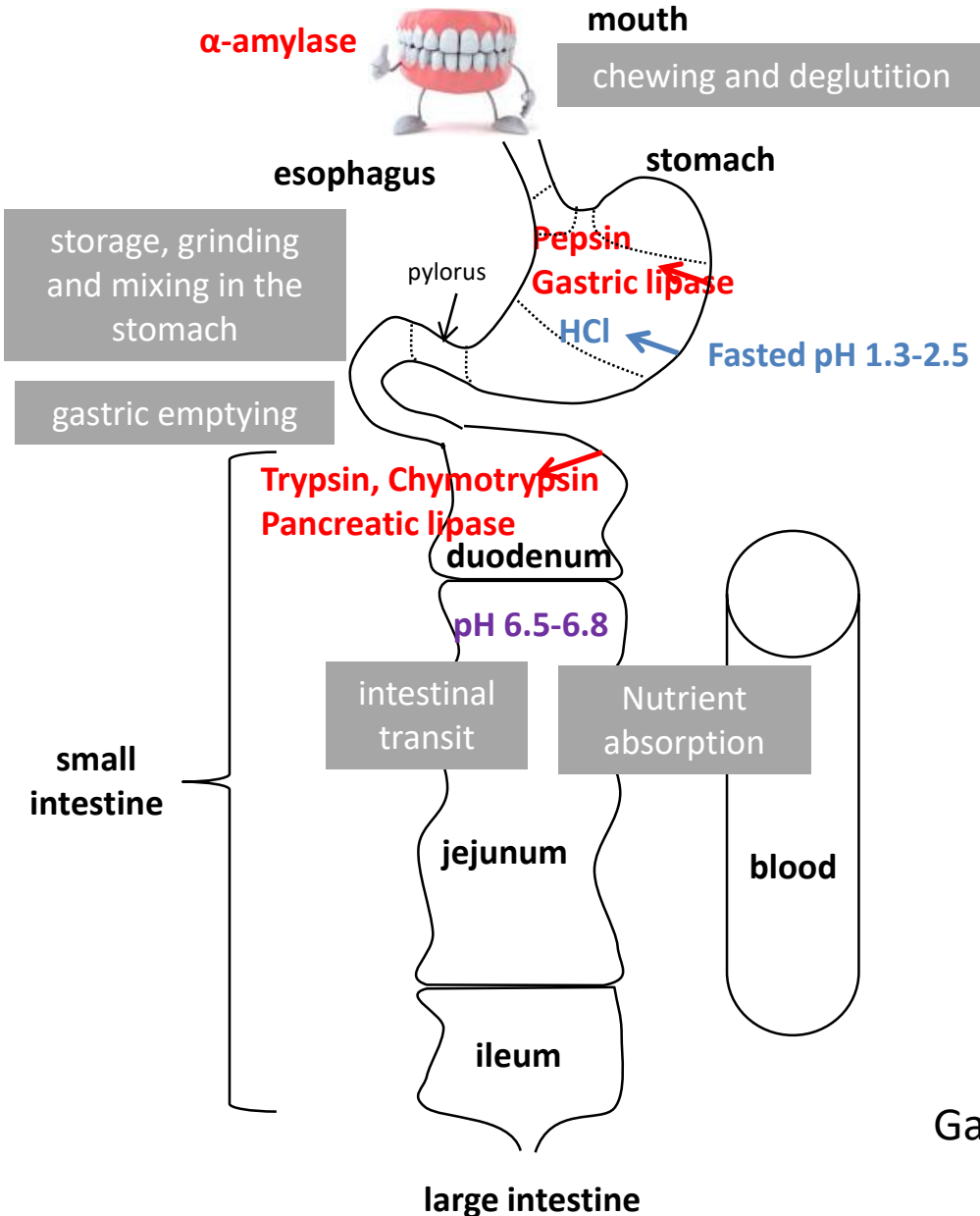
# Our goals

Healthy Adult/ Infant / Elderly

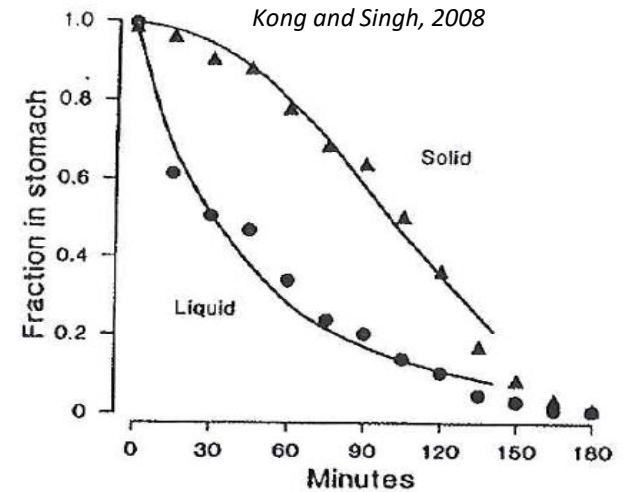


- ☞ To understand the mechanisms of breakdown of food matrices and their constituents in the gut and identify the beneficial/deleterious food components released during digestion
- ☞ To determine the impact of the structure of food matrices on nutrient bioavailability
- ☞ To model these phenomena in order to develop a reverse engineering approach

# The digestive process



From Roger Lentle, Massey Univ. NZ



Gastric phase = a very complex but crucial step for the whole digestion process

# Models available at INRAE for simulating digestion

Peng et al. 2021  
 Halabi et al. 2021  
 Giribaldi et al. 2021  
 Chauvet et al. 2023  
 Nebbia et al. 2022, 2023

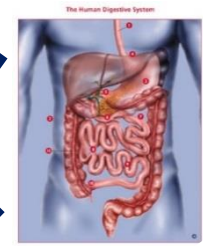
Menard et al. 2018, 2023  
 Wang et al. 2022



*In vitro static models  
 (infant, adult, elderly)*

Le Feunteun et al.  
 2014, 2020

*In silico  
 models*



*In vitro dynamic models  
 (infant, adult, elderly)*

$$\Phi_{12} = k_{12whey} \times (V_1 - m_{caswpd1} \times \alpha) + k_{12aggr} \times m_{caswpd1} \times \alpha$$



*Human  
 models*



*Animal models*



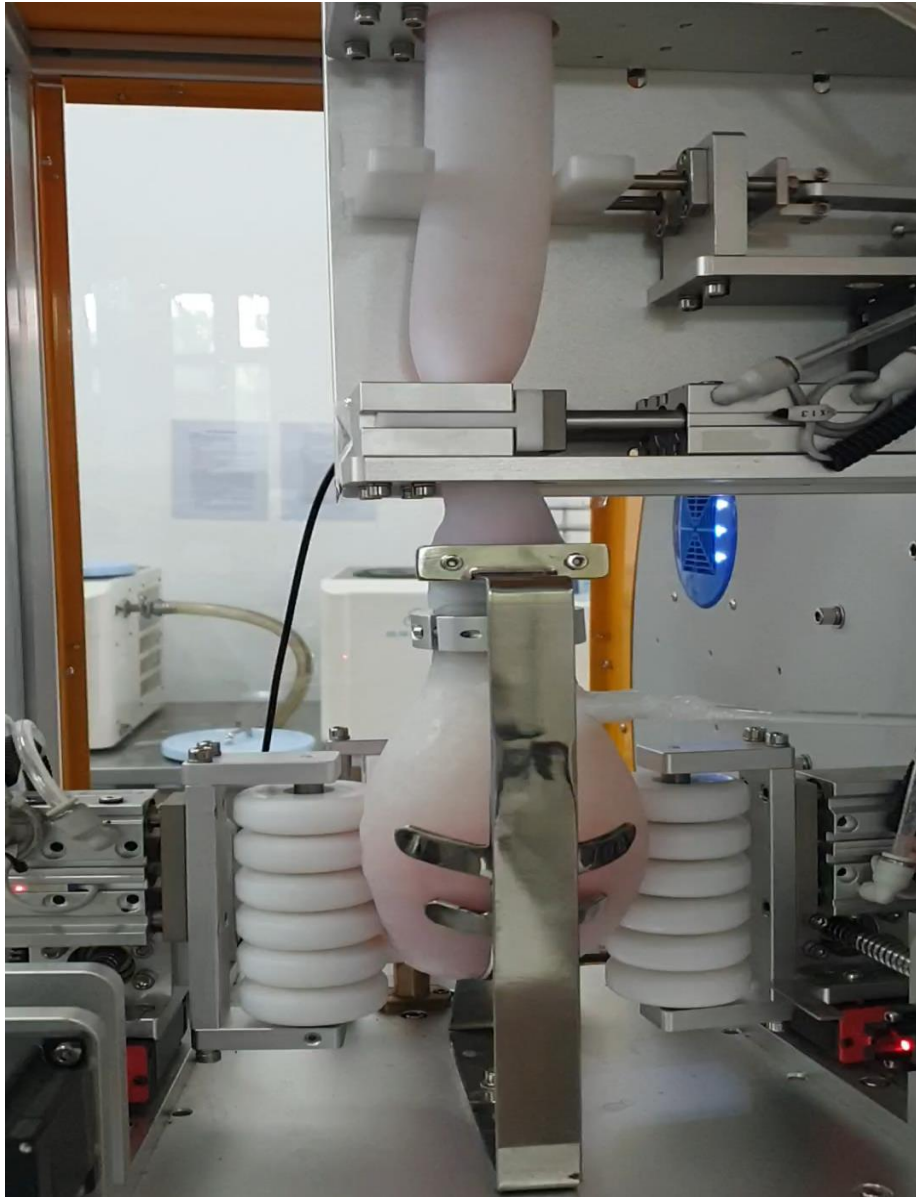
De Oliveira et al. 2016  
 De Oliveira et al. 2017  
 Buffière et al. 2020  
 Boulier et al. 2023



Lemaire et al. 2021  
 Nau et al. 2022  
 Jimenez-Barrios et al. 2023  
 Charton et al. 2022, 2023



# NERDT™ : the NEar Real Digestive Tract



Xiao Dong Pro-Health  
Smart Digestion  
Suzhou University

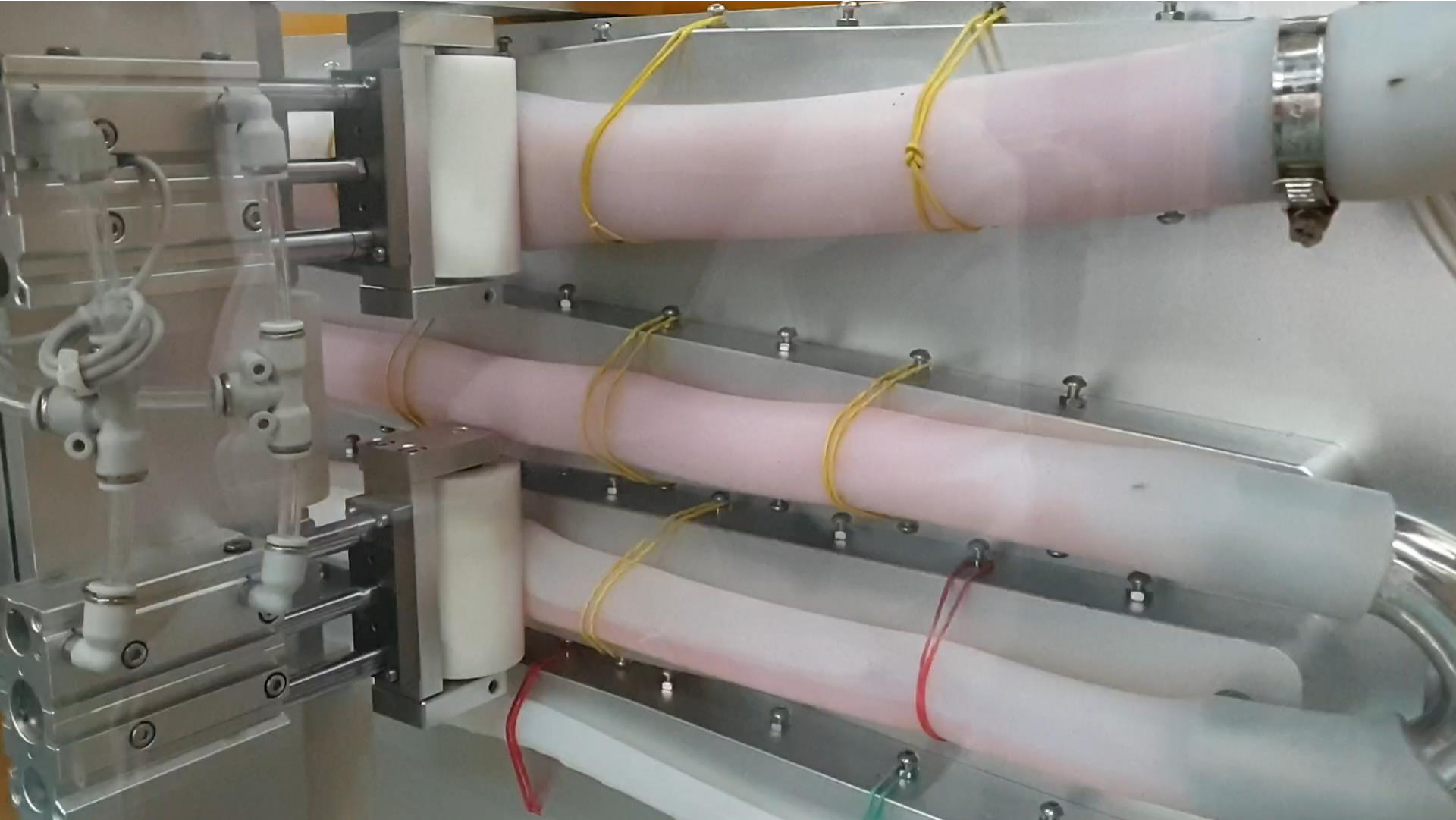


INRAE

L'INSTITUT  
agro Rennes  
Angers



# Simulating the small intestine





# The molecular structure of food protein affects the kinetics of digestion

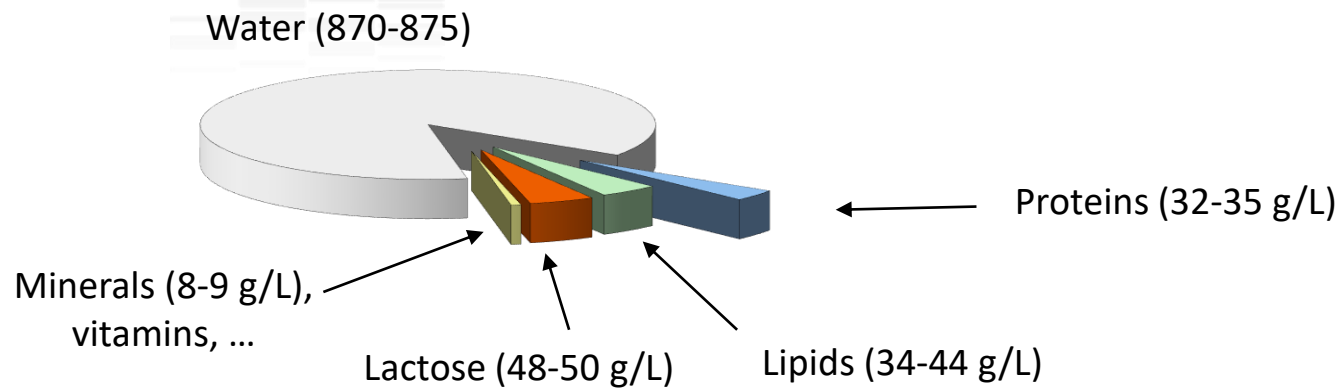
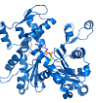


Boudry G., Henry G & Dupont D.  
INRAE, Rennes, France



# Milk

## Molecular scale

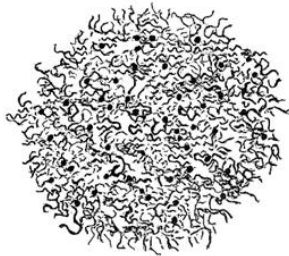


Caseins (80%)  
 $\alpha_{S1}$ ,  $\alpha_{S2}$ ,  $\beta$ ,  $\kappa$

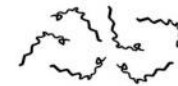


Whey Proteins (20%)  
 $\beta$ -lg,  $\alpha$ -la

$\varnothing \sim 200$  nm  
(Holt, 1994)



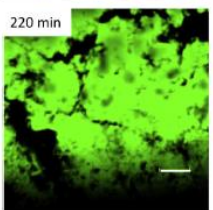
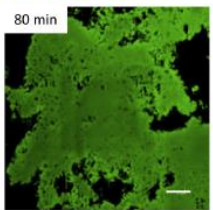
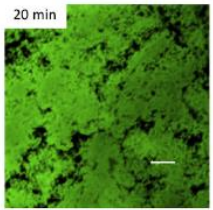
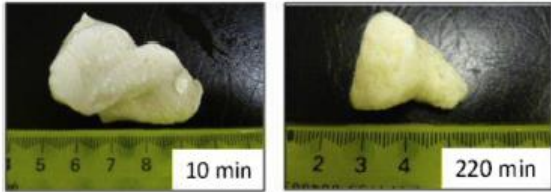
Casein organized into a supramolecular structure: the **casein micelle (CM)**



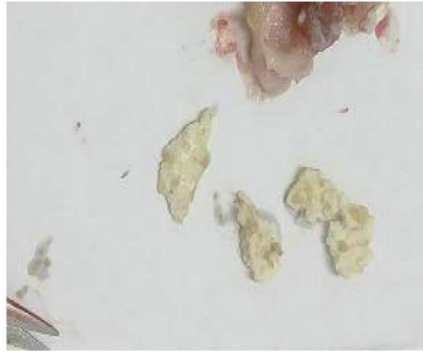
$\varnothing \sim 11$  nm  
~15 casein molecules;  
(Thomar *et al.* 2013)

Casein can also be extracted after acidification followed by neutralization: the **caseinate (CS)**

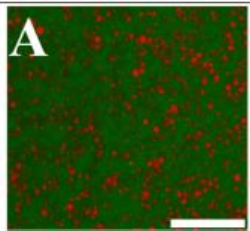
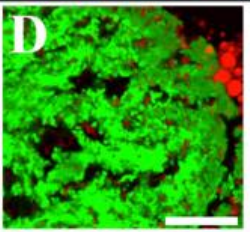

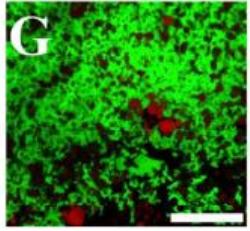

# Milk coagulates in the stomach



*In vitro* demonstration using the HGS  
Ye *et al.* 2016



Ye *et al.* 2019  
*In vivo* evidence using a rat model

	Raw	
Initial		
36min		
182 min		

Mulet-Cabero *et al.* 2019  
*In vitro* semi-dynamic model



Skim Milk Powder



Sodium Caseinate

Casein micelles and sodium caseinate form different coagulums  
Wang *et al.* 2018

The objectives of the study were to:

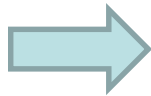
- 1 Determine whether gastric emptying of an isoproteic solution of CM and CS are different or not (exp. 1)
- 2 Characterize the structure of the resulting chyme and determine if CM and CS are differently metabolized (exp. 2)

## Experiment 1 – Determination of Gastric Emptying

96 g of CM or CS  
rehydrated in  
800 ml of water

+12 g of glucose

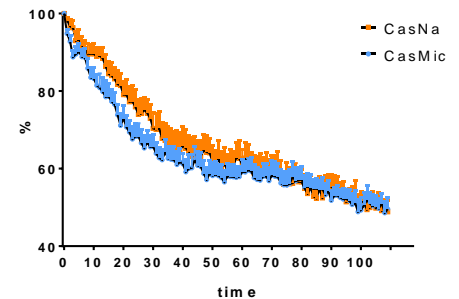
+  $^{99m}\text{Tc}$ -colloidal  
(25Mbq)



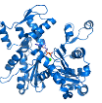
9 pigs (20-25 kg)



$\gamma$ -scintigraphy over 120 min



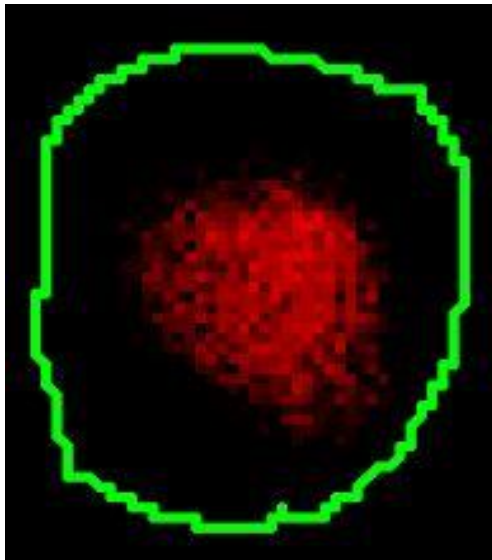
Gastric emptying half-time ( $T_{1/2}$ ) and shape of the curve ( $\beta$ )



## But a differential behaviour of CS and CM in the stomach

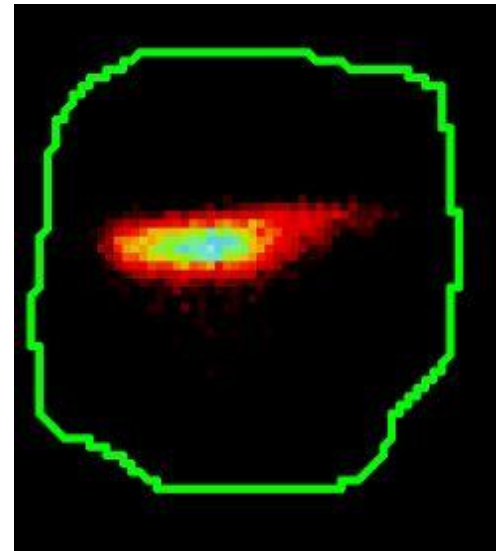
Exemple scintigraphic images at the beginning of gastric emptying (5-10 min after ingestion)

CM



Radioactivity fully fills the stomach

CS

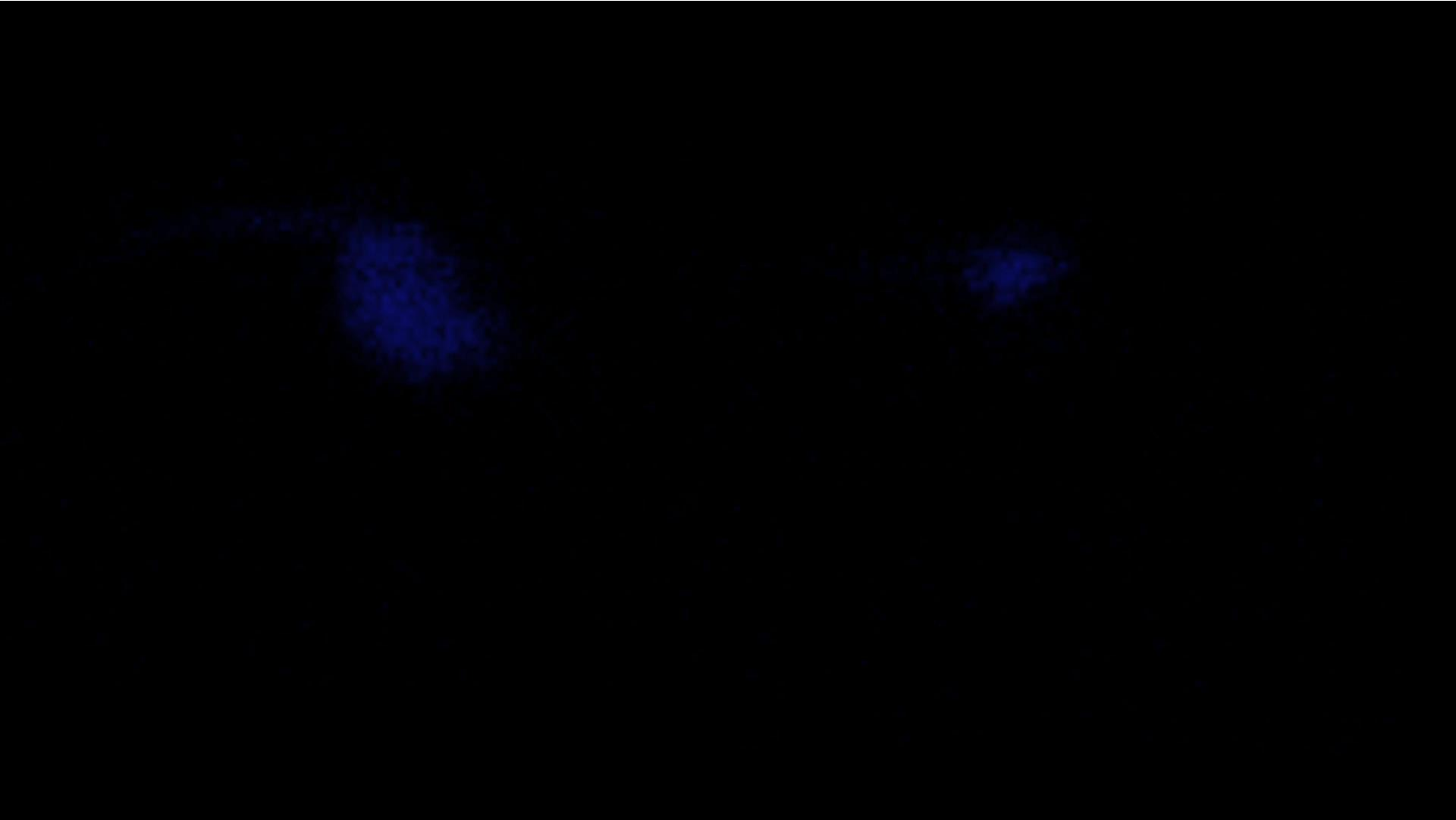
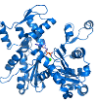


Radioactivity is highly concentrated in the proximal part of the stomach



# Gastric emptying

Molecular scale





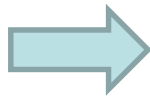
The objectives of the study were to:

- 1 Determine whether gastric emptying of an isoproteic solution of CM and CS are different or not (exp. 1)
- 2 Characterize the structure of the resulting chyme and determine if CM and CS are differently metabolized (exp. 2)

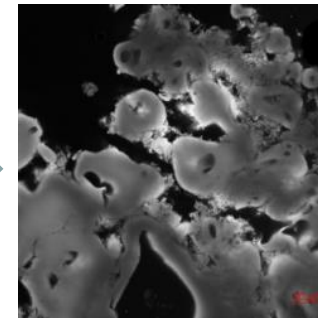
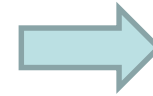
## Experiment 2 – Chyme structure and protein metabolism

96 g of CM or CS rehydrated in 800 ml of water

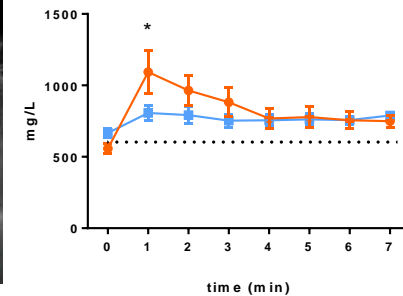
+12 g of glucose



10 catheterized pigs (20-25 kg)



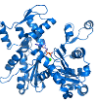
Characterization of the chyme structure (slaughtering after 10 min, n=4)



Free plasma amino acids over 7h n=6

# Collection of the stomach contents

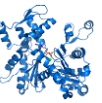
Molecular scale



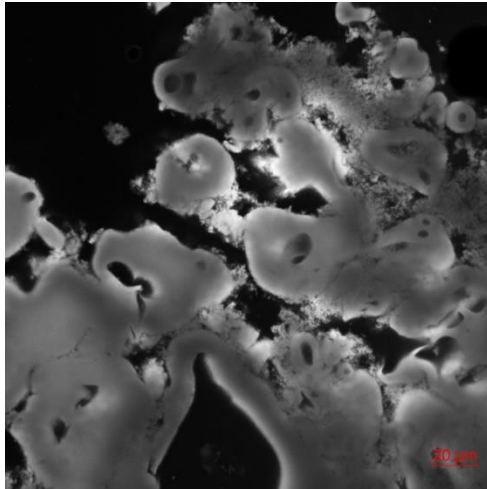
CM forms a large coagulum in the stomach whereas CS mainly remains in the liquid form

# Microstructure of gastric chymes

Molecular scale



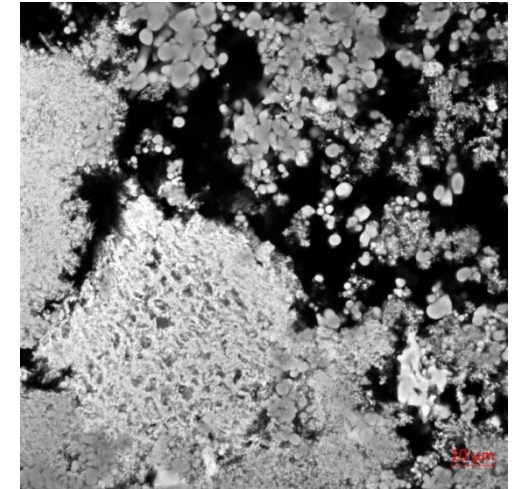
CM



CM Gels (left) are compact and dense =

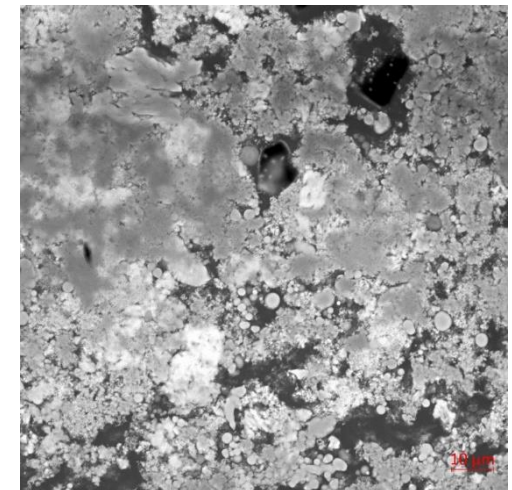
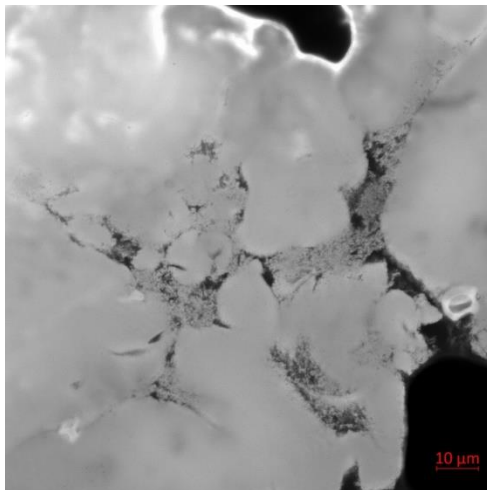
**Strong coagulum**

CS



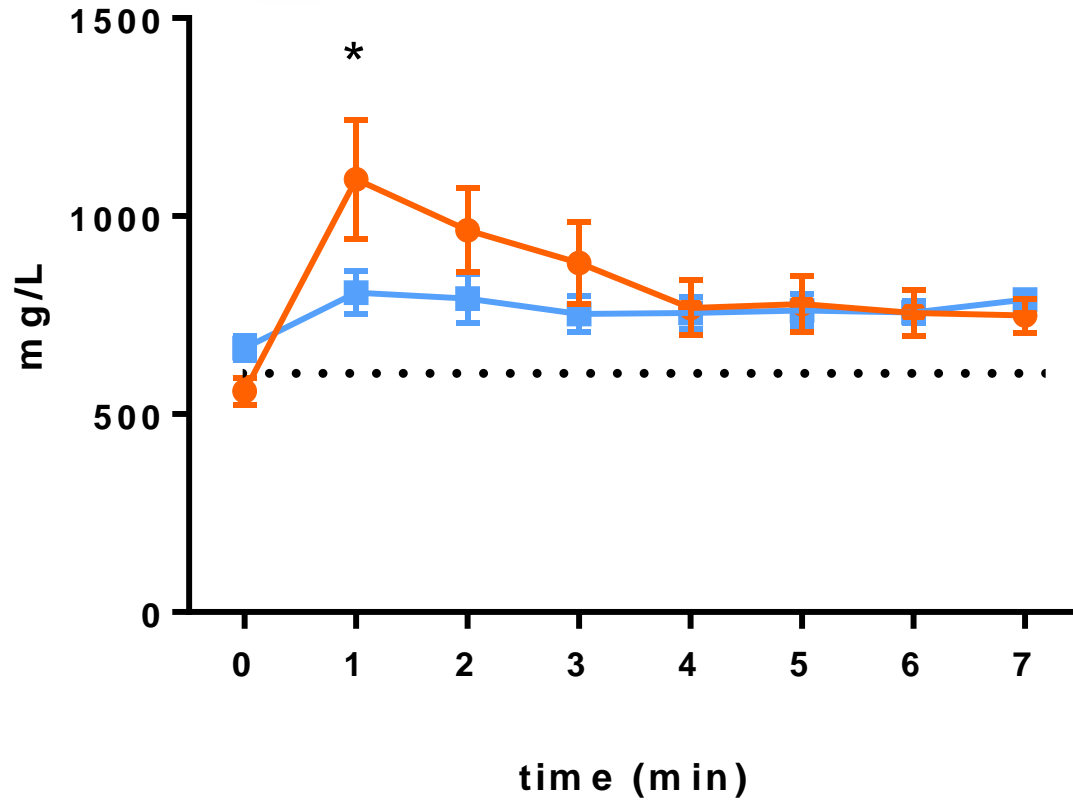
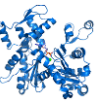
CS Gels (right) are an agglomerate of spherical particles that can easily dissociate. The gel have a very « loose » structure =

**Protein precipitate**



# Plasma amino acids

Molecular scale



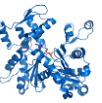
■ CasMic  
● CasNa

casein x time  
 $P = 0.0007$

**Boulier *et al.*  
Food Chem. 2023**

AA peak after 1h for CM whereas the concentration remains stable for CS





## Conclusion


- \* CM form a strong coagulum in the stomach leading to a slow and constant release of plasma amino acids up to 7 h
- \* CS form a loose precipitate in the proximal part of the stomach but most of the caseins remain solubilized in the liquid fraction
- \* CS are rapidly metabolized in the small intestine leading to the appearance of a peak of plasma amino acids one hour after protein ingestion

**CM = slow caseins, CS = fast caseins**

## Perspectives

Do some *in vivo* kinetics experiments

Use  $^{15}\text{N}$  labelled-caseins to differentiate endo/exogenous proteins



# From the protein molecular structure to the the food microstructure: The case of egg white gels



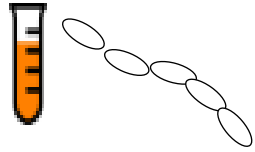
Nau F, & Dupont D.  
INRAE, Rennes, France



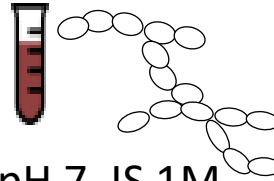
# The microstructure of egg-white gels made from different types of aggregates affects the kinetics of proteolysis

Microscopic scale

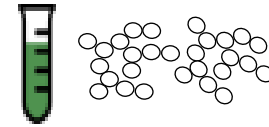
80°C/6h



pH 9, IS 1M



pH 7, IS 1M



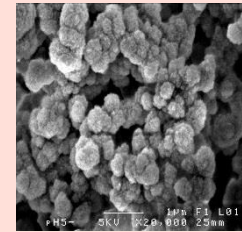
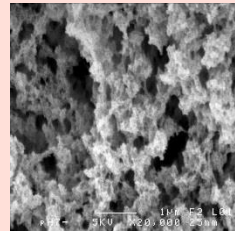
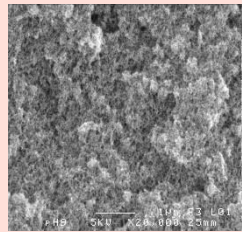
pH 5, IS 1M



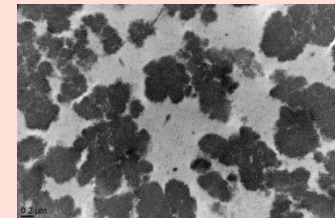
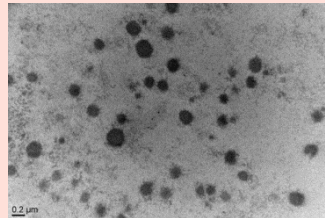
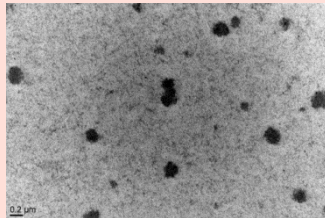
Aggregates	linear	branched	spherical
Rate of <i>in vitro</i> digestion	+++	++	+

Gels 90°C/2.5 h			
-----------------	--	--	--

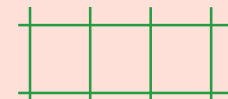
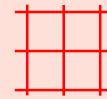
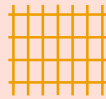
Nyemb *et al.*  
Food Hydro. 2016  
Nyemb *et al.*  
Food Res Int 2016



SEM



CRYO-TEM



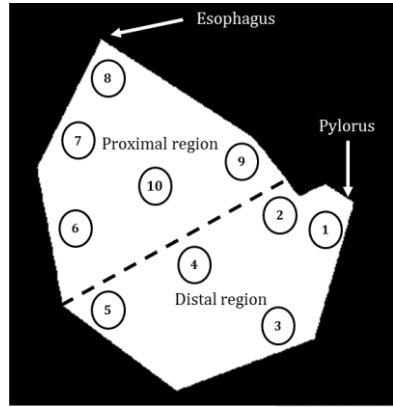
Rate of *in vitro* digestion +

++

+++

# Spatial-temporal evolution of pH during an *in vivo* digestion

Microscopic scale



20 min



60 min



120 min



240 min

pH 7.0

n=33



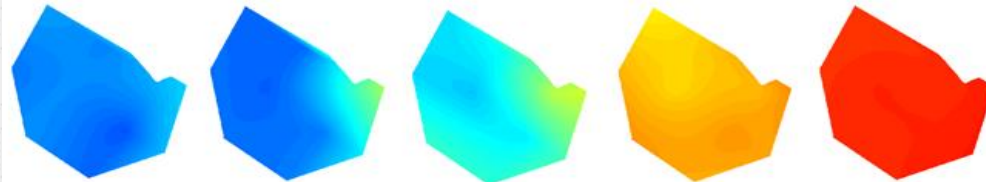
Gel pH 5



n=33



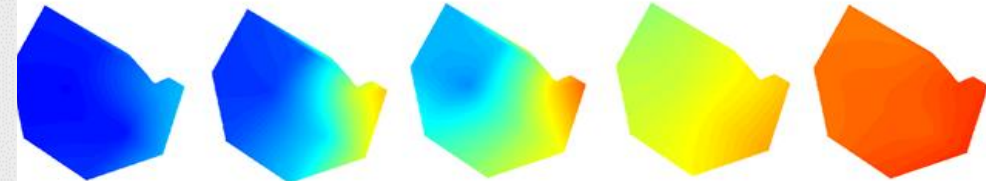
Gel pH 7



n=33



Gel pH 9



20 min

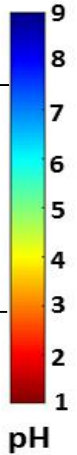
60 min

120 min

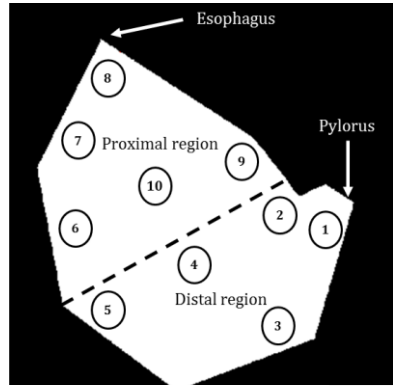
240 min

360 min

Digestion time



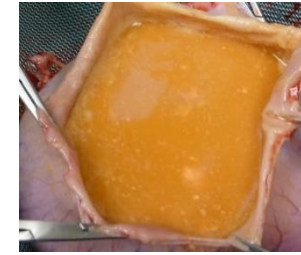
# Spatial-temporal evolution of pH during an *in vivo* digestion



20 min



60 min

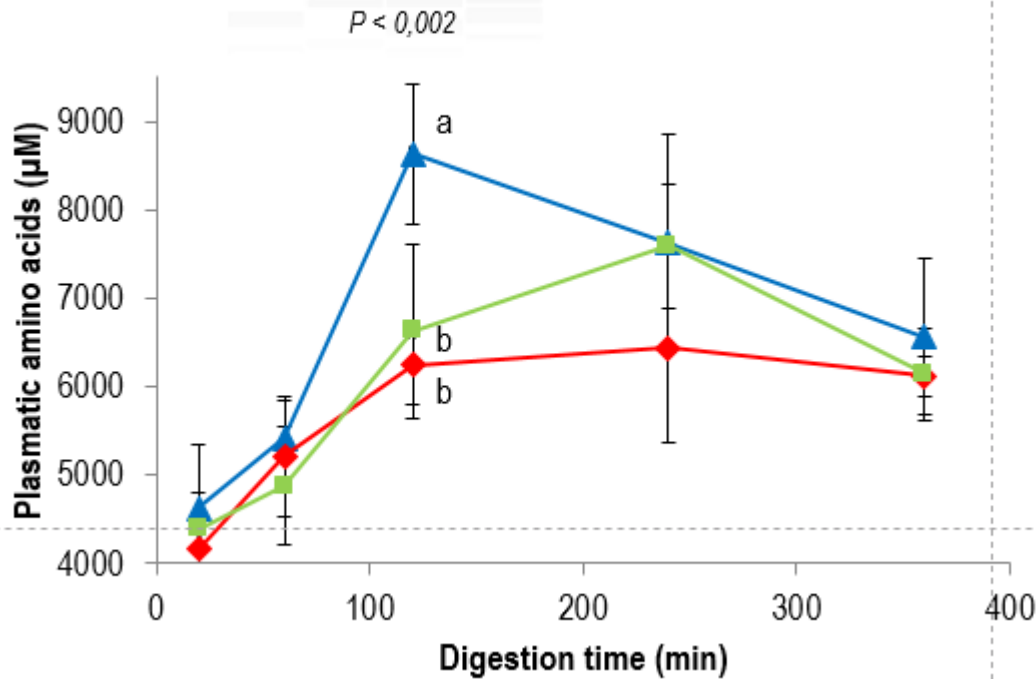


120 min



240 min

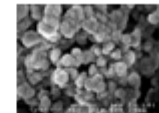
pH 7.0



▲ Granular-spongy EWG

■ Smooth-rigid EWG

◆ Intermediate EWG



Microscopic scale





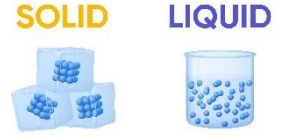
# Food structure as modified by processing affects the kinetics of food digestion



Le Feunteun S, Menard O, Dupont D.  
INRAE, Rennes, France



# Comparison of 6 dairy products of identical composition but different structure



Macroscopic scale

Ultra Low Heat powder



rehydration in water 14.5%

unheated milk ("raw" milk)



heat treatment 90°C-10 min

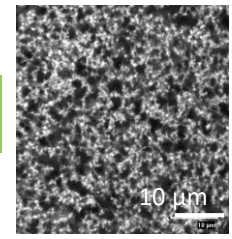
heated milk



macrostructure

rennet gel

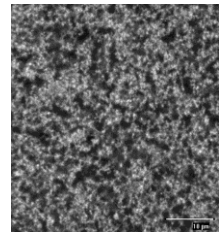
24h-20°C, rennet 0.003 % v/w



pH 6.6

rennet gel

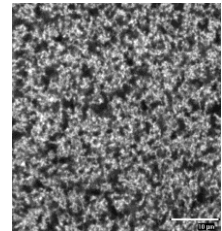
24h-20°C, rennet 0.3 % v/w



pH 6.6

acid gel

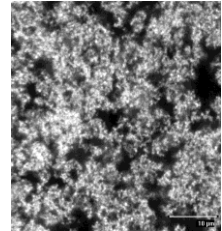
24h-20°C, GDL 3 % w/w



pH 4

stirred acid gel

24h-20°C, GDL 3 % w/w + mixer 2 min



pH 4

Fat-free matrices:

40 g/L caseins, 10 g/L whey proteins, 95 g/L lactose and minerals

+ marker of the meal transit (Cr<sup>2+</sup>-EDTA) → Gastric emptying half-time

# The multi-cannulated mini-pigs

SOLID

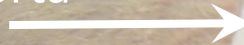
LIQUID



Macroscopic scale

6 minipigs (20 ± 1kg)

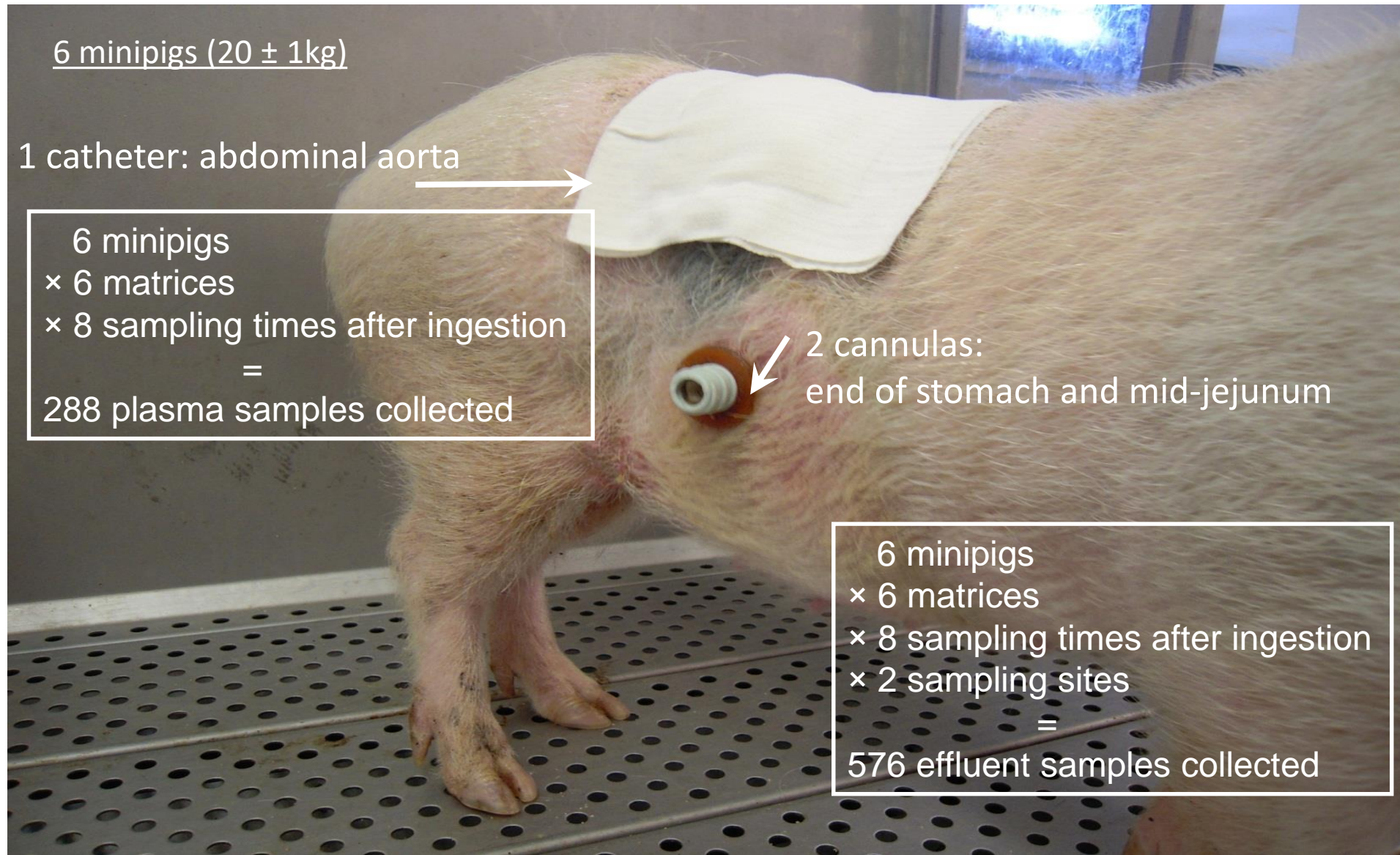
1 catheter: abdominal aorta



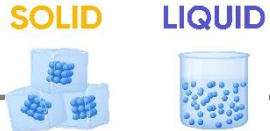
6 minipigs  
× 6 matrices  
× 8 sampling times after ingestion  
=  
288 plasma samples collected

2 cannulas:  
end of stomach and mid-jejunum

6 minipigs  
× 6 matrices  
× 8 sampling times after ingestion  
× 2 sampling sites  
=  
576 effluent samples collected

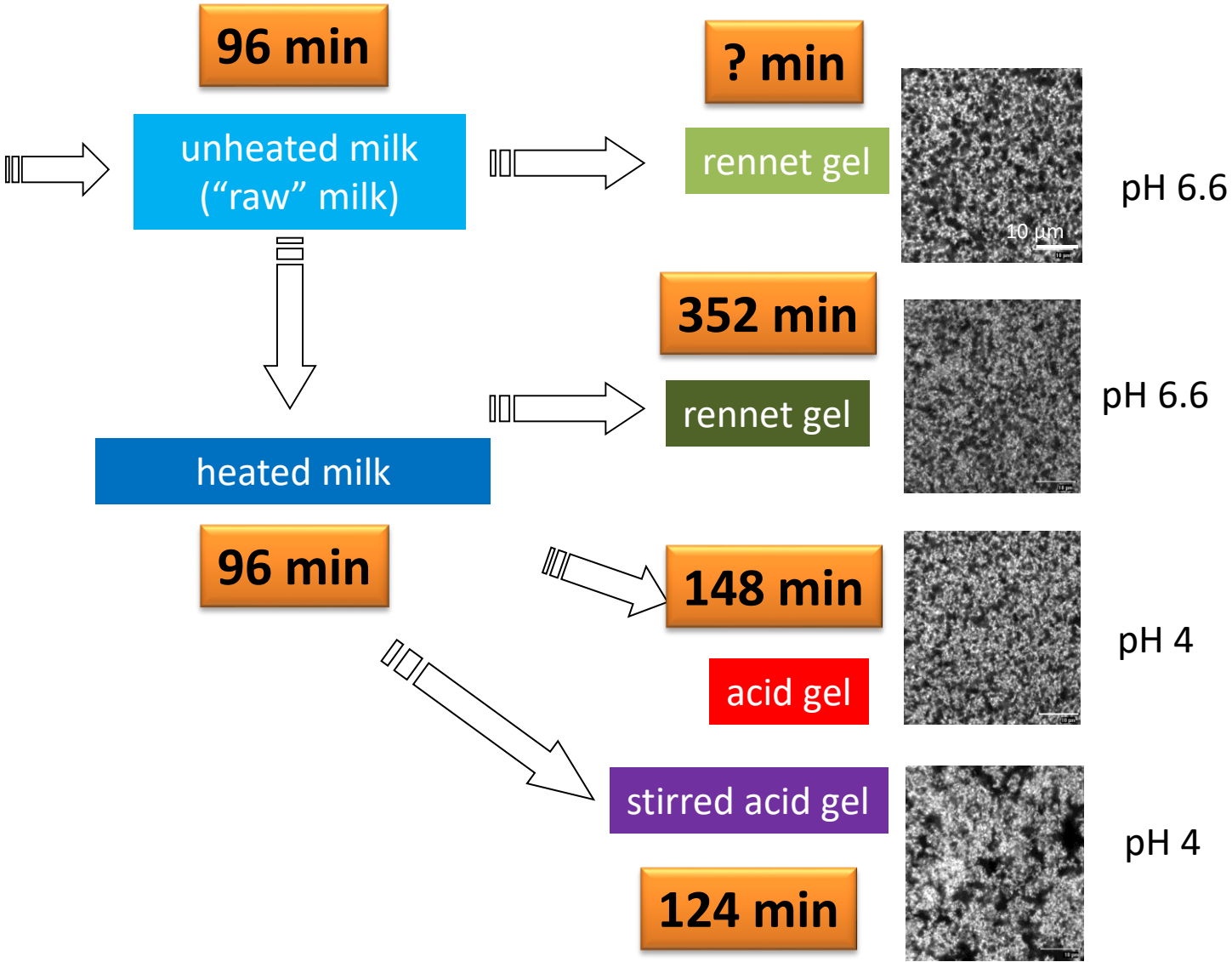


# Gastric emptying half time



Macroscopic scale

Ultra Low Heat powder



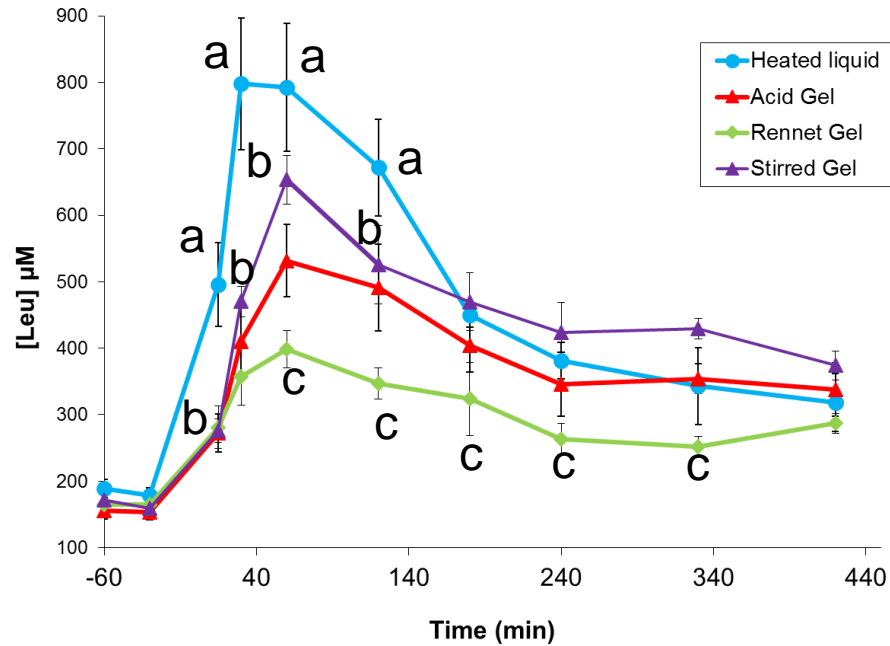


# The liquid-gel transition



Macroscopic scale

## Effect on absorption

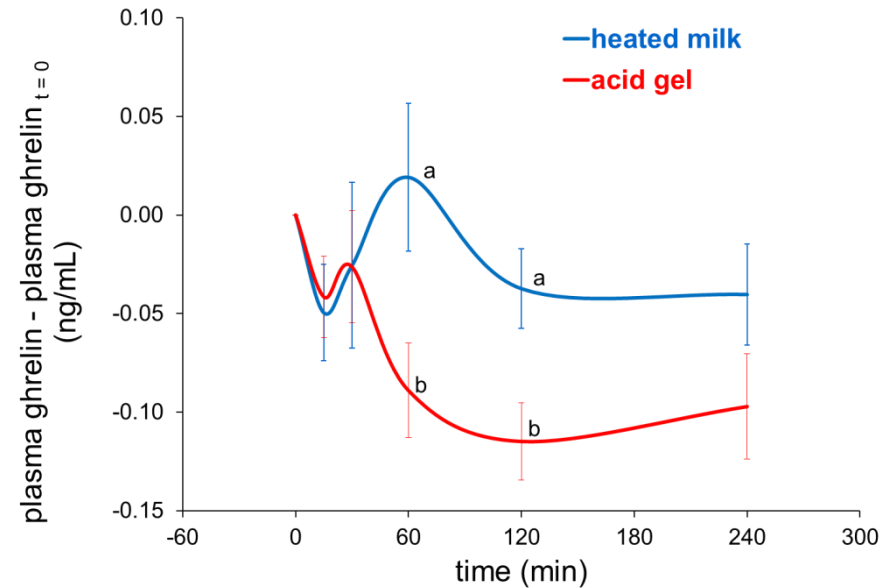


milk gelation:

→ delayed proteins transit → delayed AA absorption  
maximal AA concentration in the plasma

## Potential effect on satiety

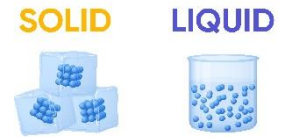
ghrelin (gastrointestinal hormone → appetite stimulation)



milk gelation:

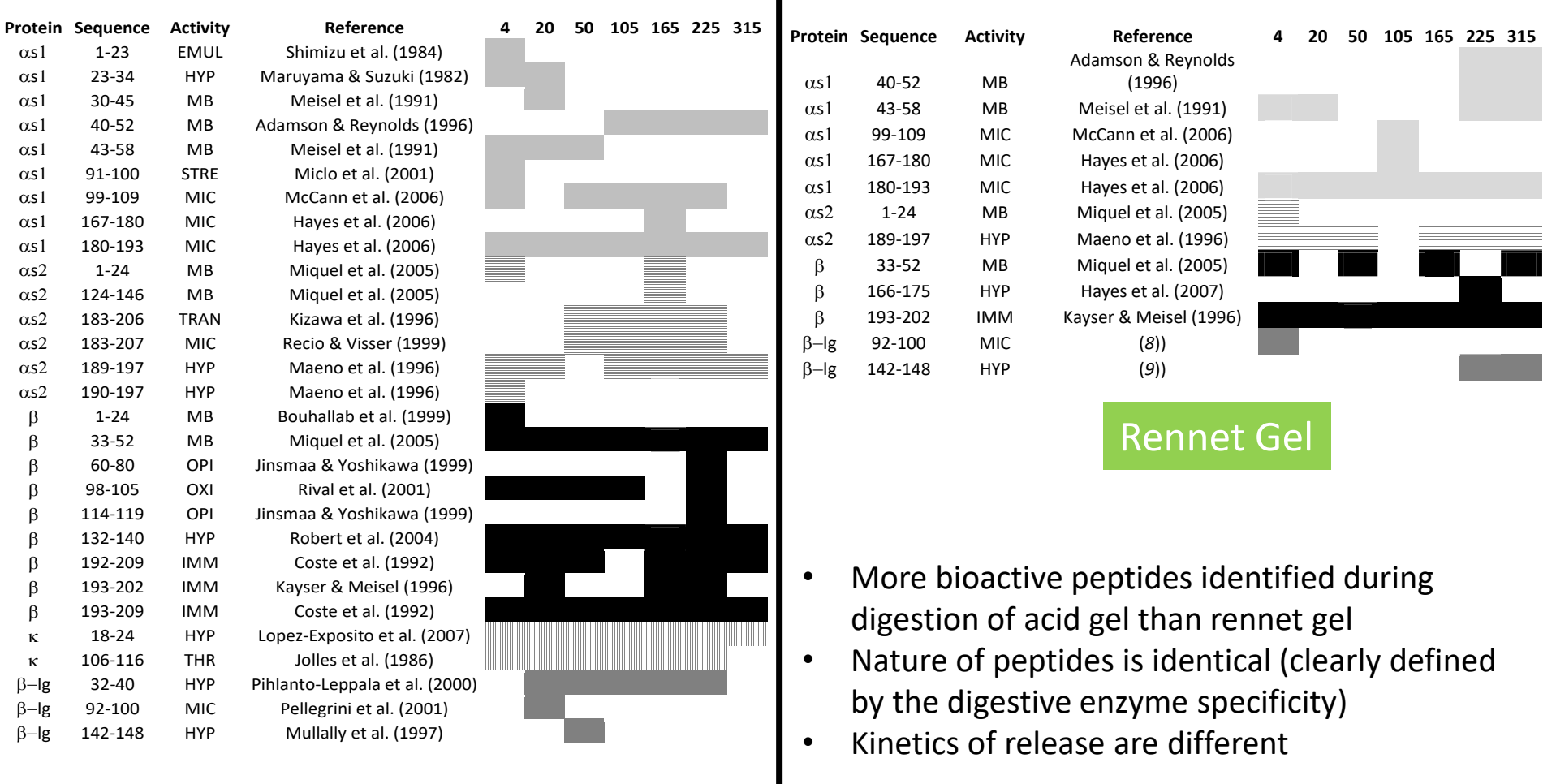
↘ postprandial ghrelin concentration =  
↑ satiety ?

# Bioactive peptides released during digestion differ from one matrix to another



Macroscopic scale

More than 16000 peptides identified by LC-MS-MS in the jejunum



Acid Gel

Rennet Gel

- More bioactive peptides identified during digestion of acid gel than rennet gel
- Nature of peptides is identical (clearly defined by the digestive enzyme specificity)
- Kinetics of release are different

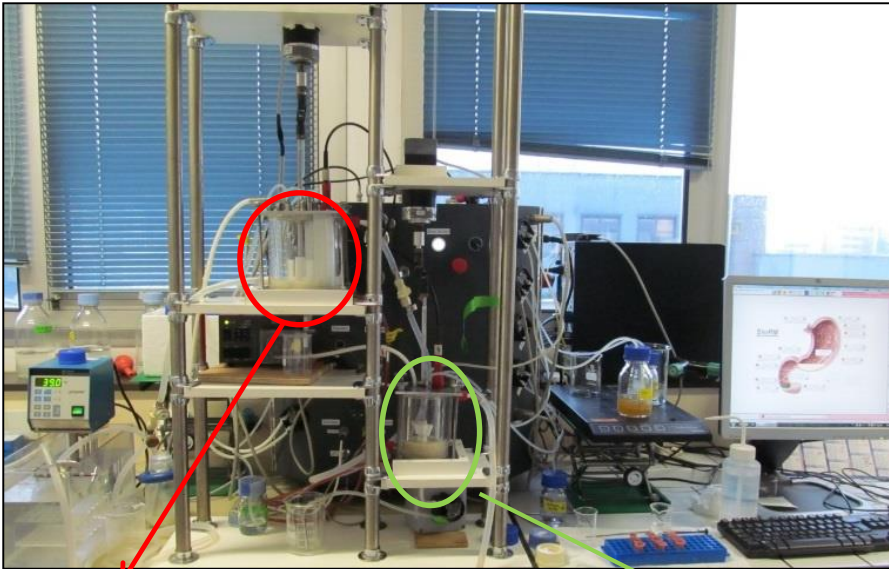
Barbé et al. 2014  
Food Res Int



# Differential behaviour of acid/rennet gels in gastric conditions

- ☞ Acid/Rennet gel: identical composition, similar rheological properties and pore size
- ☞ ≠ Time of residence in the stomach (Acid 148 min /Rennet 352 min)
  - ☞ How can we explain this difference? Dynamic *in vitro* digestion of the 2 gels

**Ménard *et al.*  
Food Chem 2014**



**DIDGI®**

**StoRM® software**

**Stomach**

**Small intestine**

- Pepsine
- Gastric lipase
- Simulated gastric fluid
- HCl



**Emptying :  
Elashoff's model**



- Pancreatin
- Bile
- Simulated intestinal fluid
- NaHCO<sub>3</sub>

**Emptying :  
Elashoff's model**

# Behaviour of acid and rennet gels in the stomach during *in vitro* dynamic digestion

Barbé et al.  
Food Chem. 2014

Acid Gel



Rennet Gel



Formation of a strong coagulum with rennet gel → slow down the gastric emptying of caseins

**The structure that a food adopts in the stomach is essential to understand its digestion**

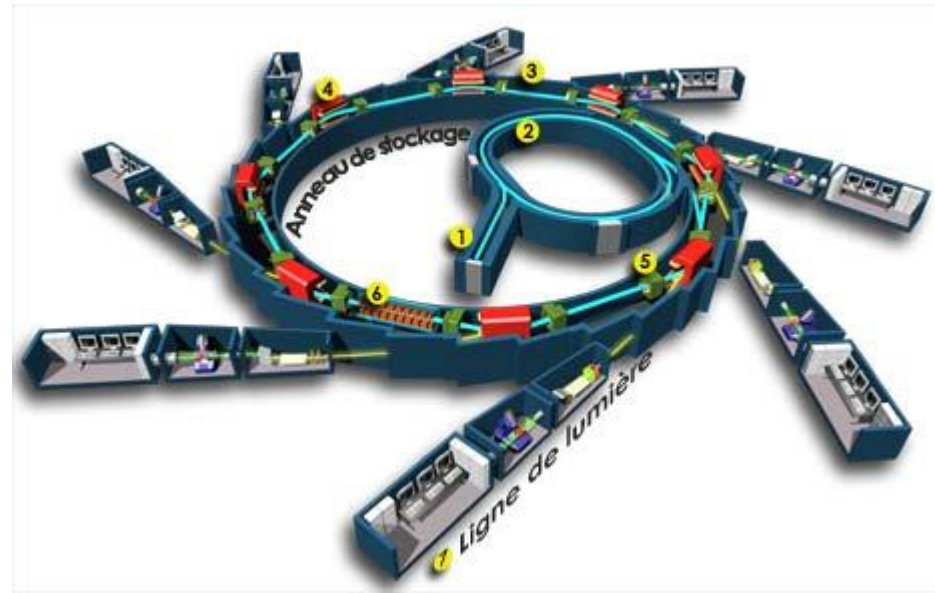


Soleil is a particle (electron) accelerator that produces the synchrotron radiation, an extremely powerful source of light that permits exploration of inert or living matter

DISCO

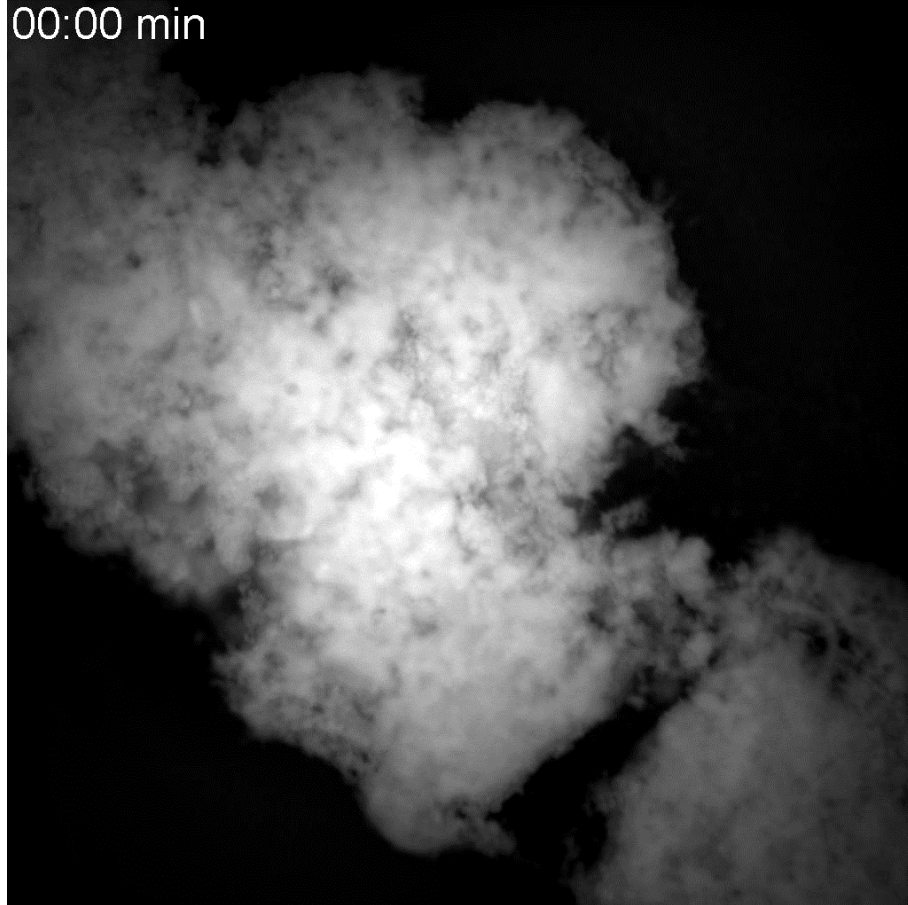
DISCO is a VUV to visible beamline dedicated to biochemistry, chemistry and cell biology. The spectral region is optimized between 60 and 700 nm with conservation of the natural polarization of the light

☞ Allow the imaging of protein intrinsic fluorescence with a UV microscope

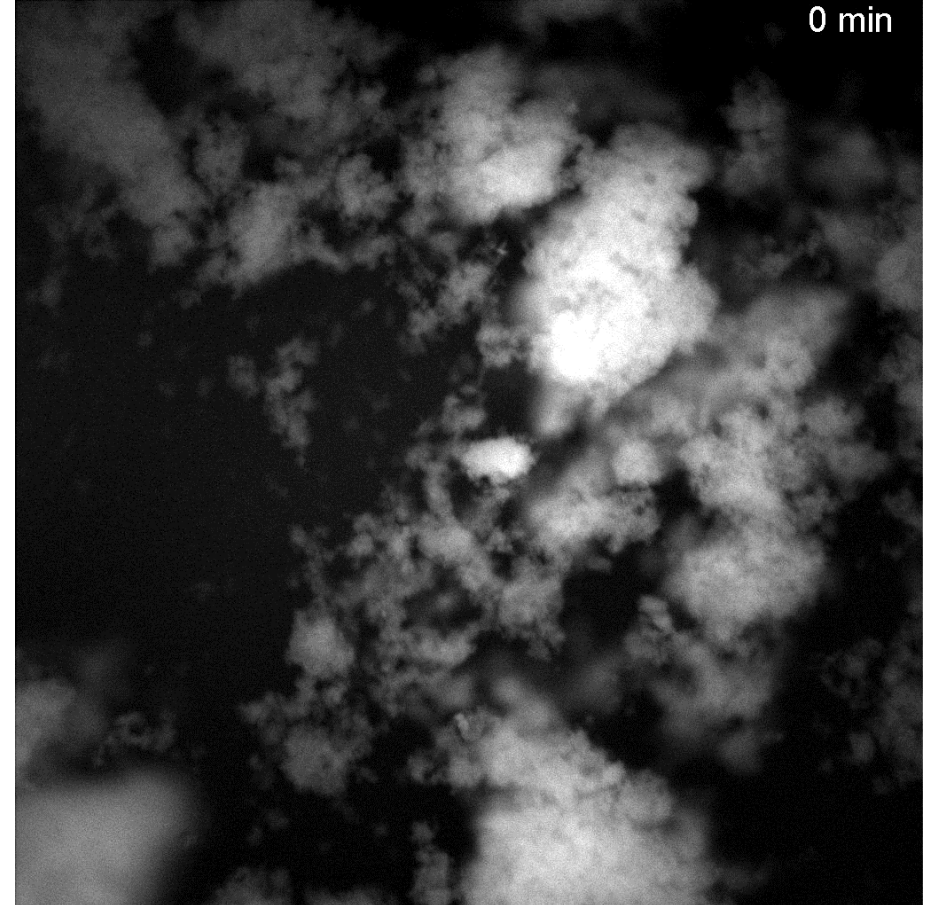




# Kinetics of gel particles disintegration



**Rennet Gel**



**Acid Gel**

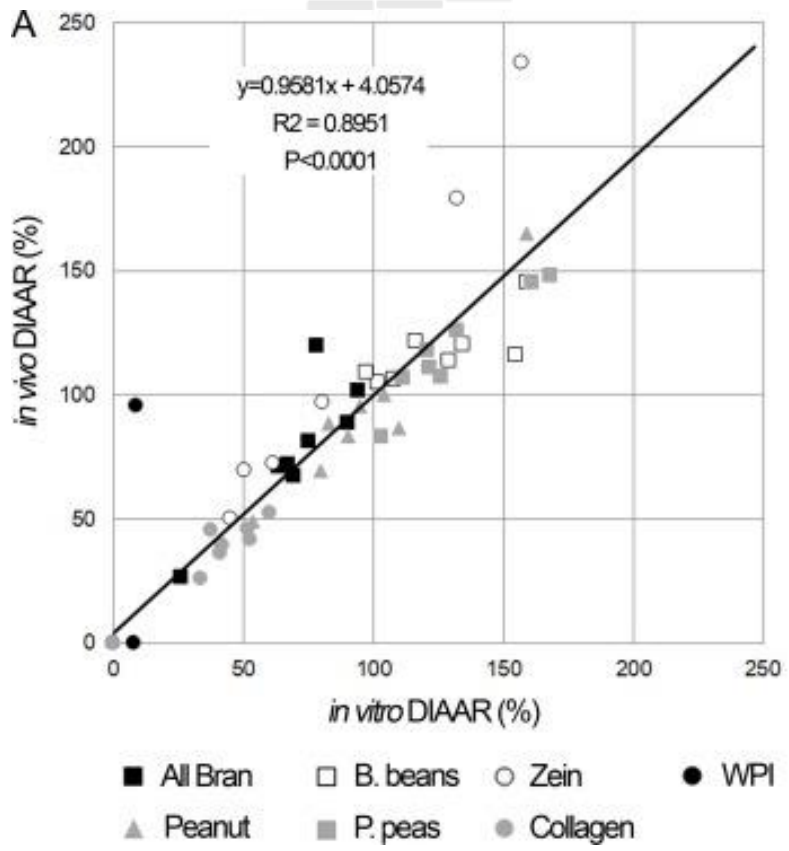
# Can we estimate plant protein digestibility with in vitro digestion models?



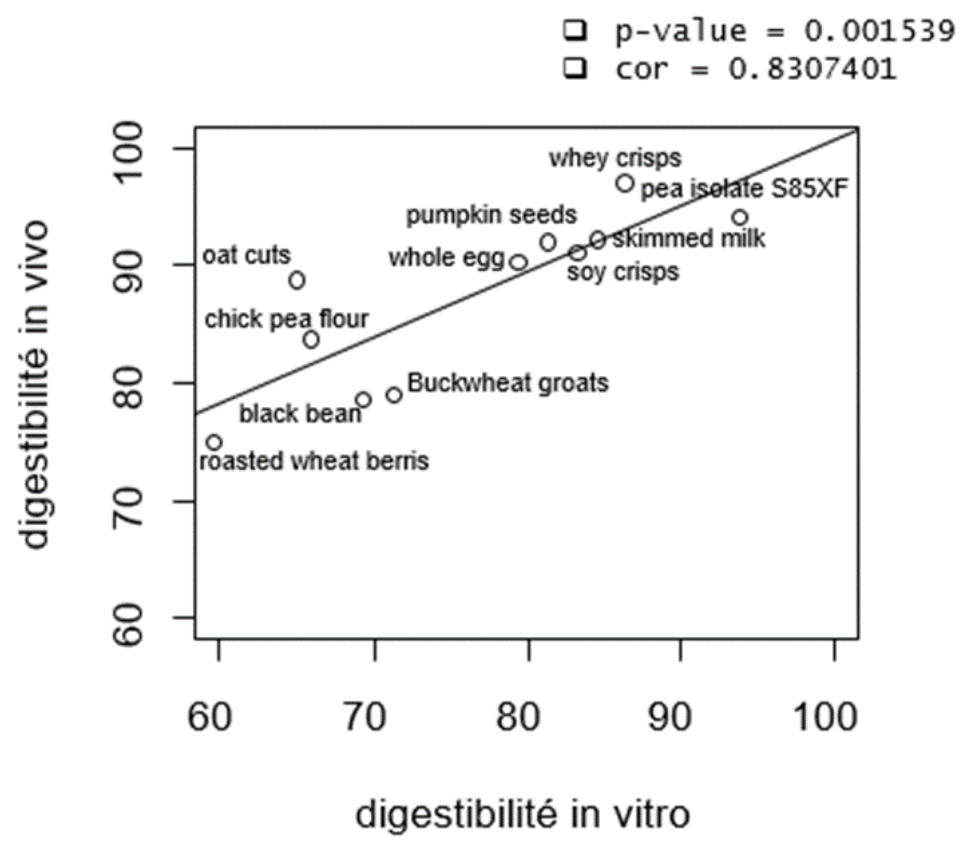
Le Feunteun S, Menard O, Dupont D.  
INRAE, Rennes, France



# In vitro/ in vivo correlation for protein digestibility measurement



Sousa et al. 2023



Nau et al. unpublished

Overall, good correlation are observed but some differences between studies persist



# Protein digestibility with a dynamic *in vitro* digestion model

Study of 4 plant-based foods: 2 solids / 2 liquids

Tofu



Soymilk



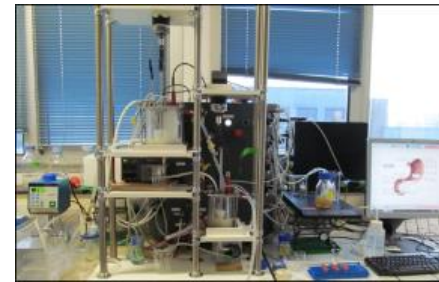
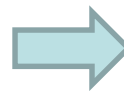
Seitan



Pea Emulsion



**Reynaud et al. 2021  
Food Chem. 341**

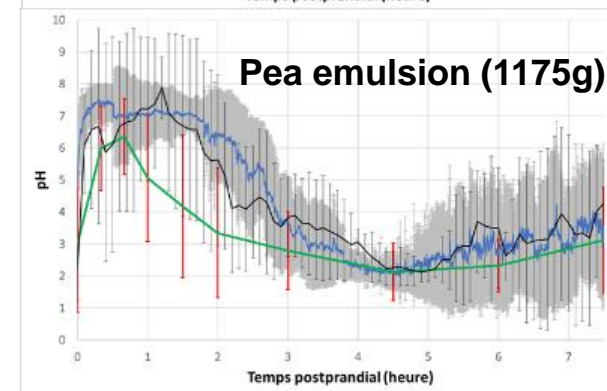
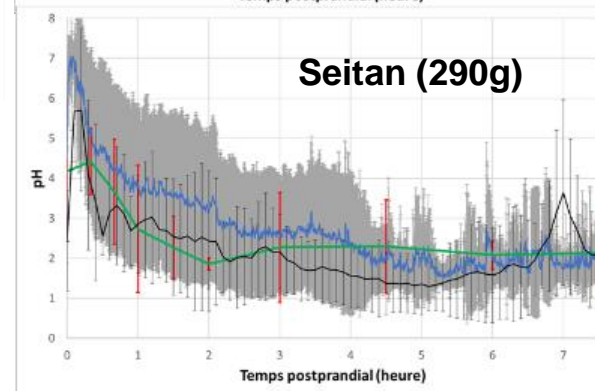
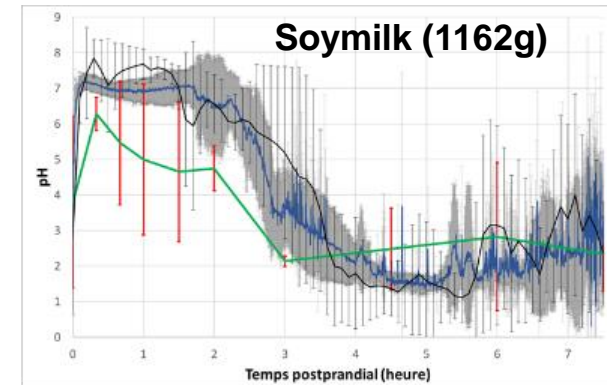
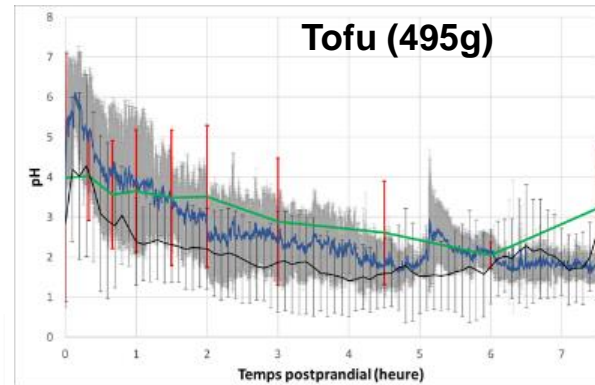
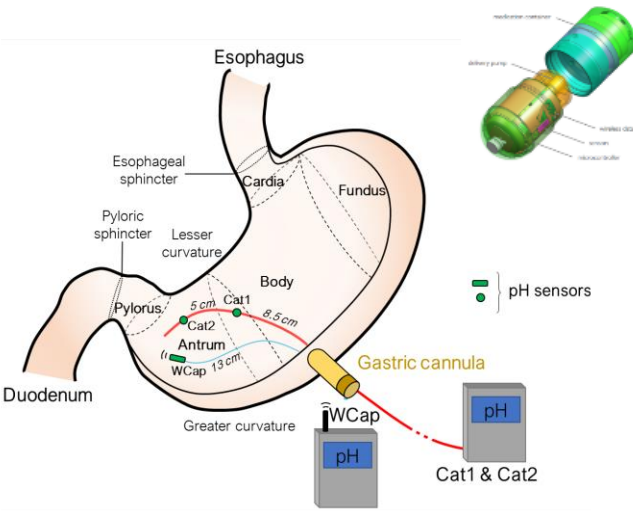


*In vitro*  
digestibility  
(%)

Dynamic *in vitro* digestion DiDGi®

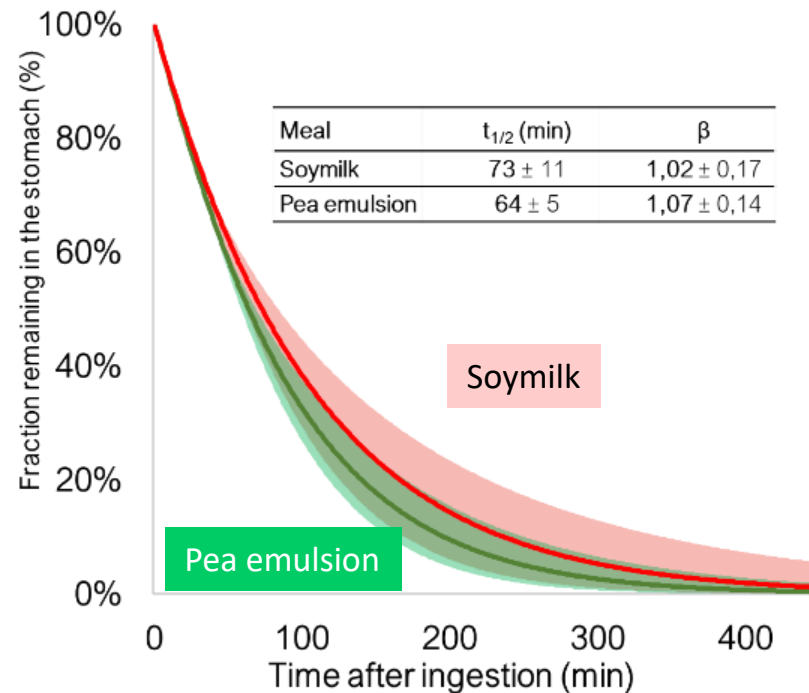
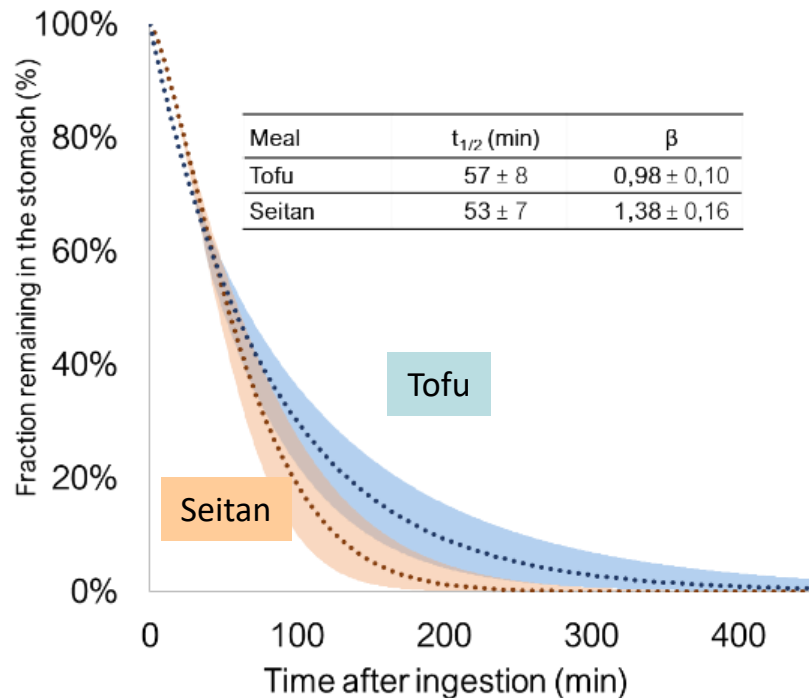
# In vivo data are needed to program the digestion simulator

## Evolution of gastric pH



# In vivo data are needed to program the digestion simulator

## Gastric emptying



**Reynaud et al. 2020**  
**Food Res Int, 128**

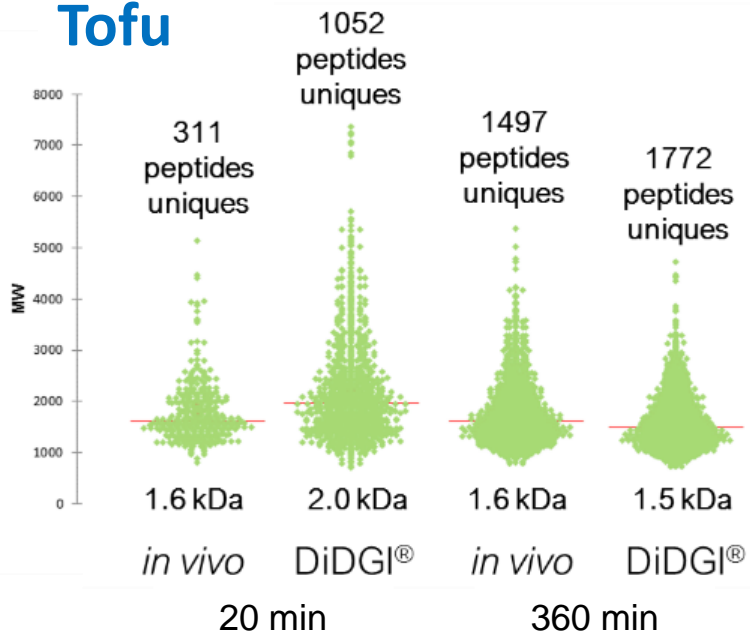
# Comparison between pig and *in vitro* data



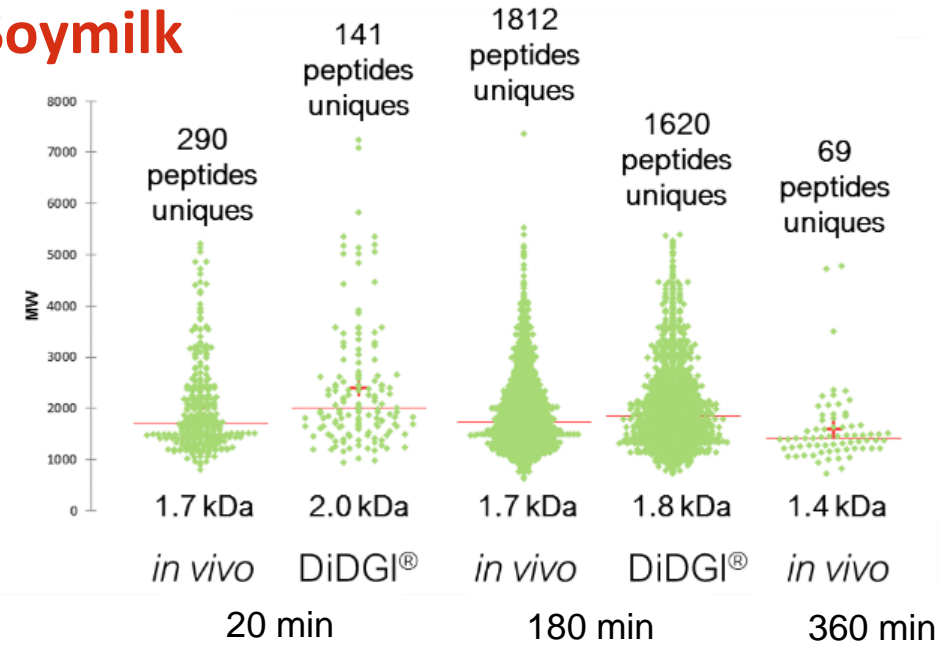
Model	Digestibility	Tofu	Soy milk
<i>in vivo</i>	True	97.1 ± 4.8%	99.4 ± 2.2%
	Apparent	56.5 ± 7.8% <sup>b</sup>	71.3 ± 2.5% <sup>a</sup>
<i>in vitro</i>	Apparent simulated	63.7 ± 3.5% <sup>b</sup>	72.7 ± 1.4% <sup>a</sup>

## Comparison of the gastric peptidome

### Tofu



### Soy milk



# Improving DHA delivery by encapsulation and design of functional foods

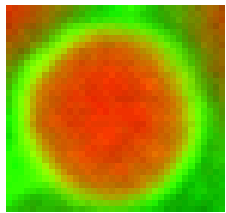
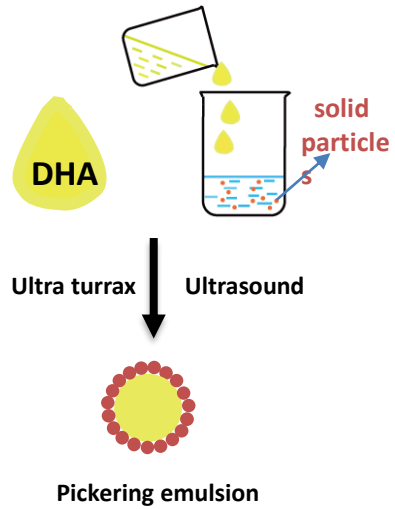


Wang J, Pedrono F, & Dupont D.  
INRAE, Rennes, France



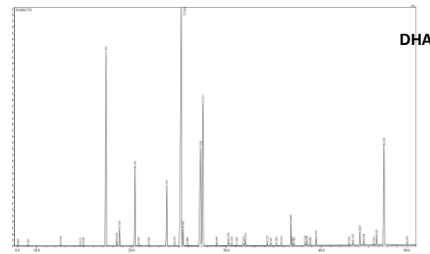
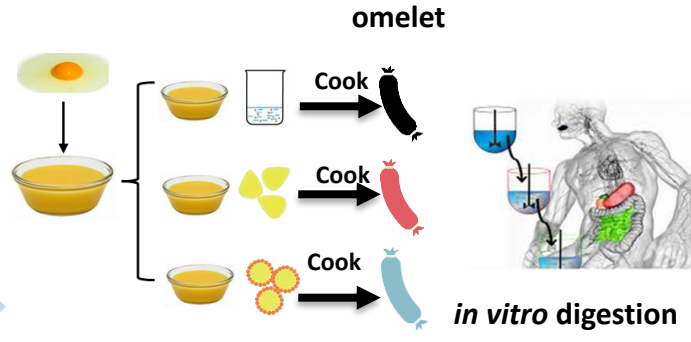
# General strategy

## DHA oil encapsulation



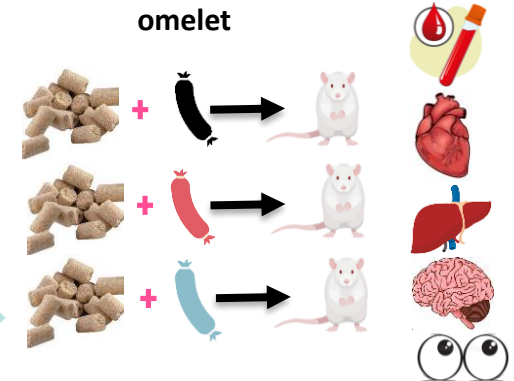
Characterization by CLSM

## DHA bioaccessibility



GC-MS

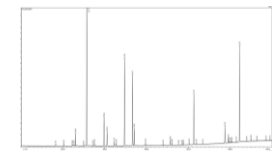
## DHA bioavailability, accretion and metabolism



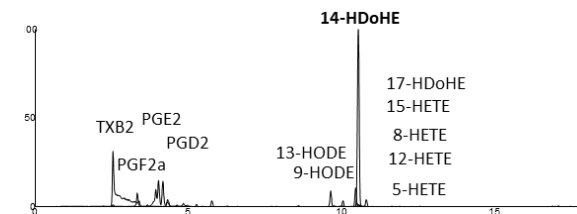
in a rat model



GC-MS



LC-QQQ



Wang et al.  
Food Res Int 2022

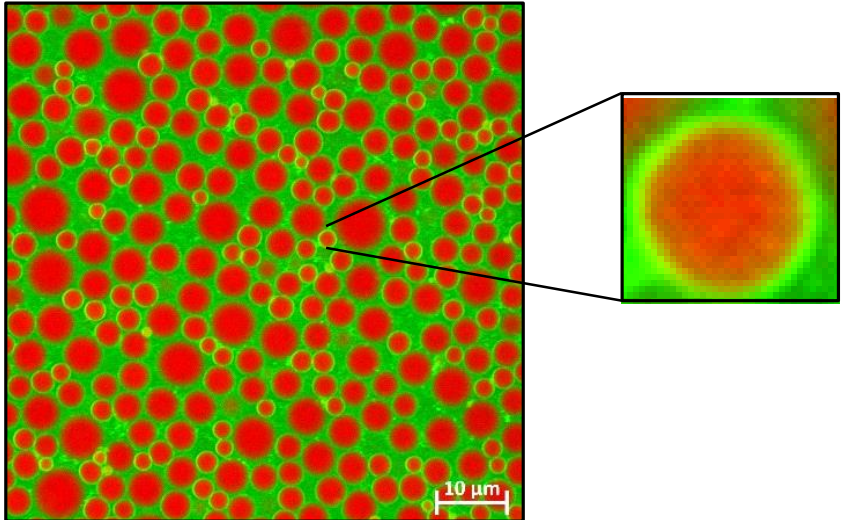
Wang et al.  
Frontiers in Nutr  
2022

Wang et al.  
Nutrients 2023



# DHA oil in emulsion and omelet

## In emulsion

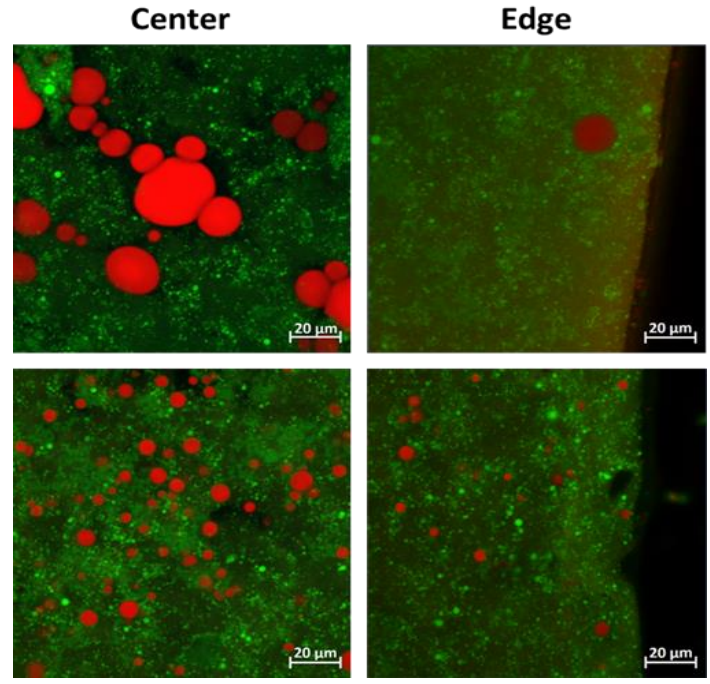


### Encapsulated DHA oil with heat-denatured WPI

DHA oil stained with Nile Red and proteins stained with Fast Green.

The particle size of heat-denatured WPI on average is 42 nm.

## In omelet

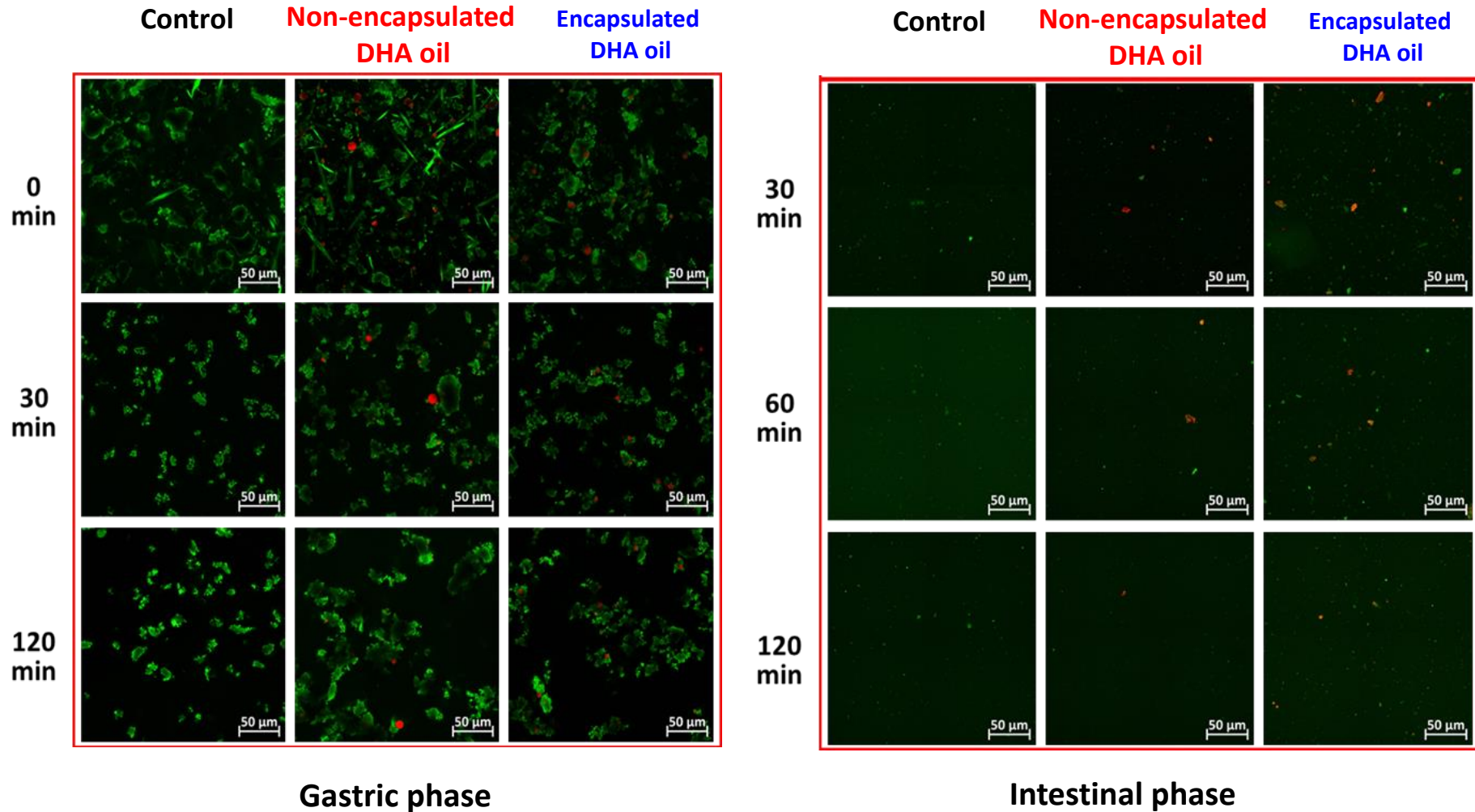


Non-encapsulated DHA oil

Encapsulated DHA oil

### Distribution of non-encapsulated and encapsulated DHA oil in omelets.

# DHA oil in omelet during digestion

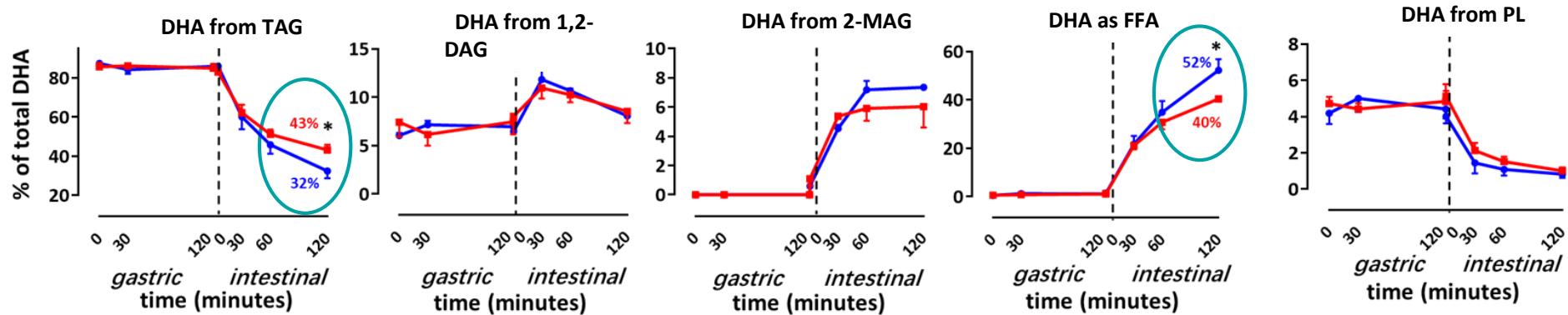


**DHA oil and proteins were mainly hydrolyzed in intestinal phase**

# The release of DHA from different lipid species during digestion

Non-encapsulated DHA oil —■—

Encapsulated DHA oil —●—



The evolution DHA from different lipid species during digestion.

In gastric phase (pepsin and RGE):

- ❑ DHA oil was not hydrolyzed in gastric phase.

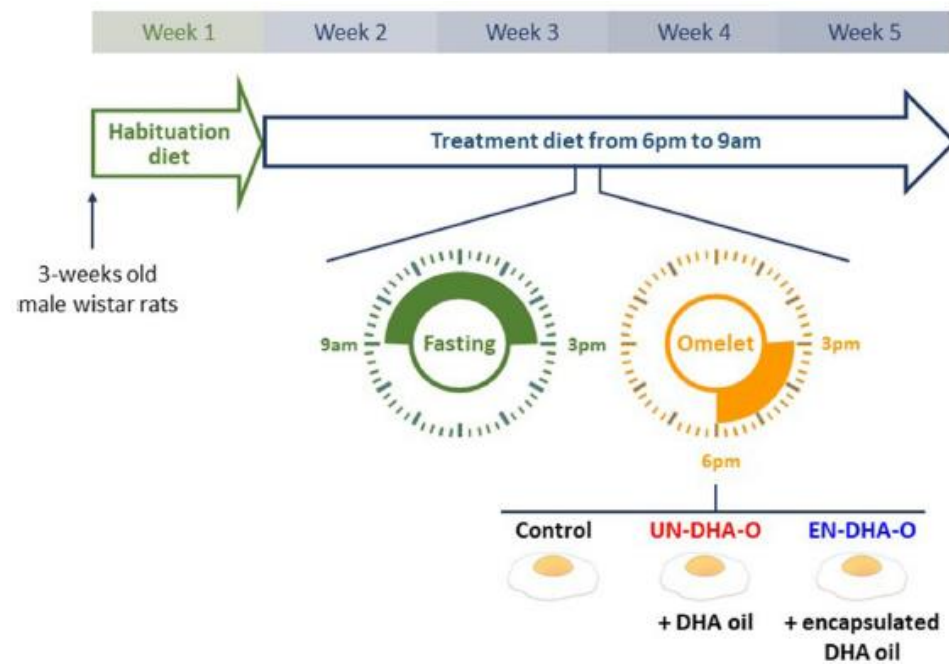
In intestinal phase (bile salt and pancreatin):

- ❑ Hydrolyzed TAG and released FFA encapsulation > unencapsulation

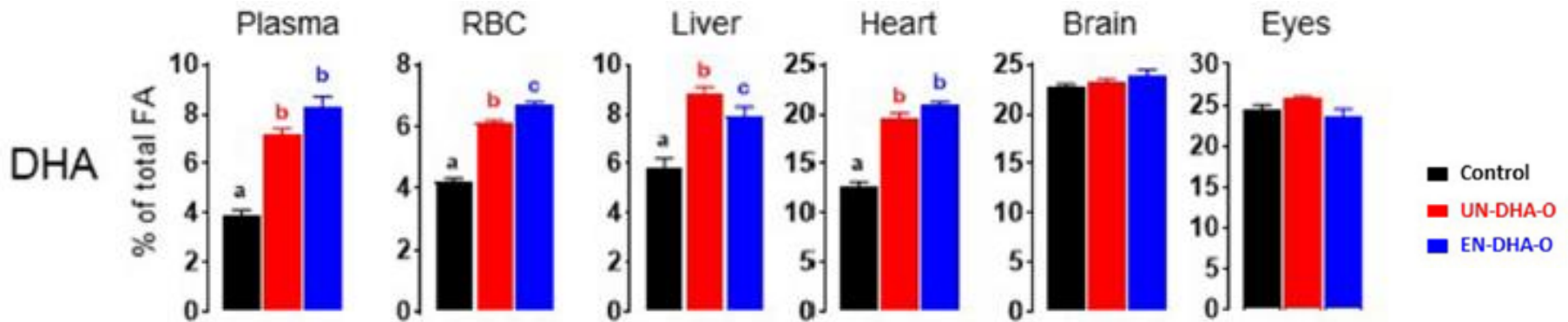
- ❑ Larger interaction area between DHA oil and pancreatic lipase made by emulsification (Maljaars, 2012).

- ❑ Around 10-25% and 40-70% of ingested TAG can be hydrolyzed in gastric and intestinal phase, respectively (Bauer et al; Carriere et al., 1993).

# What happens *in vivo*?



Encapsulation increased DHA concentration in plasma and red blood cells



Encapsulation increased oxylipins in heart and brain (precursors of protectins and maresins that limit inflammation and infection)



# Food structure affects the delivery of hydrophilic and lipophilic micronutrients



Hiolle M., Dupont D. & Nau F.

INRAE, Rennes, France





# Objective of the study:

Demonstrate the effect of the food macrostructure on the bioavailability of Vit B9, Vit B12, Vit D and lutein

## Development of food matrices

Identical composition on dry matter:

- 17 % proteins
- 30 % lipids
- 52 % carbohydrates

Enriched in micronutrients



Biscuit  
DM = 97 %



Sponge cake  
DM = 57 %



Pudding  
DM = 51 %



Custard  
DM = 31 %

## Characterisation of the matrices

Extensive biochemical characterization

Macrostructure : texture analysis

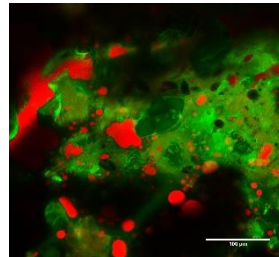
Microstructure : confocal microscopy

## Clinical study

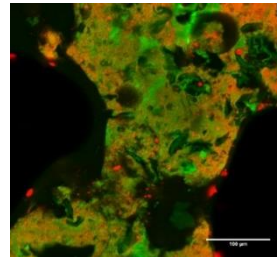
12 healthy volunteers (20-30 y)  
Randomized, controlled, crossover study

Postprandial analyses over 8h

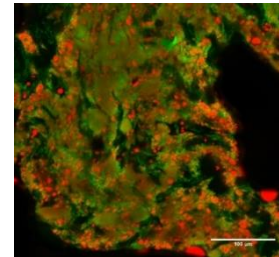
Quantification in the plasma :  
- Vit B9, B12, D and lutein



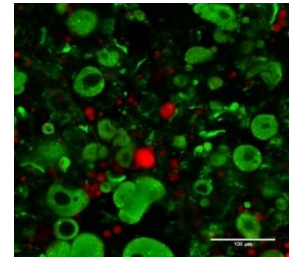
Biscuit



Sponge cake



Pudding



Custard

■ : lipids (Nile Red)

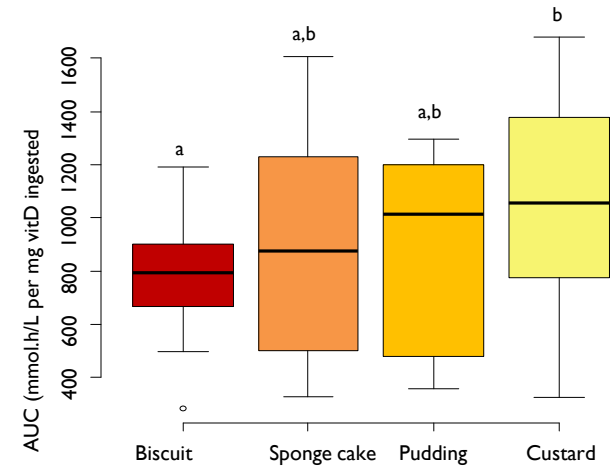
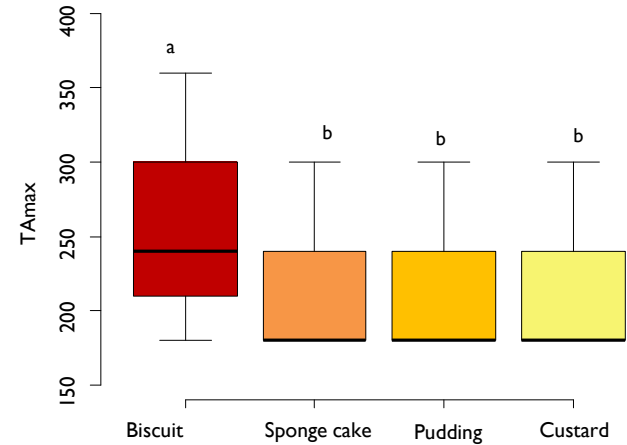
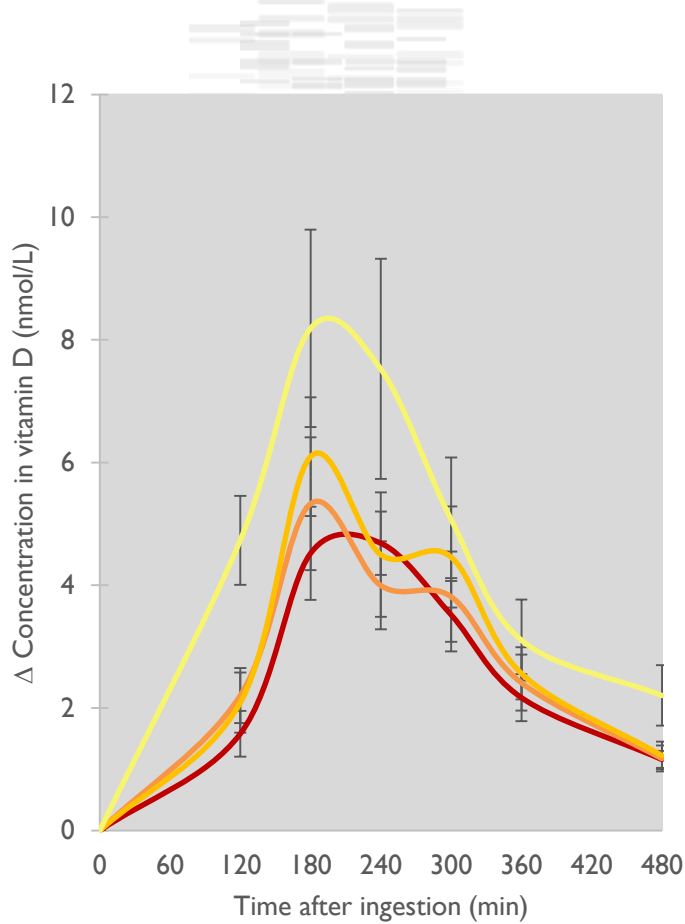
■ : proteins (Fast Green)

Hiolle *et al.*

Food Chem. 2020



# Food structure affects micronutrient kinetics of release and bioavailability



**Buffière *et al.***  
**Food Chem. 2020**

Increased AUC for vitamin D when provided via a custard

# Understanding human milk digestion to design new infant formulas that will have the same behaviour in the GI tract



Deglaire A., Menard O., De Oliveira S., Bourlieu C.  
& Dupont D.

INRAE, Rennes, France



# Human/ bovine milk / Infant Formula

## Lipid globule structure

Human milk

Bovine milk

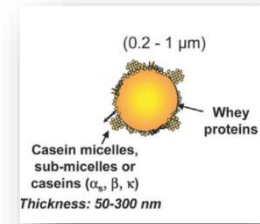
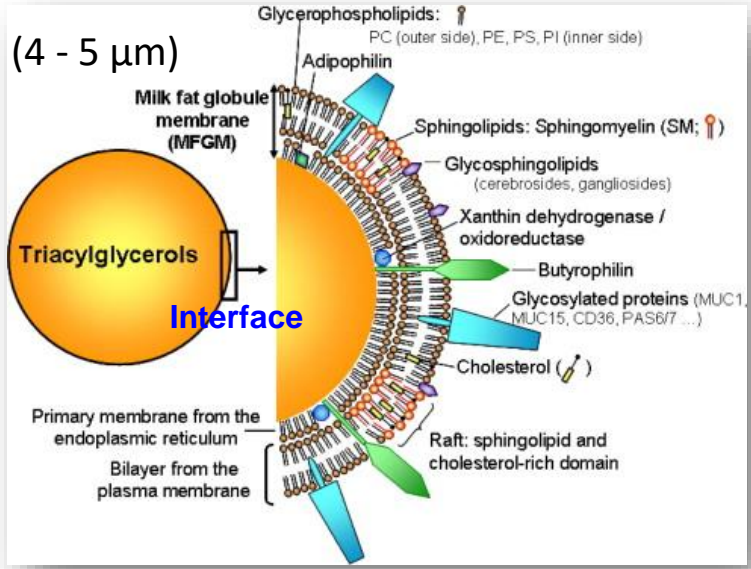
Infant Formula



Native milk fat globule

Lipid droplets

Triacylglycerols



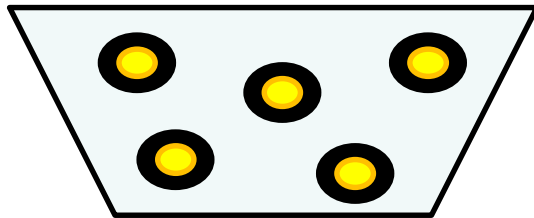
(0,2 - 1 μm)

(Lopez and Briard-Bion, 2007)

(Lopez, 2010)

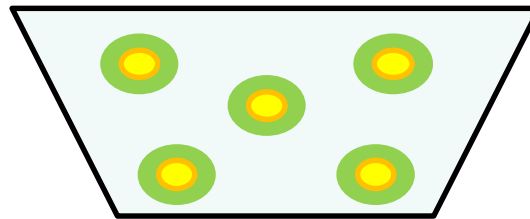
# Infant formulas: can we create lipid structures biomimetic on the native fat globule?

Formula  
T1



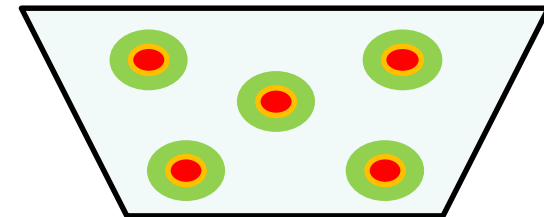
Interface 100 % Proteins  
100% vegetable oil

Formula  
T2

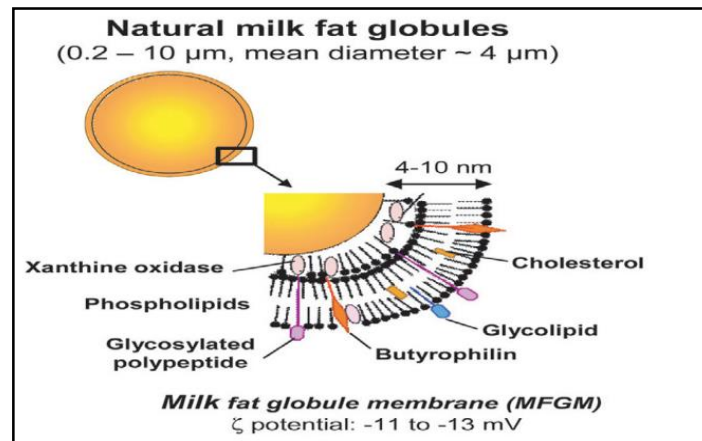


Interface 100 % phospholipids  
100% vegetable oil

Formula  
T3

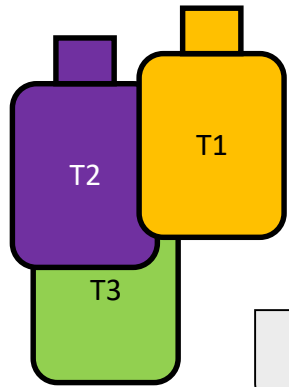


Interface 100 % phospholipids  
40% vegetable oil + 60% milk fat



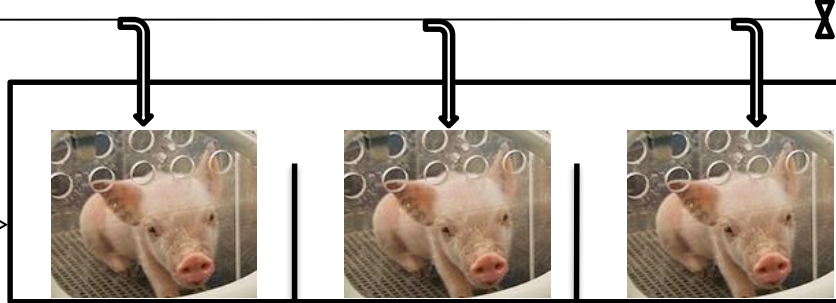
Lopez, (2007)

# Can the composition of infant formula modulate the physiological response of the neonate?



- Veg
- Veg + PL
- Dairy Fat + PL

Automatic meal delivery (10 meals/ day)



+  
Mother-fed piglets  
(MF = + control)



Slaughtering after

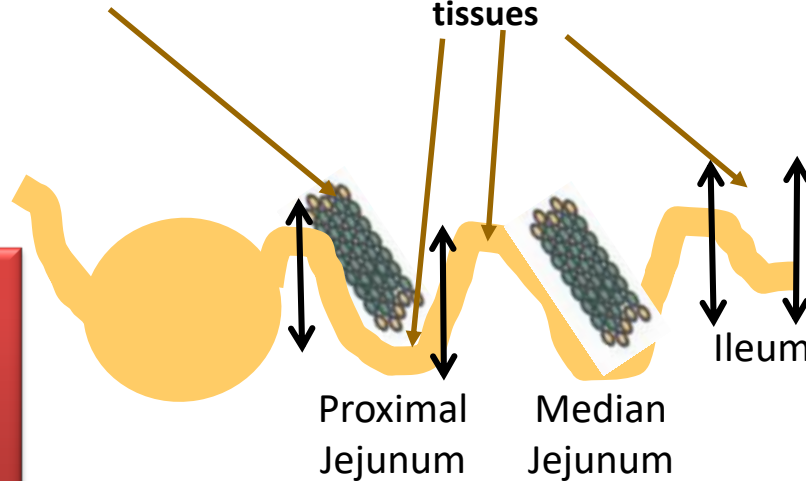
7 days

28 days

(90 min postprandial)

Mesenteric Lymph Nodes (MLN)

Collect of effluents and tissues



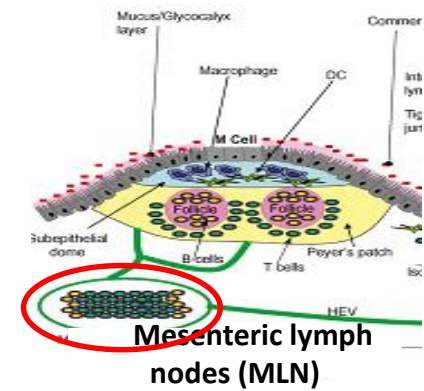
Effluents:  
-SDS-PAGE  
-Elisa

Tissues:

- Morphometry
- Enzyme Activities
- Intestinal Permeability
- Local immune response
- Microbiota

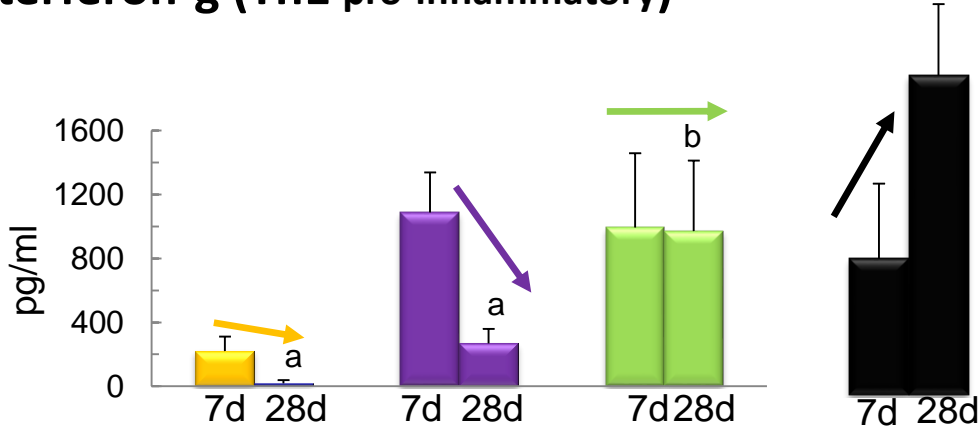


# Secretory activity of MLN

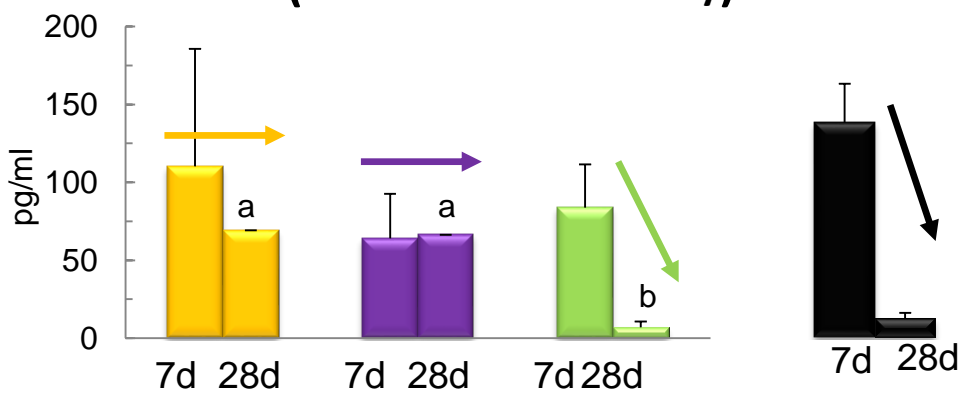


- Veg
- Veg + PL
- Dairy Fat + PL
- Porcelets SM

## Interferon-g (Th1 pro-inflammatory)



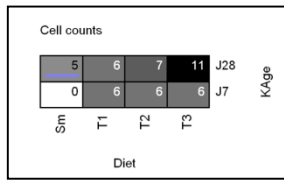
## Interleukine-10 (Th2 anti-inflammatory)



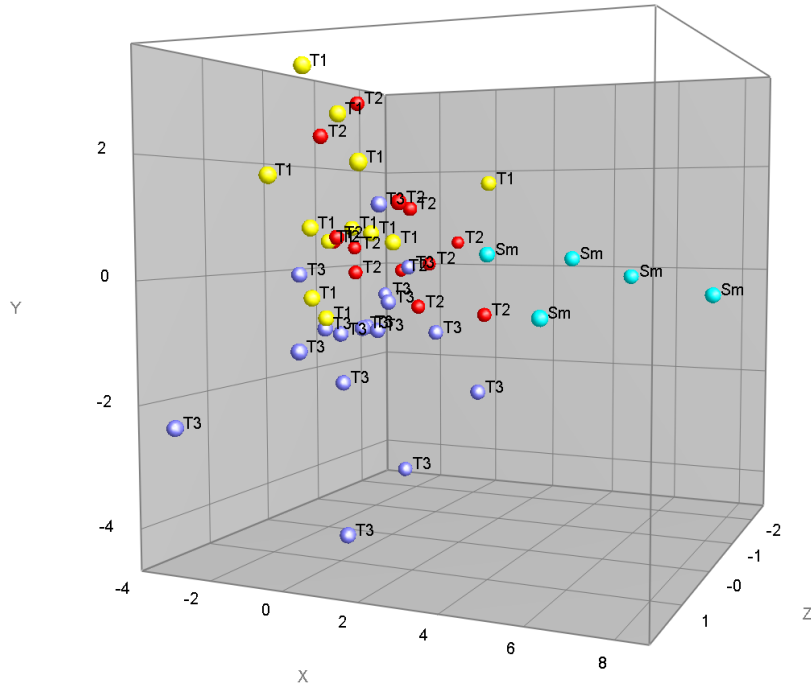
Milk lipids → maturation of the piglet's immune system more similar than with sow's milk

Le Huerou et al.  
Eur J Nutr 2017

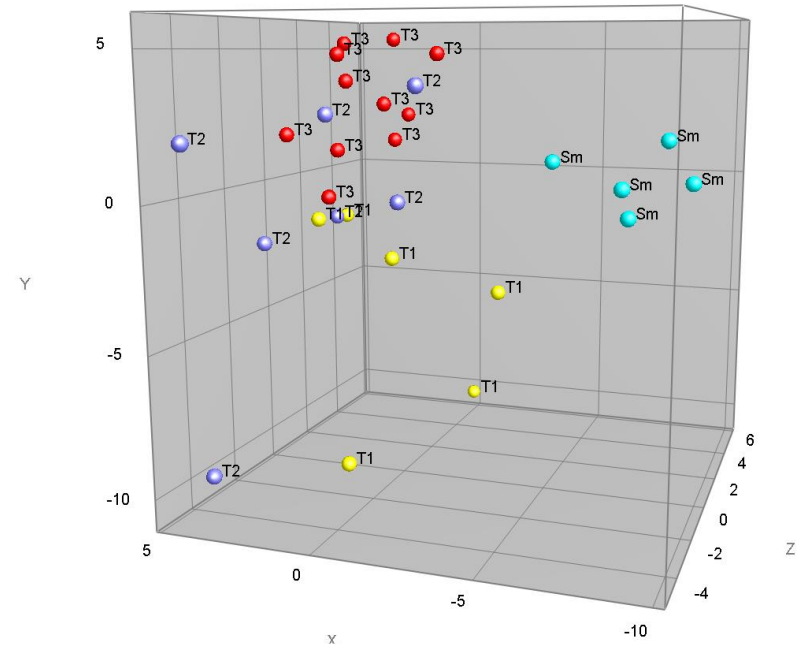
# Microbiota by DHPLC



D7 & D28

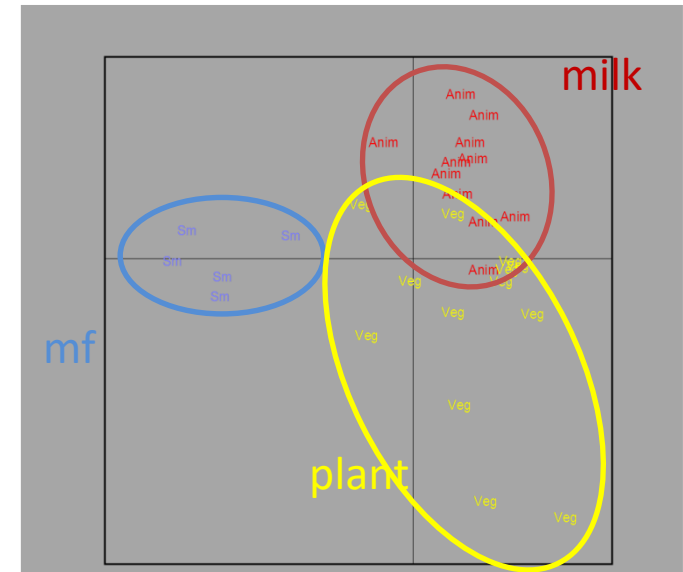


D28

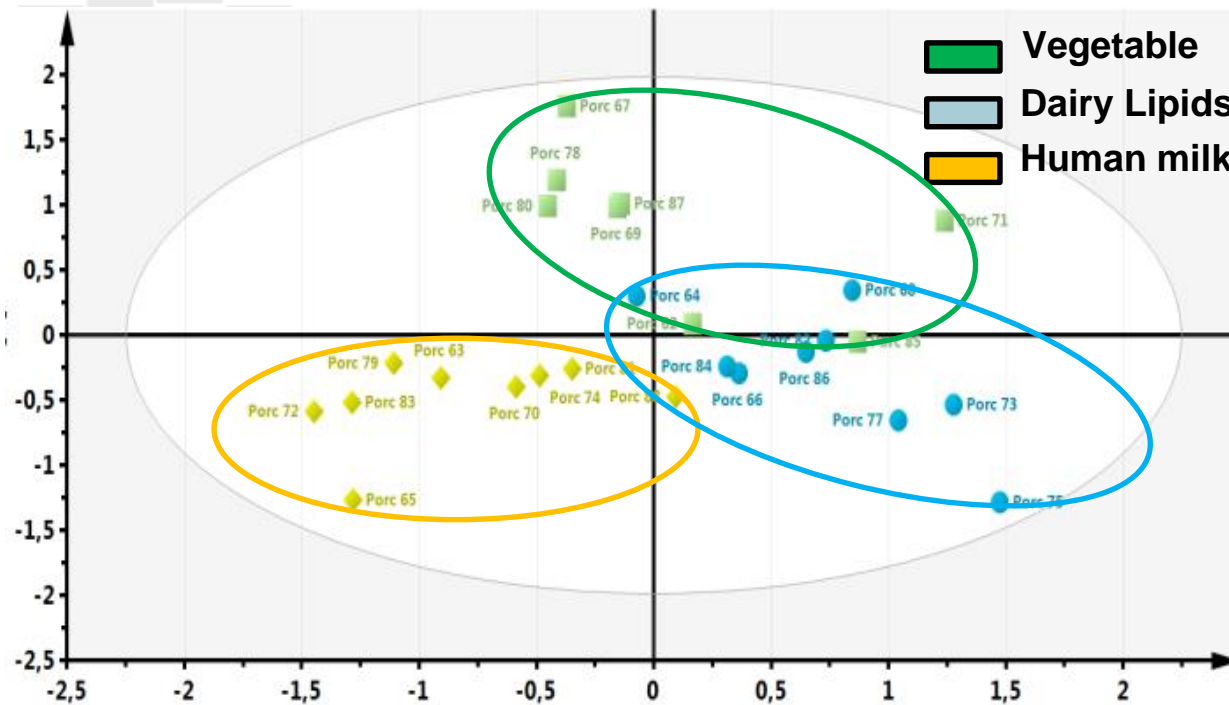


The composition/structure of the infant formula « orientates » the microbiota

More Proteobacteria with milk fat /  
More Firmicutes with plant oil



# What happens when they become older (140 d)?



If animals are submitted to a nutritional stress (high fat/sugar diet), some differences remain in:

- \* the microbiota composition
- \* the fecal metabolome with different metabolites (including propionate)
- \* the immune system with a reduced susceptibility to inflammation with milk lipids

# Conclusion

The structure/composition of food regulate the kinetics of protein digestion and the release of amino acids in the bloodstream

Gastric emptying rate will highly depend on the structure that the product will adopt in the stomach cavity.

Understanding the mechanisms of food particle breakdown in the stomach is critical to control the structure a food will adopt in gastric conditions

Being able to design food structures for controlling the kinetics of hydrolysis of macronutrients will allow to obtain food particularly adapted to specific population



# The Bioactivity & Nutrition team at INRA Rennes



## Head

Didier DUPONT - Senior Scientist

## Scientists

Amélie DEGLAIRE – Lecturer

Juliane FLOURY – Lecturer

Catherine GUERIN - Lecturer

Steven LE FEUNTEUN – Senior Scientist

Joëlle LEONIL – Senior Scientist

Martine MORZEL – Senior Scientist

Françoise NAU – Professor

Frédérique PEDRONO – Lecturer

Xiaoxi YU – Post-doc

Guilherme FURTADO – Post-doc

## PhD students

Yohan REYNAUD (2016-2019)

Amira HALABI (2017-2020)

Jun WANG (2018-2021)

Lea SALLELES (2018-2021)

Elise CHARTON (2019-2022)

Lucile CHAUVET (2019-2022)

Ousmane SUWAREH (2019-2022)

## Technicians

Gwenaële HENRY

Yann LE GOUAR

Nathalie MONTHEAN

## Engineers

Julien JARDIN

Olivia MENARD

Jordane OSSEMOND

## Masters students



# Improving health properties of food by sharing our knowledge on the digestive process

## International Network

Dr. Didier DUPONT, Senior Scientist, INRAE, France

●  
**INFOGEST**  
●



# Main objective: understanding the mechanisms of food digestion

- Develop new *in vitro*, *in vivo* and *in silico* digestion models including some for specific populations (infant, elderly)
- Harmonize the methodologies and propose guidelines for performing experiments
- Validate *in vitro* models towards *in vivo* data (animal and/or human)
- Identify the beneficial/deleterious components that are released in the gut during food digestion
- Determine the effect of the matrix structure on the bioavailability of food nutrients and bioactive molecules



# Industry involvement

~ 60 private companies are following INFOGEST







Chair

Didier Dupont - France  
didier.dupont@inrae.fr



Vice-chair

Alan Mackie - UK



[www.cost-infogest.eu](http://www.cost-infogest.eu)

In vitro  
models of  
digestion  
**WG1**

Food  
interaction –  
meal digestion  
**WG2**

Absorption  
models  
**WG3**

Digestive  
lipases and  
lipid digestion  
**WG4**

Digestive  
amylases and  
starch  
digestion  
**WG5**

In silico  
models of  
digestion  
**WG6**



Isidra Recio



Pasquale Ferranti



Linda Giblin



Myriam Grundy



Daniela Freitas



Choi-Hong Lai



Andre  
Brodkorb



Lotti Egger



Uri Lesmes



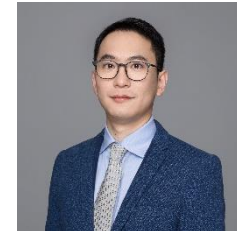
Brigitte Graf



Frederic  
Carriere



Anabel  
Mulet-Cabero



Bin Zhang



Steven Le Feunteun



# Some outputs

## *In vitro* gastrointestinal digestion Consensus INFOGEST protocol

Minekus et al. 2014  
Food & Function, 5, 1113-1124  
**3125 citations**

### Oral phase

Mix 1:1 with Simulated Salivary Fluid (SSF)  
salivary amylase (75 U/mL)  
2 min, pH 7

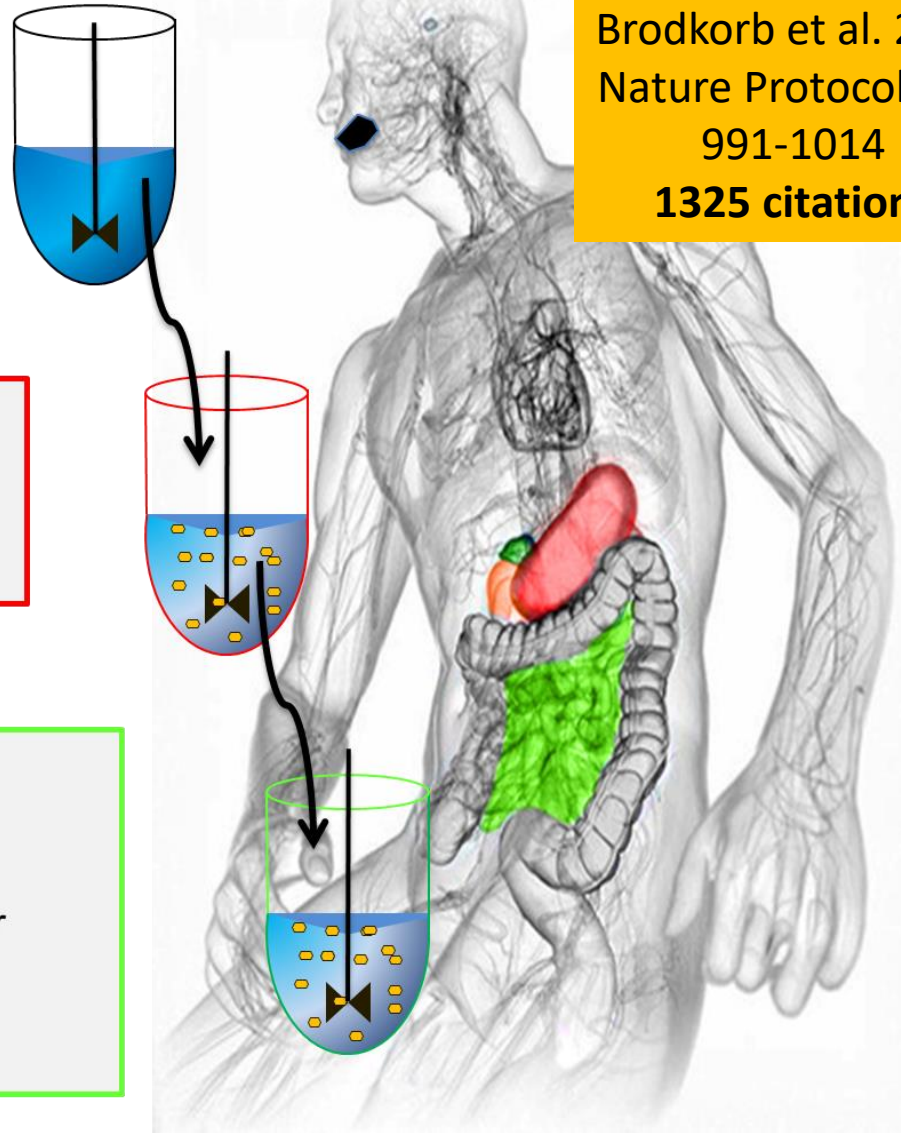
### Gastric Phase

Mix 1:1 with Simulated Gastric Fluid (SGF)  
Pepsin (2000 U/mL)  
2h, pH 3

### Intestinal Phase

Mix 1:1 with Simulated Intestinal Fluid (SIF)  
Enzymes  
Pancreatin (based on trypsin 100 U/mL) or  
Pure enzymes  
Bile (10mM)  
2h, pH 7

Brodkorb et al. 2019  
Nature Protocol, 14,  
991-1014  
**1325 citations**





We are pleased to announce the next  
**8<sup>th</sup> International Conference on Food Digestion**



**in Porto, Portugal, 9-11 April 2024**