PURPAN TOULOUSE FIELD-FLOW FRACTIONATION CENTER





Projet cofinancé par le Fonds Européen de Développement Régional



FIELD-FLOW FRACTIONATION TECHNIQUES FOR MILK PROTEIN CHARACTERIZATION

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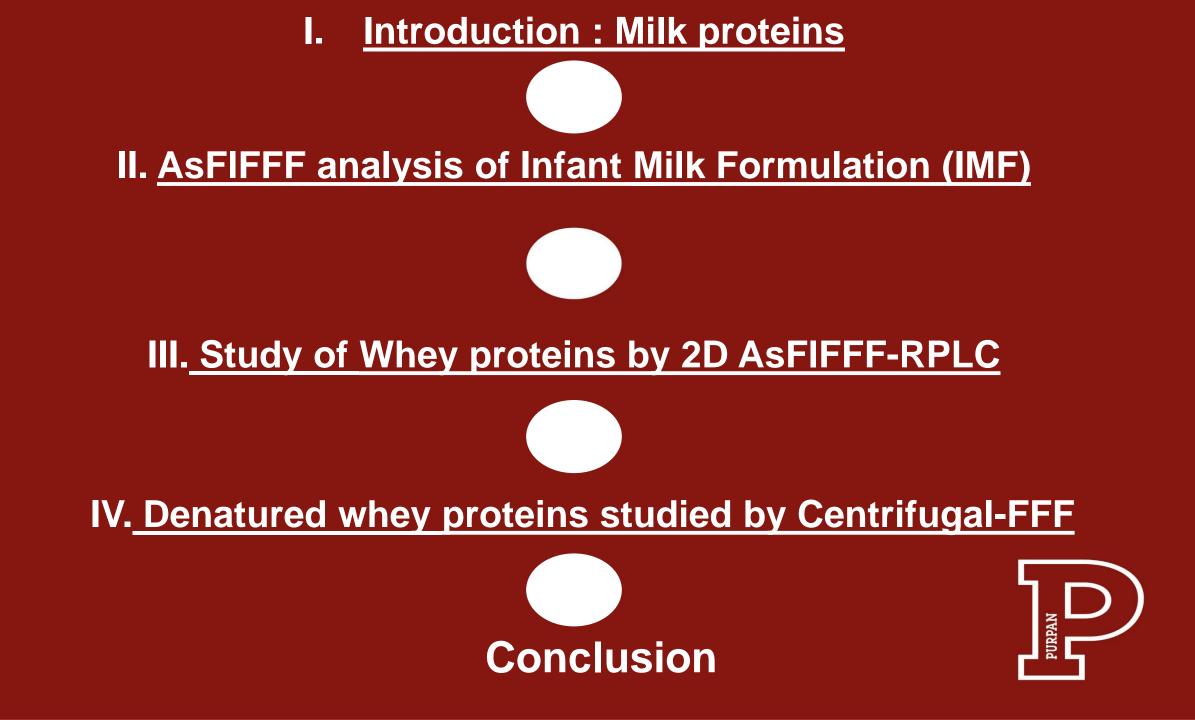
20th International Symposium on Field- and Flow-Based Separations

23–27th February 2020 University of Vienna









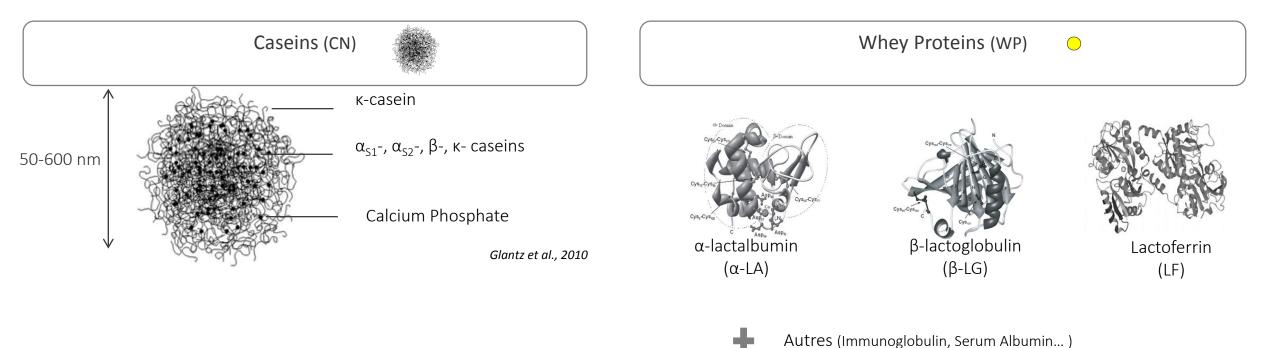




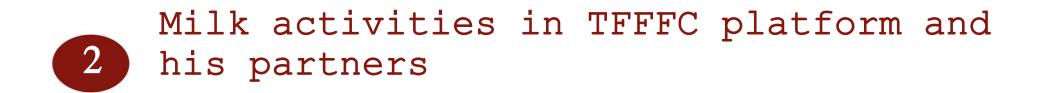




2 categories of milk proteins

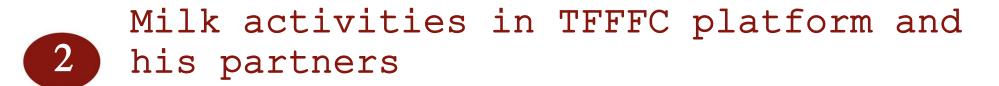


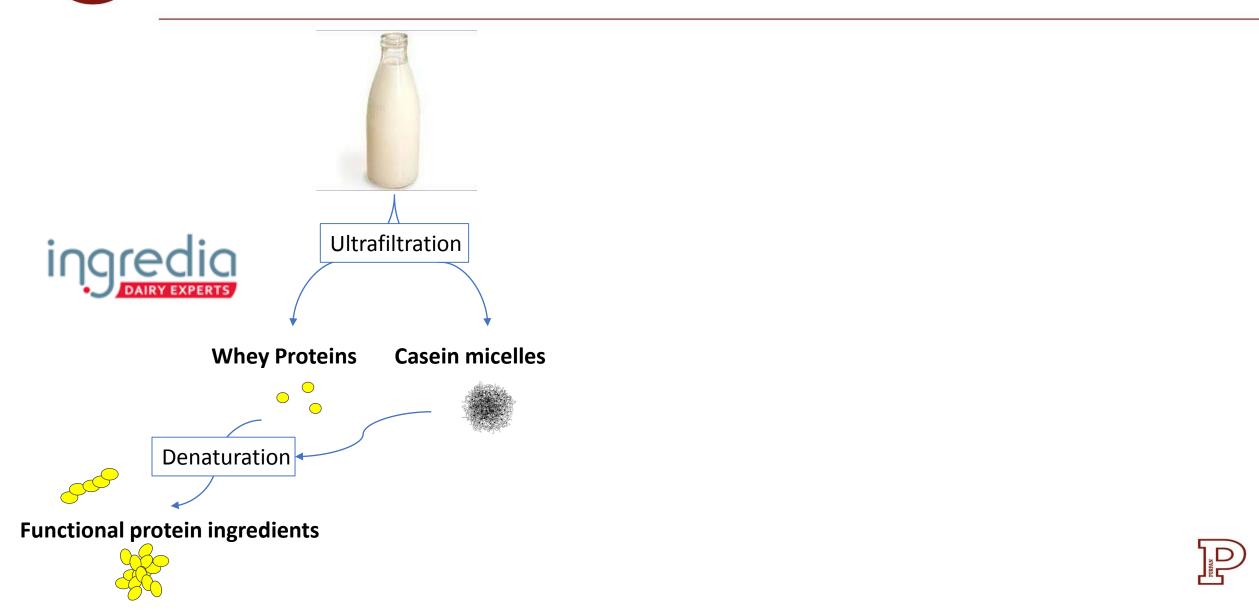


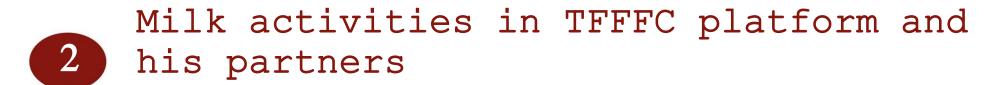


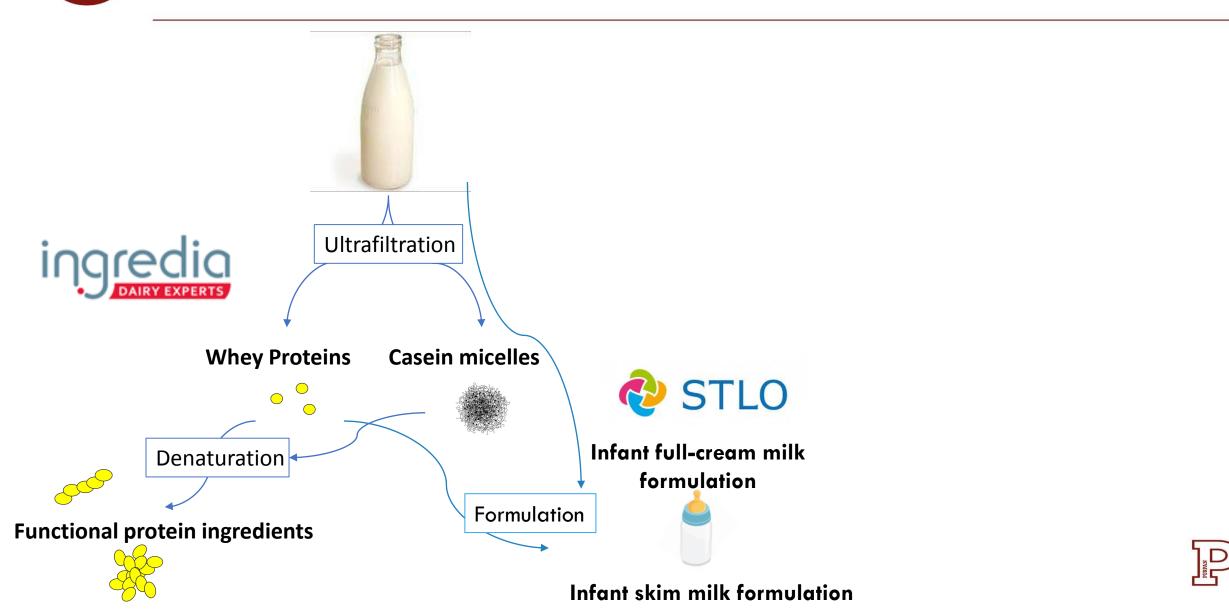


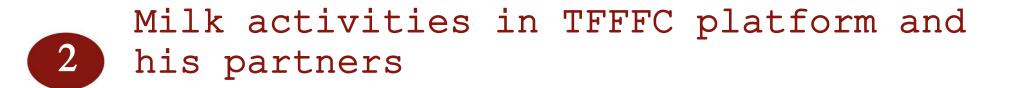




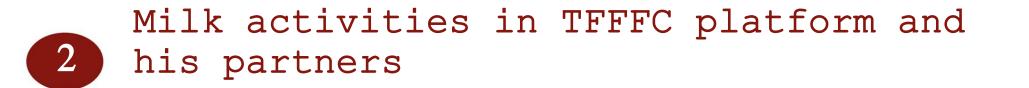






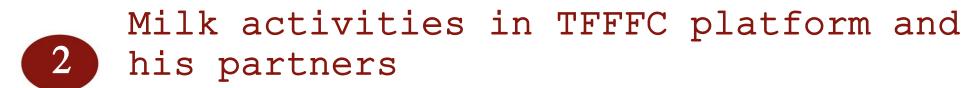










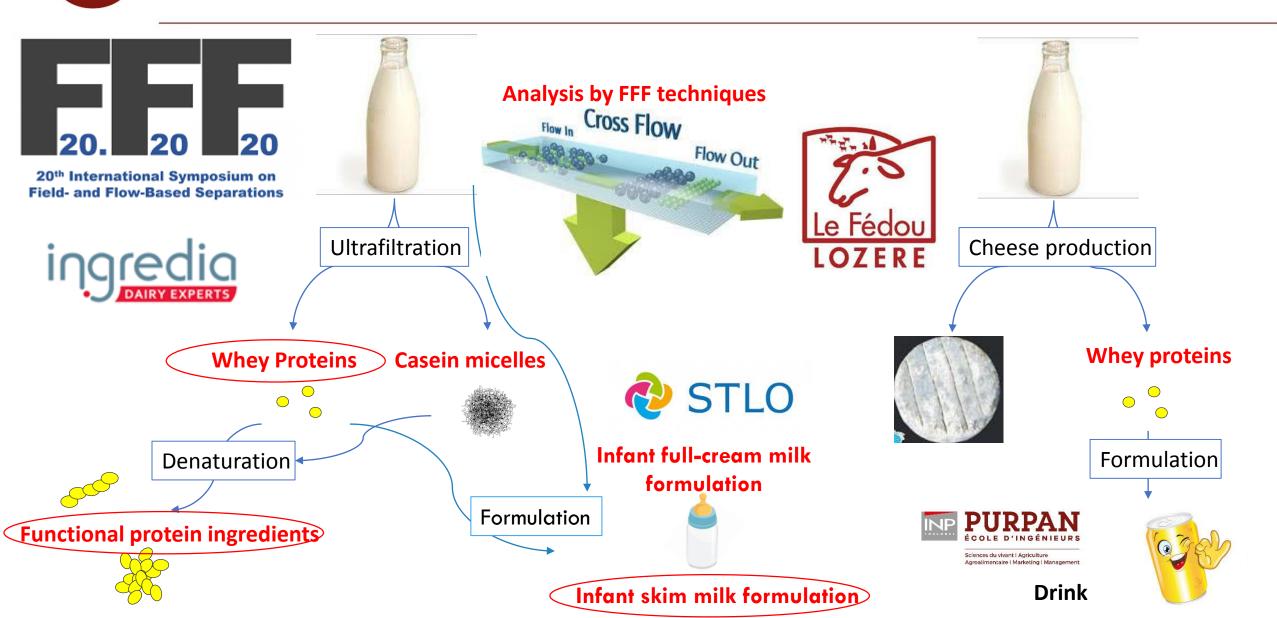




Infant skim milk formulation

Milk activities in TFFFC platform and his partners





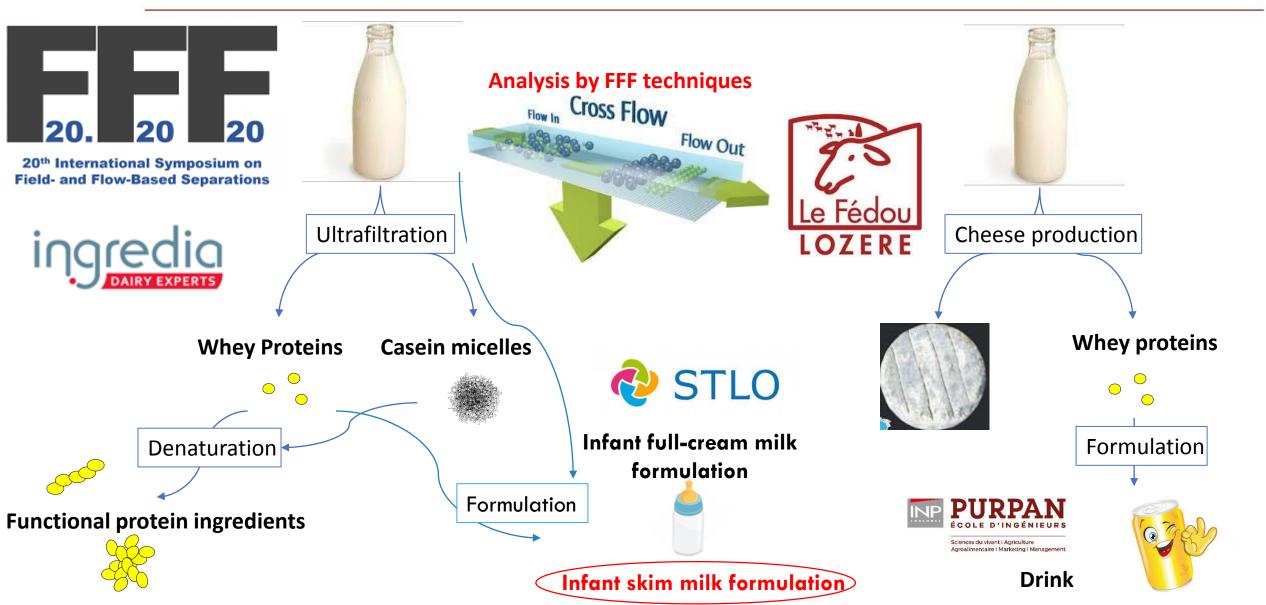


<u>As-FlFFF analysis of Infant Milk</u> <u>Formulation (IMF)</u>

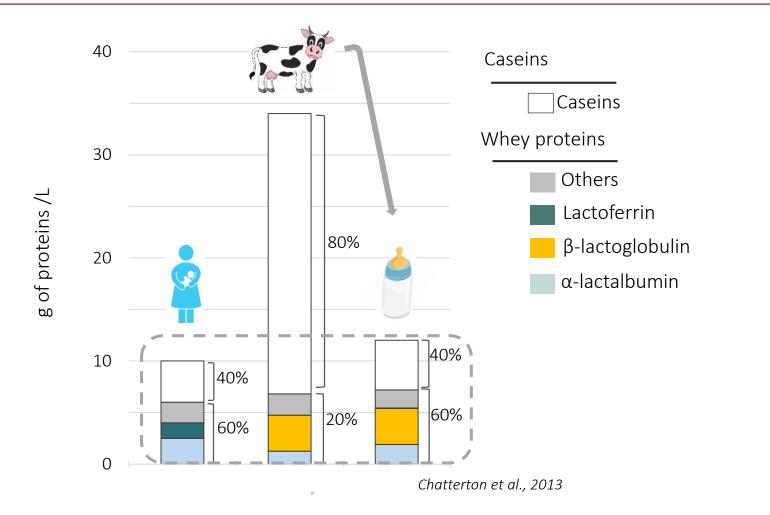




As-FlFFF analysis of Infant Milk Formulation (IMF)

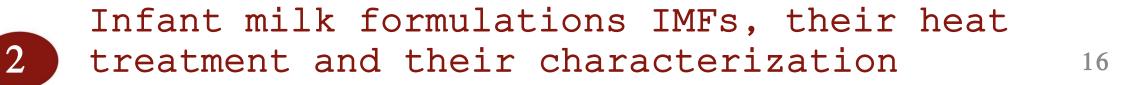


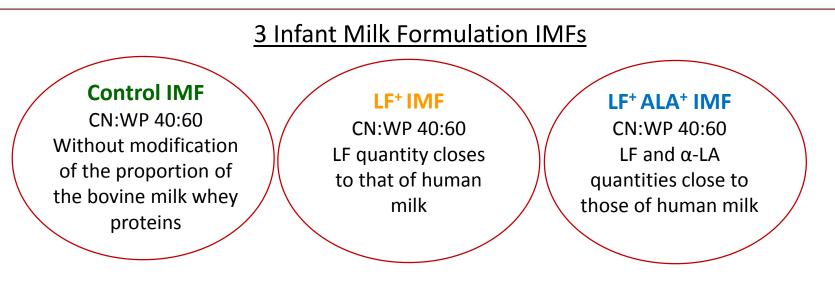




The CN:WP ratio and the WP composition have to be modified for the Infant milk formulation (IMF) to mimic the human milk composition

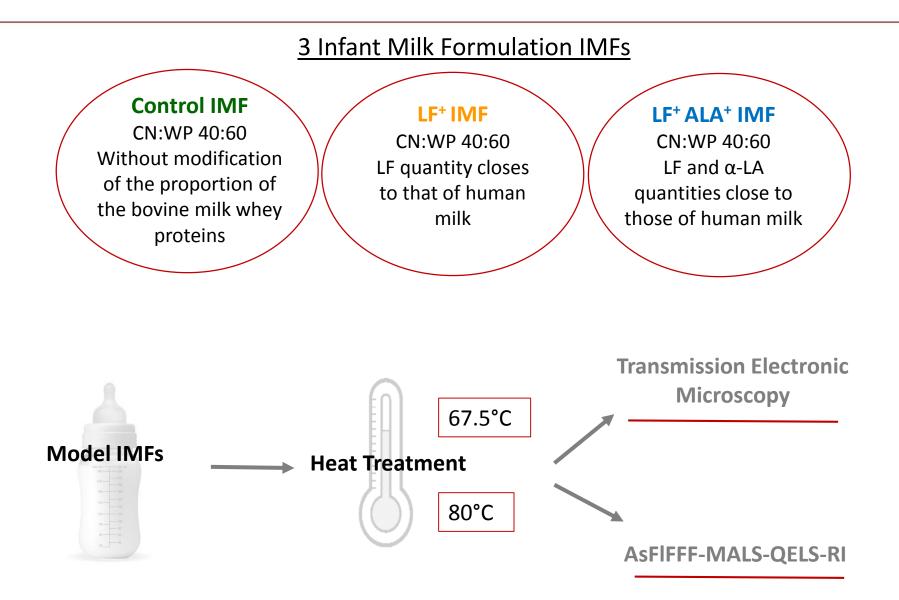






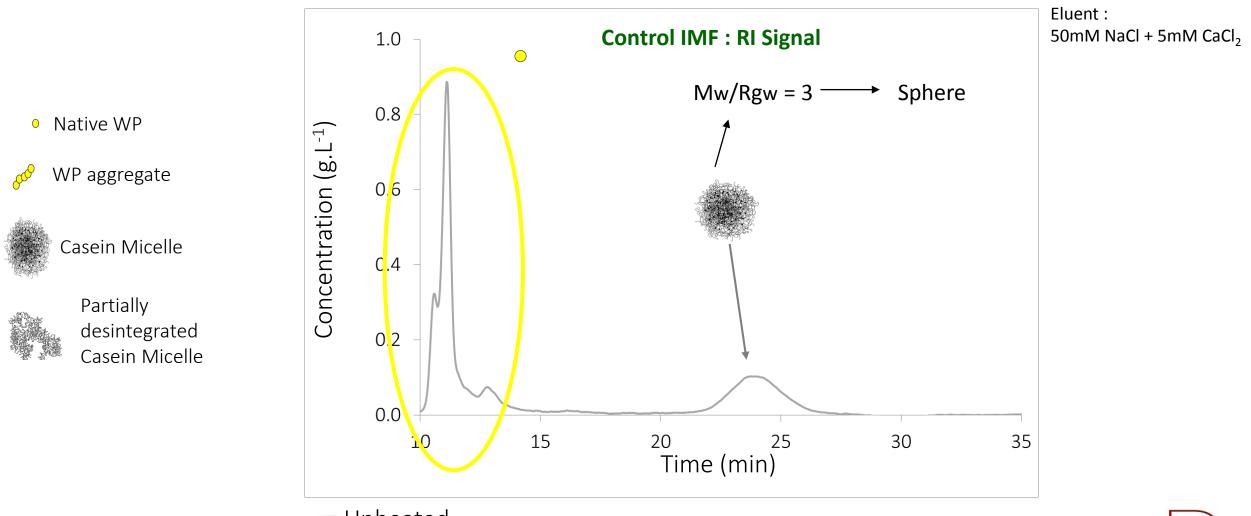


Infant milk formulations IMFs, their heat treatment and their characterization 17



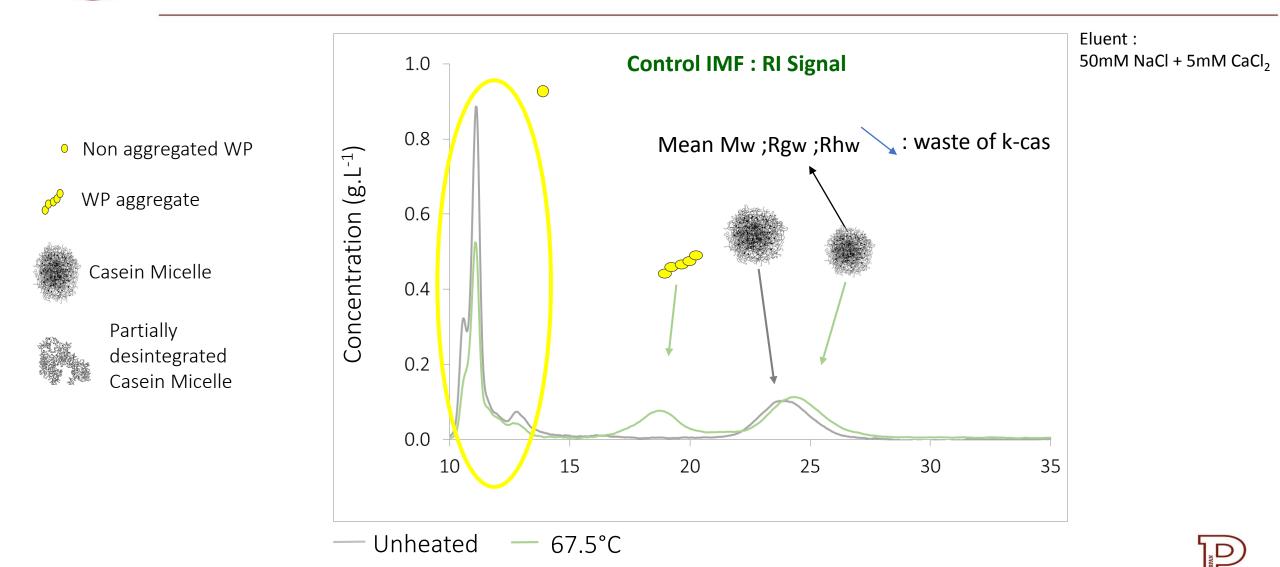


and heated IMFs at 67.5° C and 80° C : Control IMF

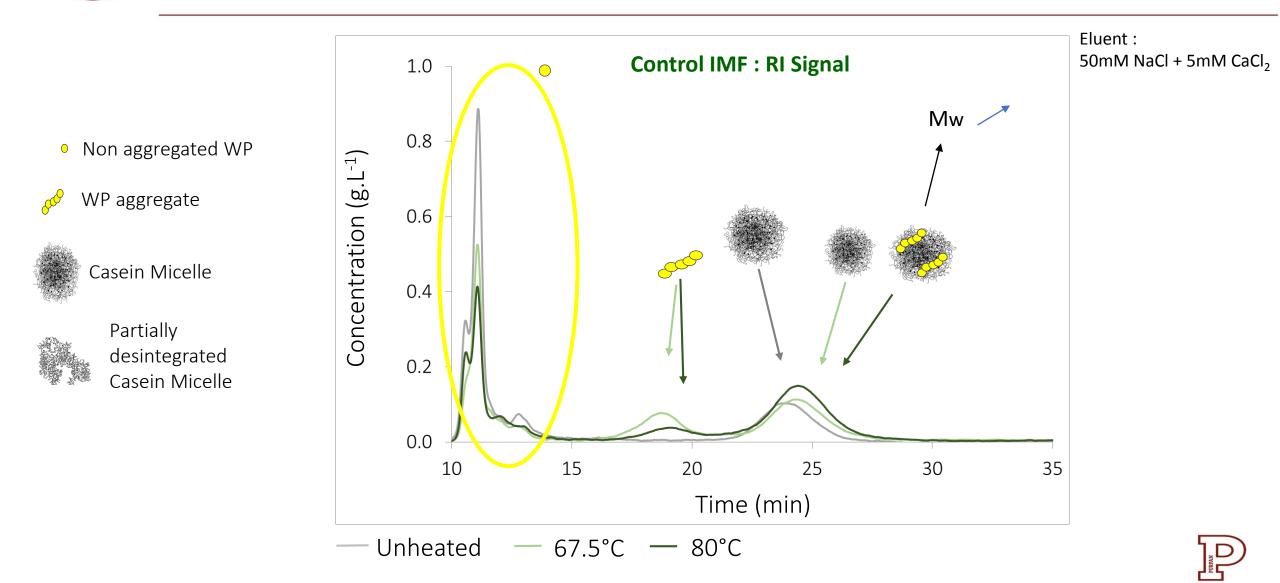


- Unheated

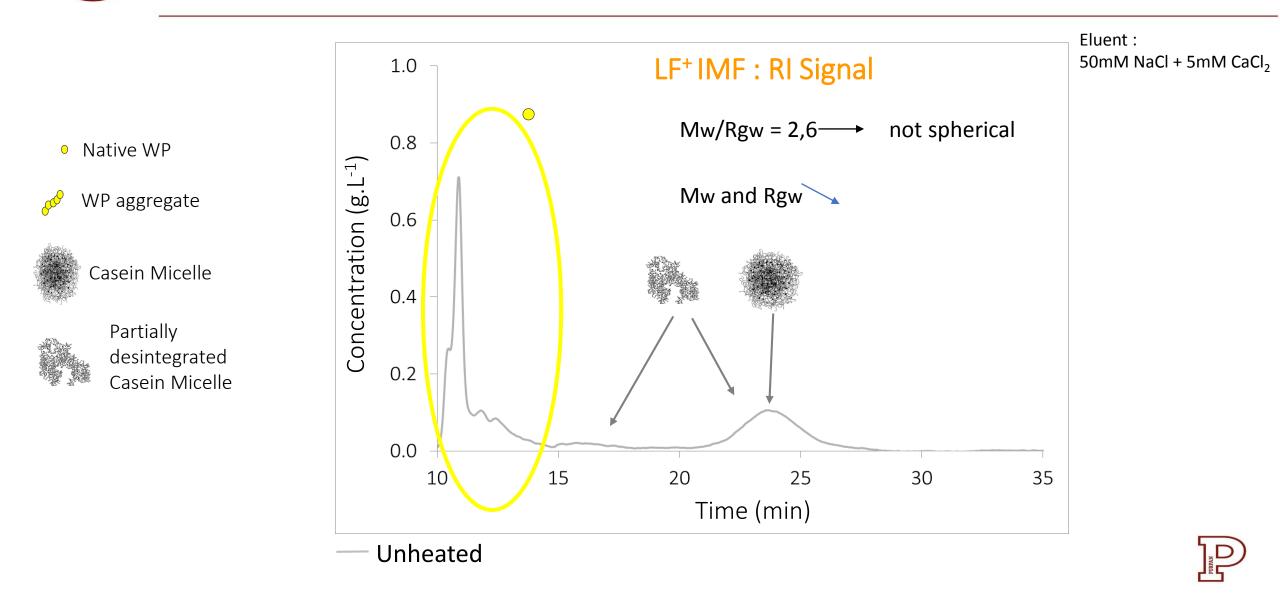
and heated IMFs at 67.5° C and 80° C : Control IMF



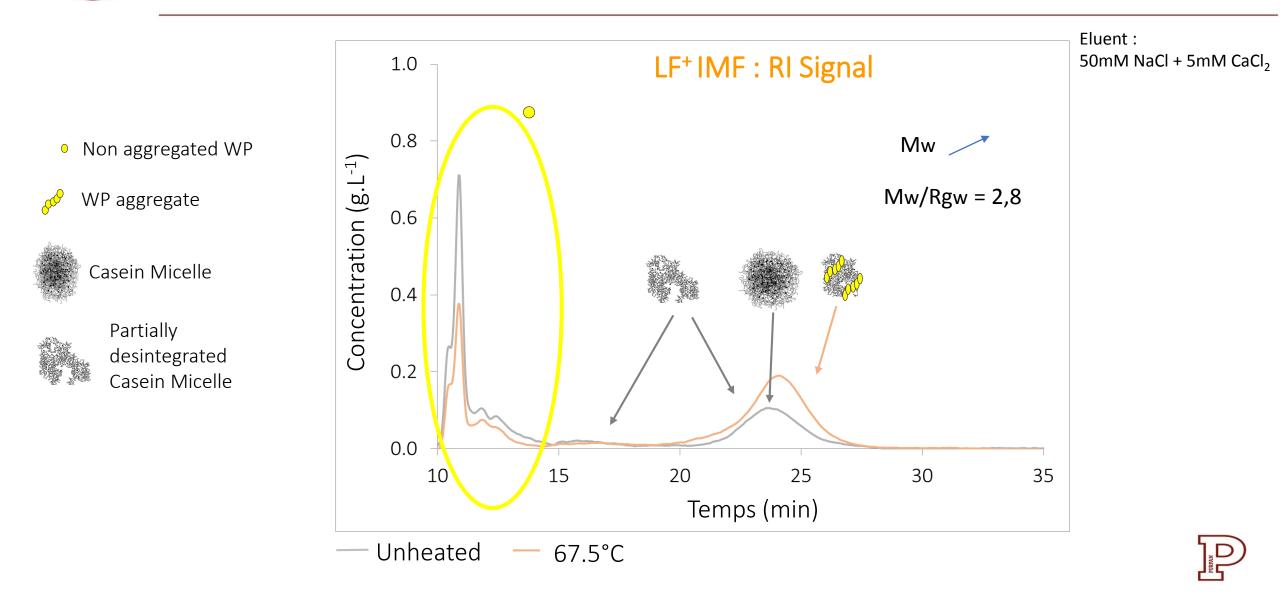
and heated IMFs at 67.5° C and 80° C : Control IMF 20



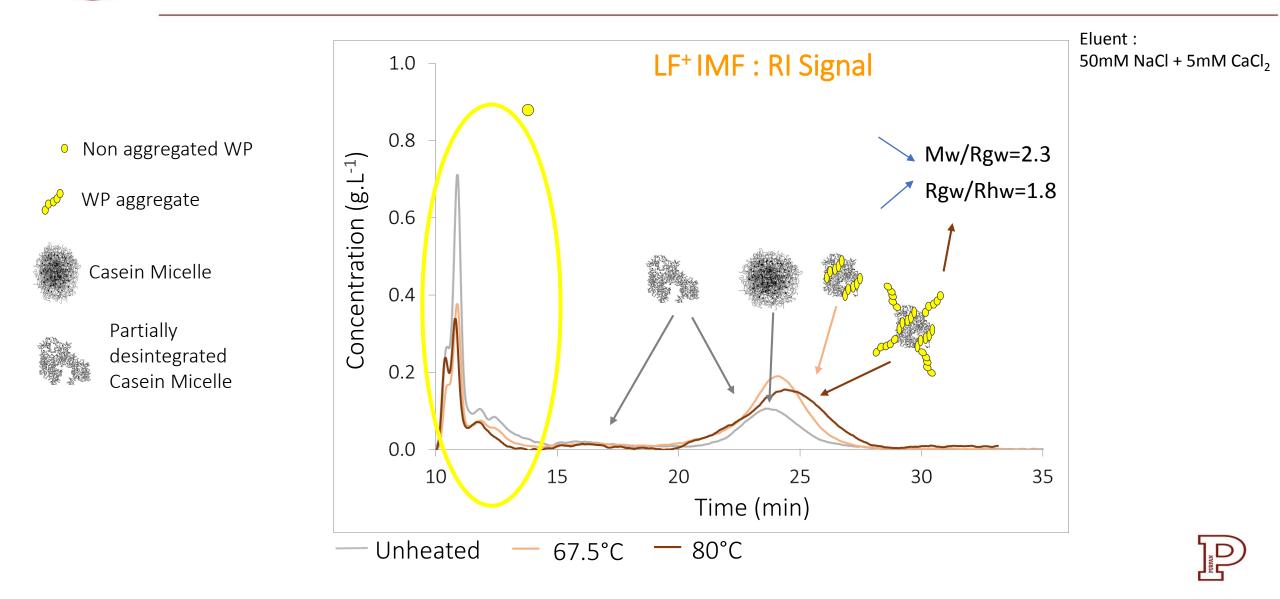
AF4-MALS-QELS-RI fractograms of the unheated and heated IMFs at 67.5° C and 80° C : LF⁺ IMF 21



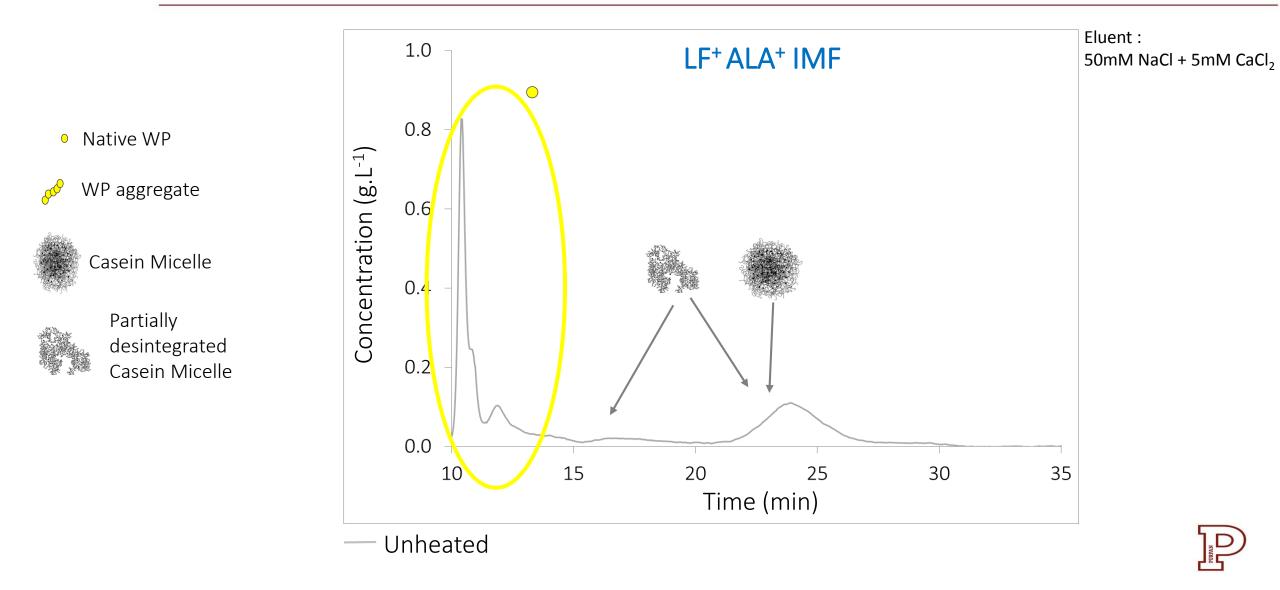
AF4-MALS-QELS-RI fractograms of the unheated and heated IMFs at 67.5° C and 80° C : LF⁺ IMF 22



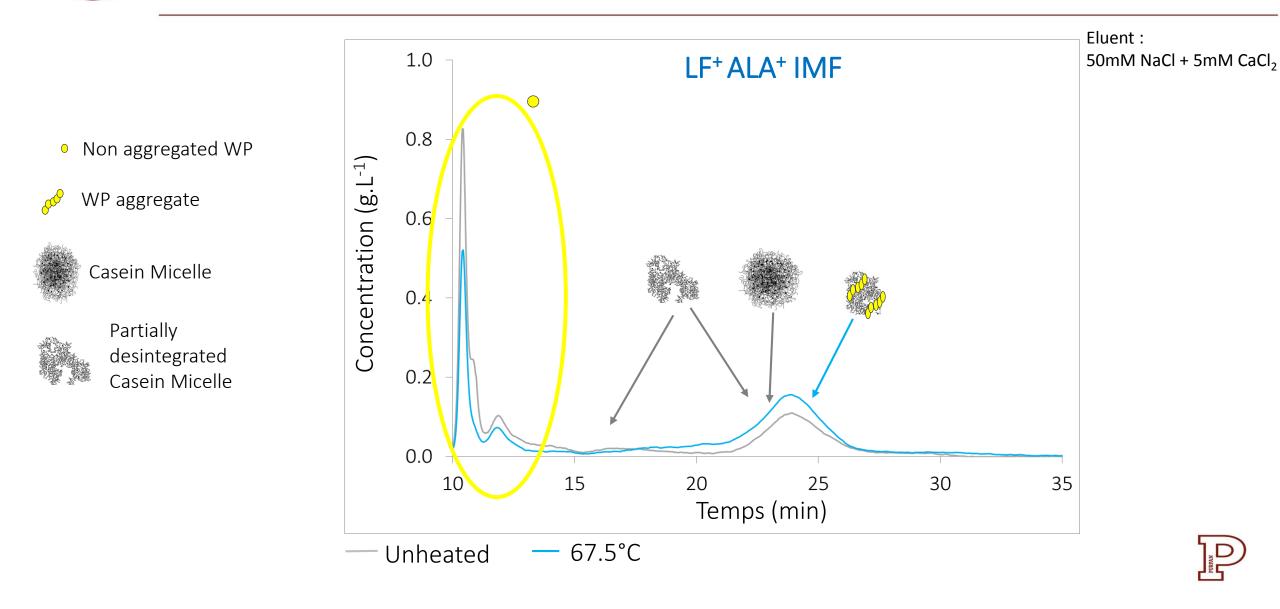
AF4-MALS-QELS-RI fractograms of the unheated and heated IMFs at 67.5° C and 80° C : LF⁺ IMF 23



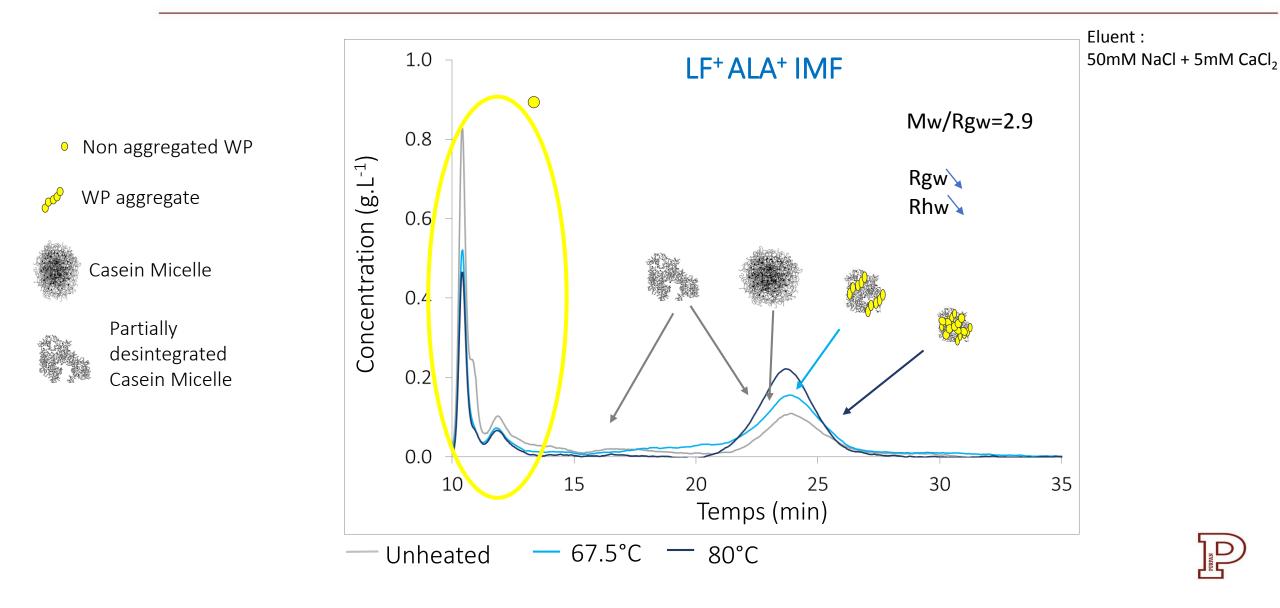
and heated IMFs at 67.5° C and 80° C : LF⁺ ALA⁺ IMF 24



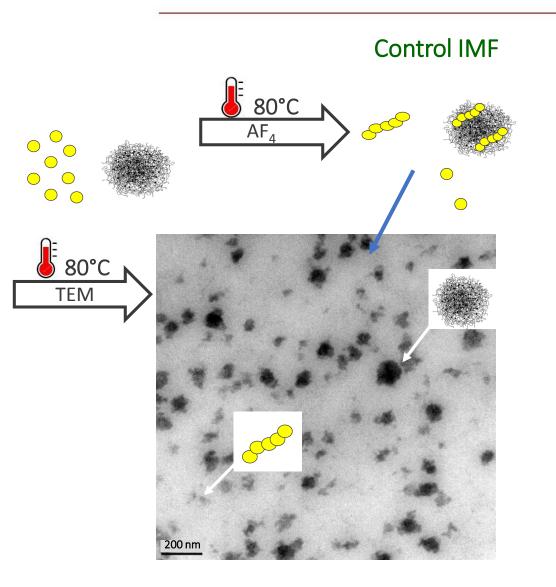
and heated IMFs at 67.5° C and 80° C : LF⁺ ALA⁺ IMF 25



and heated IMFs at 67.5° C and 80° C : LF⁺ ALA⁺ IMF 26



5 Microscopy



Non aggregated WP

⁹ WP aggregate



0

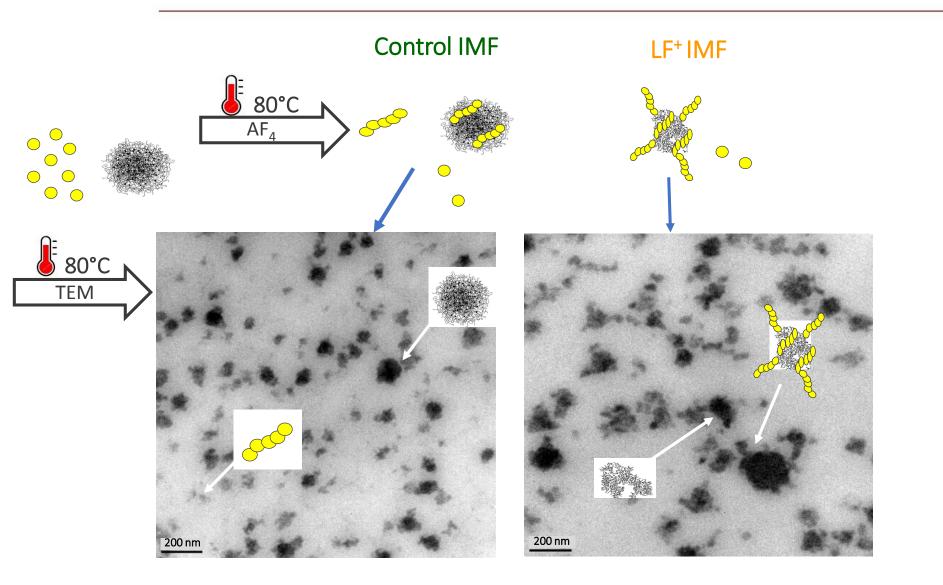
Casein Micelle

Par des

Partially desintegrated Casein Micelle



5 Microscopy



Non aggregated WP

⁹ WP aggregate

C C

Casein Micelle

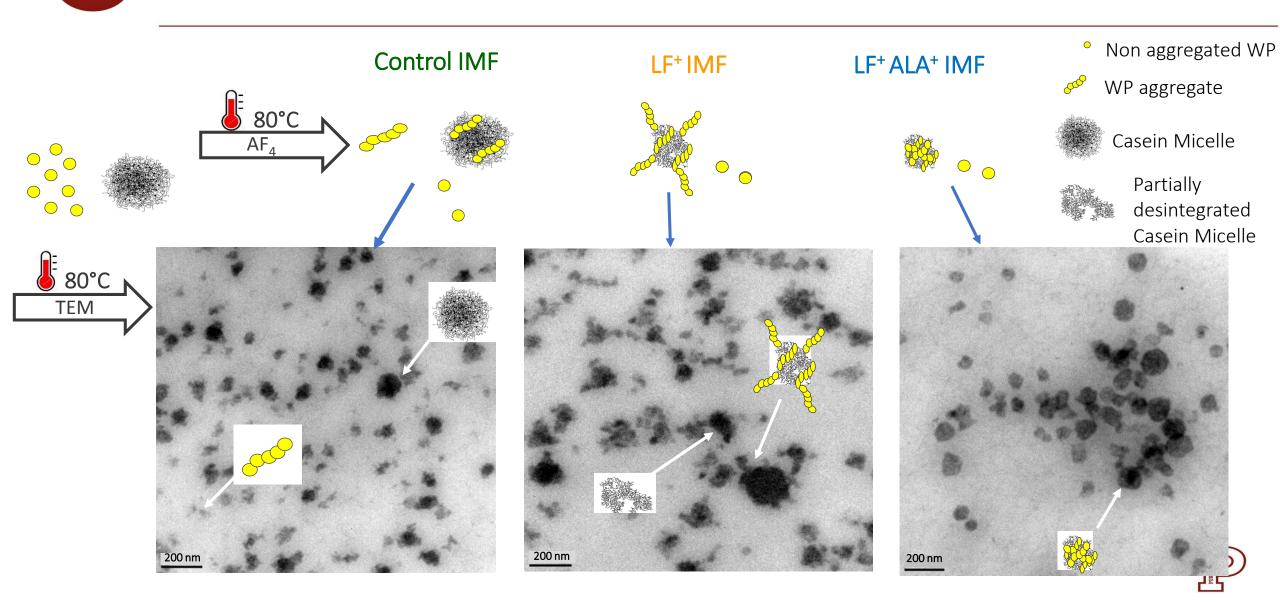
P d

Partially desintegrated Casein Micelle



5 Microscopy



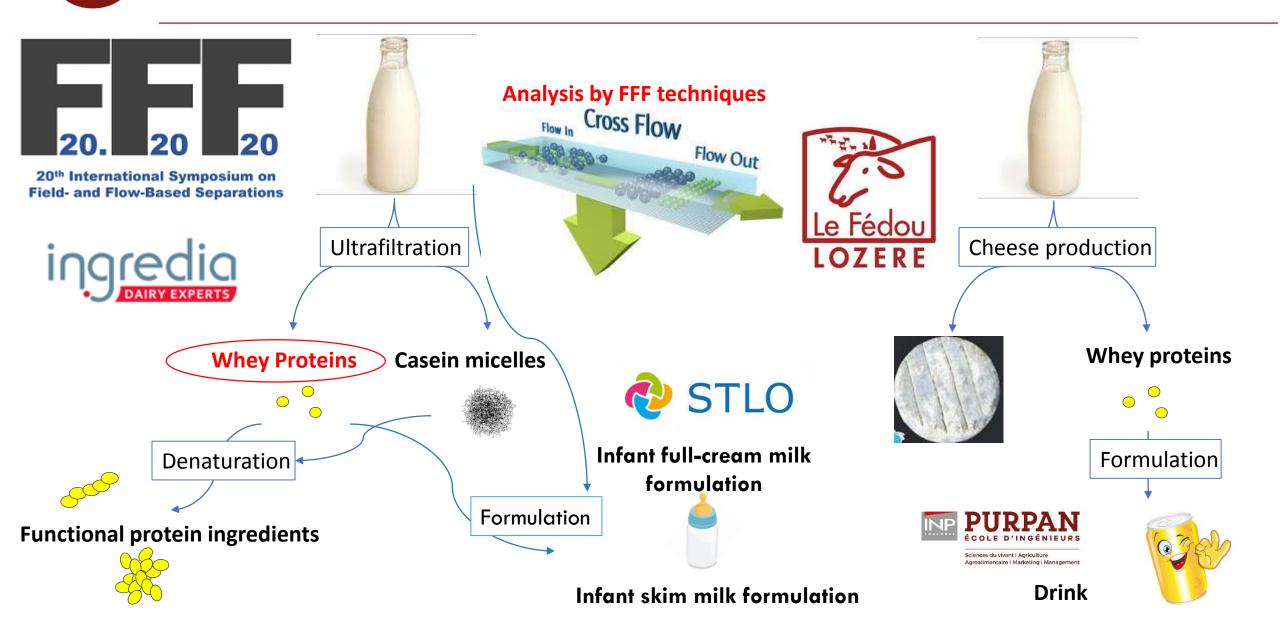




Study of Whey proteins by 2D AsFlFFF-RPLC



Study of Whey proteins by 2D-AsFlFFF-RPHPLC



Whey Proteins Analysis by **1** As-FlFFF (1D)

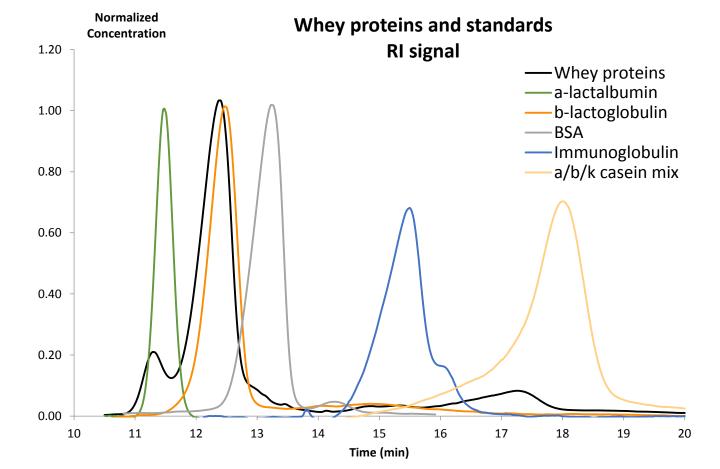
32

Industrial whey protein isolate : they're few percents of caseins inside Objective : separation of all proteins



Whey Proteins Analysis by 1 As-FlFFF (1D)

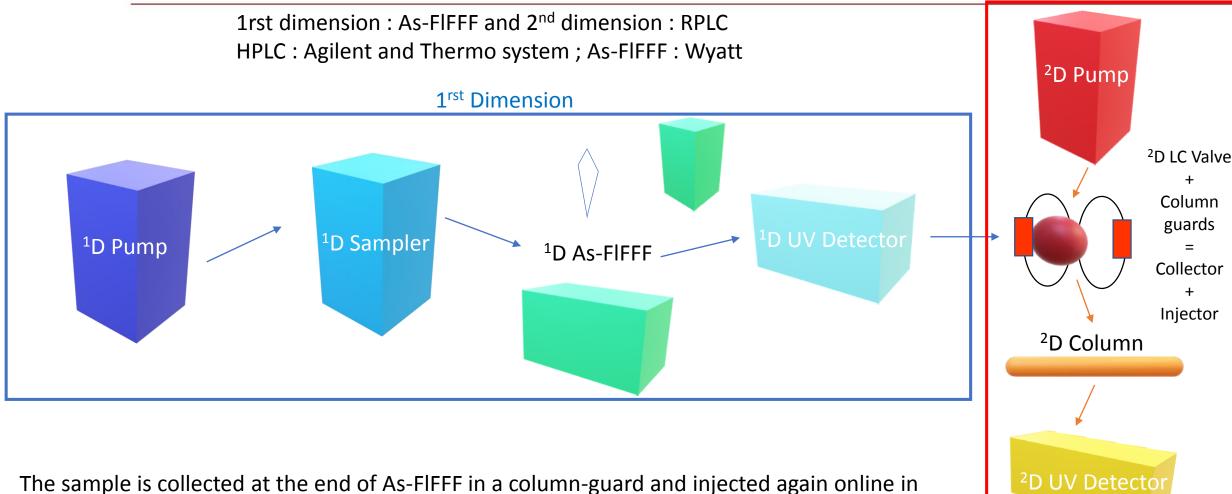
Industrial whey protein isolate : they're few percents of caseins inside Objective : separation of all proteins



- The different whey proteins are identified but not very good separated by AsFIFFF
- By HPLC it's not possible to separate caseins from whey proteins



Whey Proteins Analysis by 2D 2 AsFlFFF-RPLC



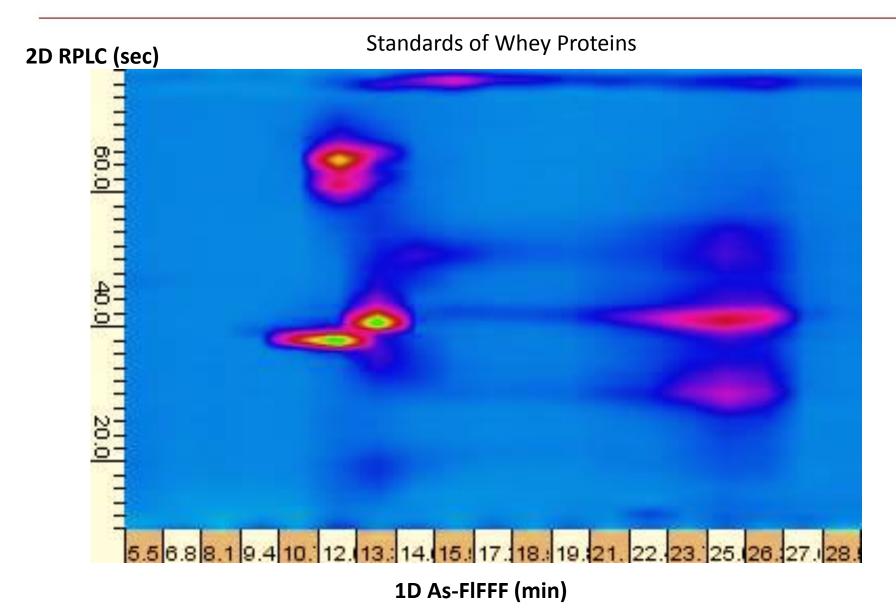
The sample is collected at the end of As-FIFFF in a column-guard and injected again online in the column every 80 seconds

34

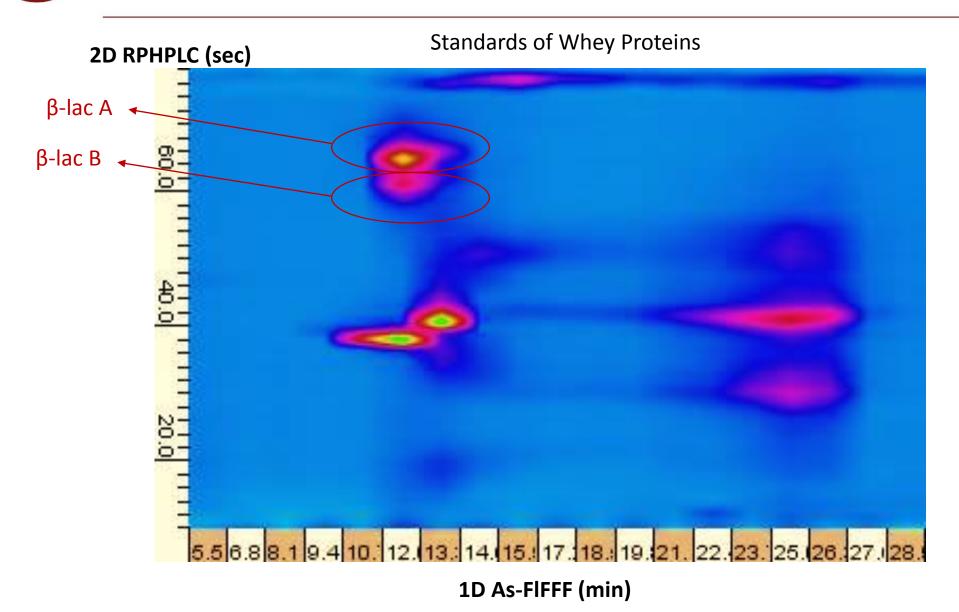
Dimension

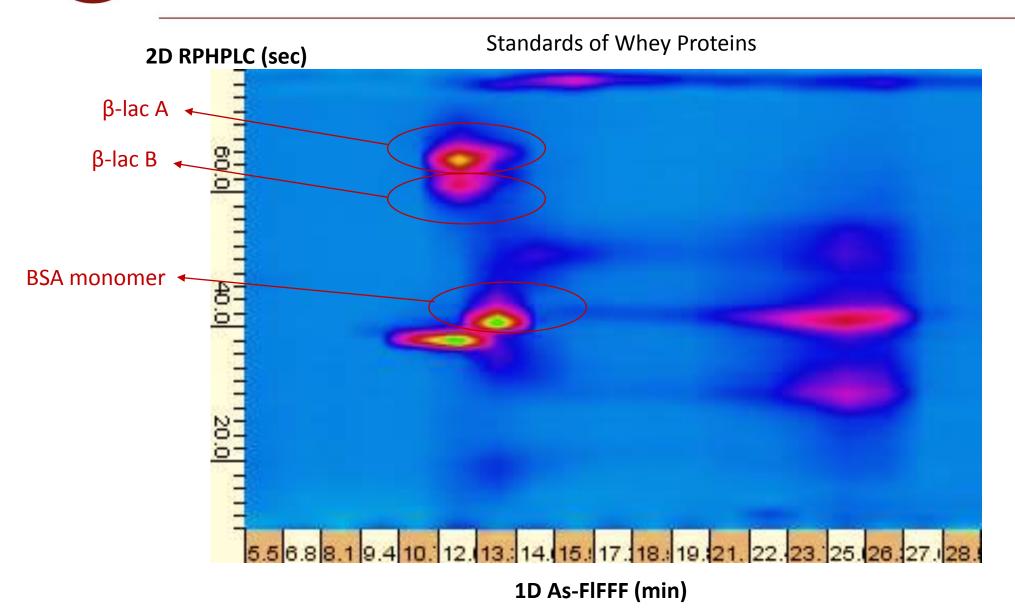
2nd

Whey Proteins Analysis by 2D AsFlFFF-RPLC : UV signal mapping $_{35}$

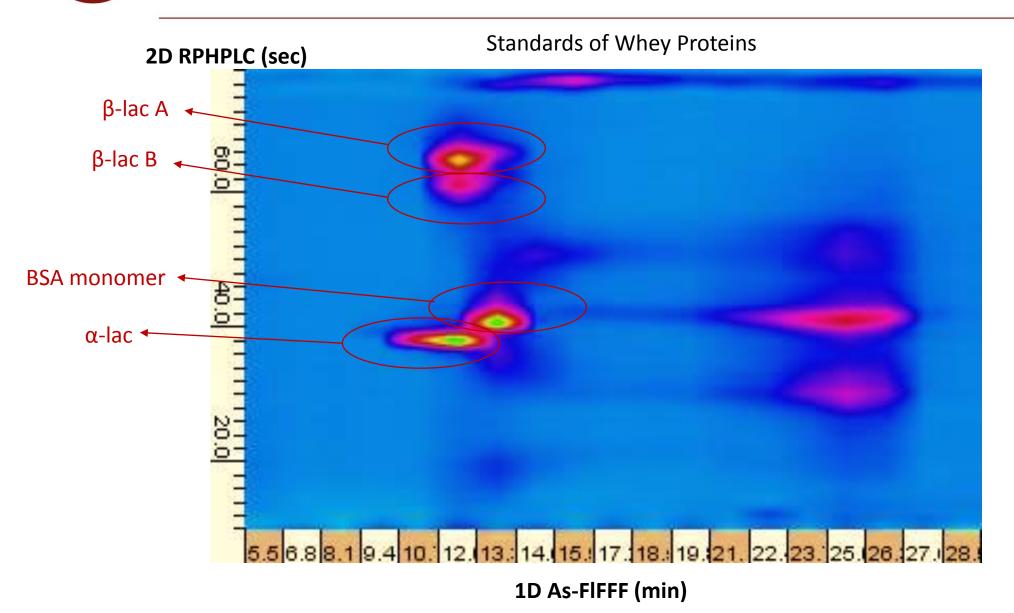


Whey Proteins Analysis by 2D-AsFlFFF-RPHPLC : UV signal mapping

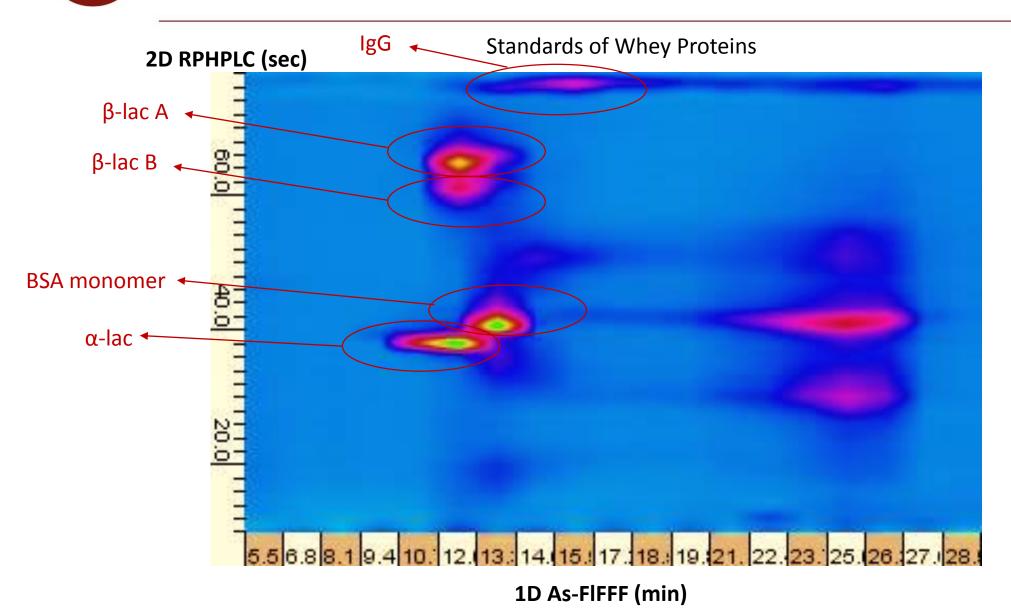




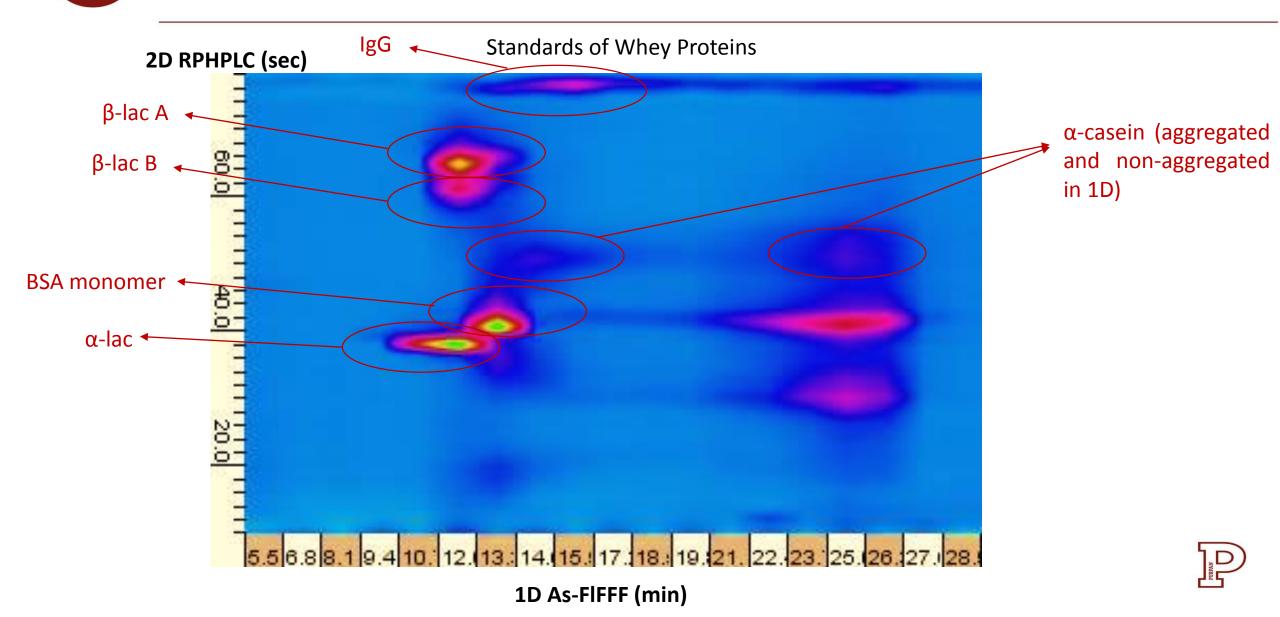


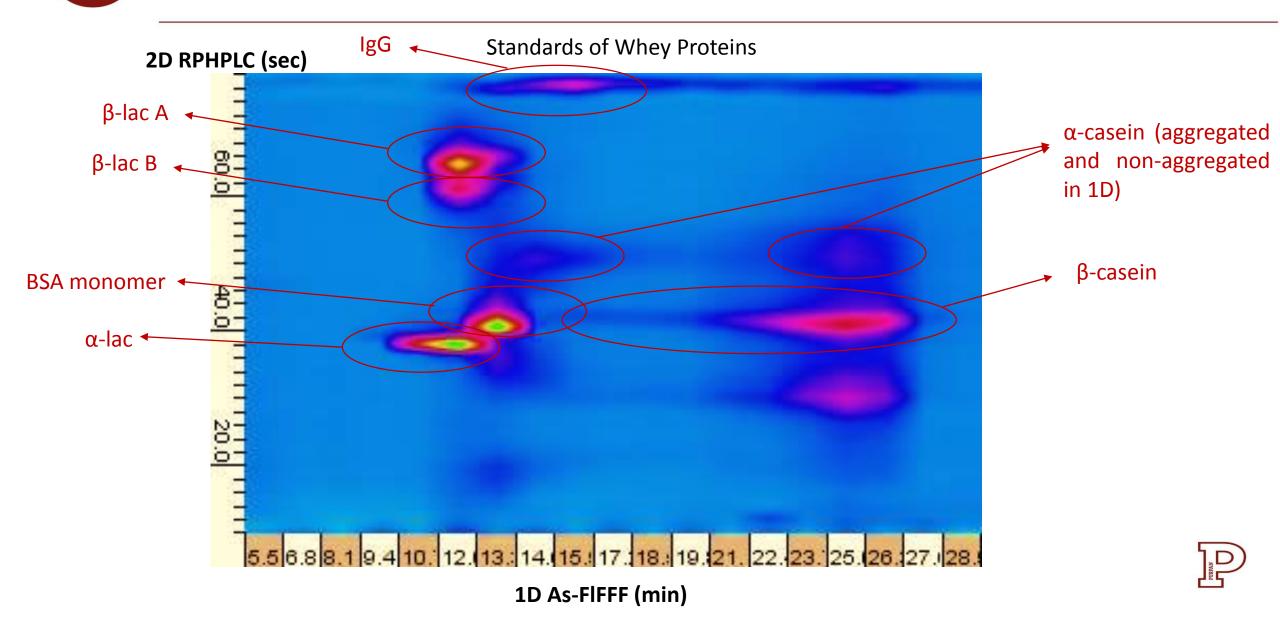


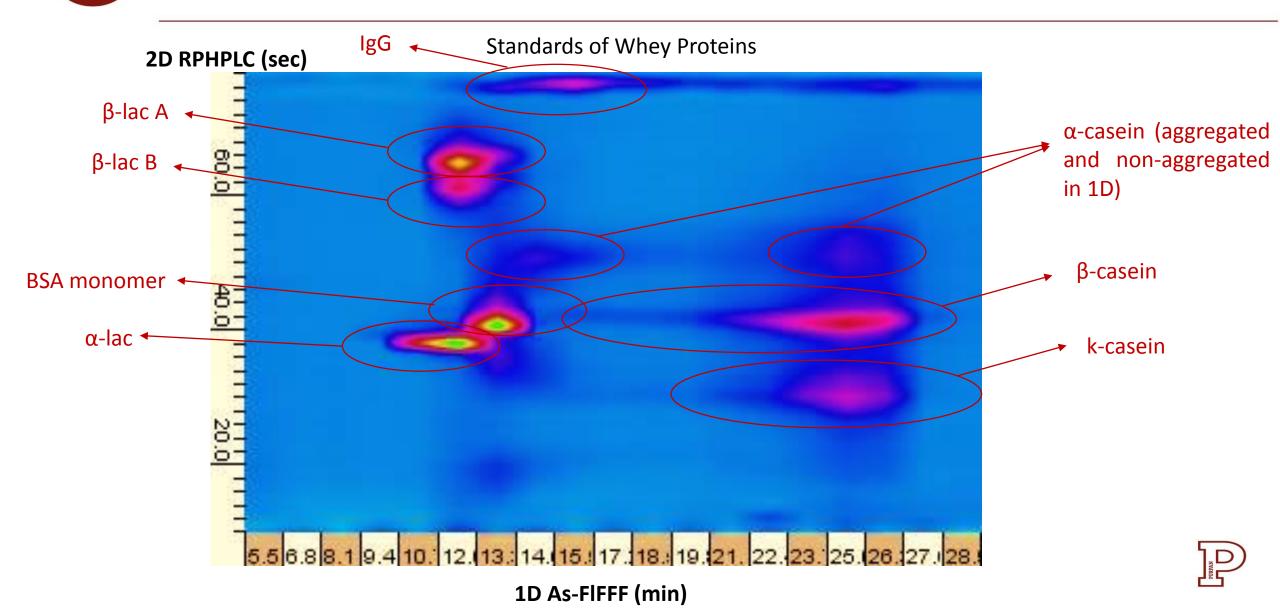














<u>Denatured whey proteins studied by</u> <u>Centrifugal-FFF</u>

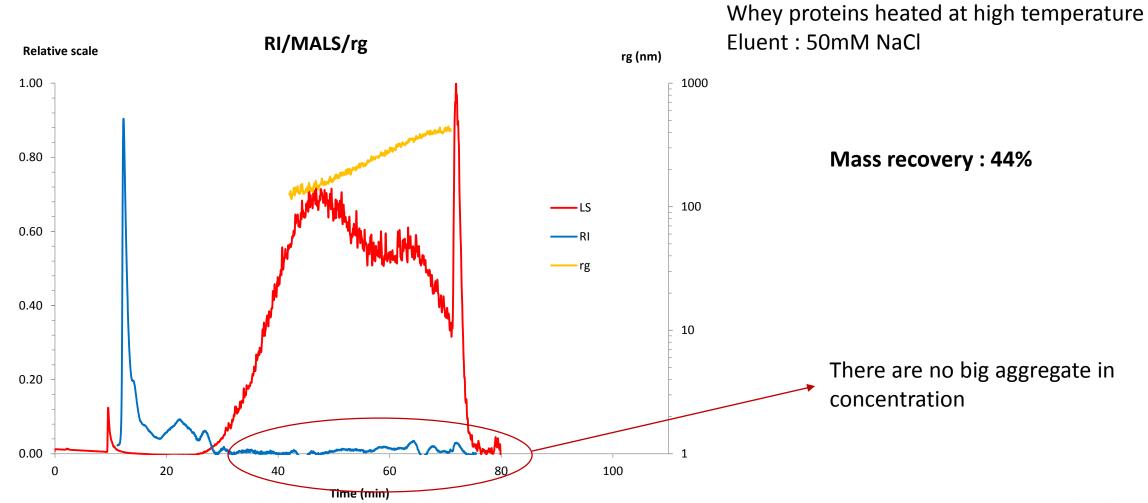






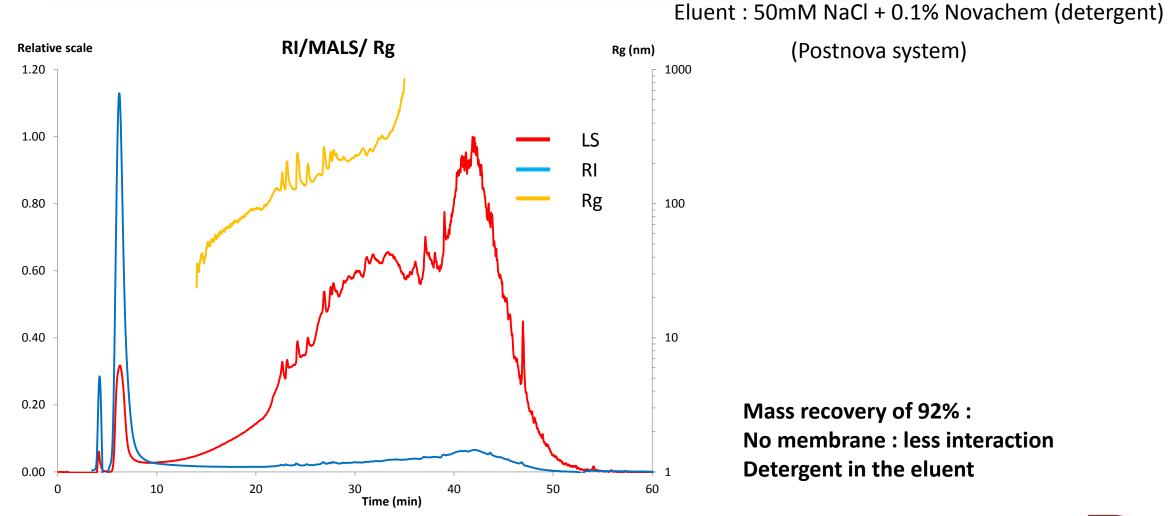


Denatured whey proteins studied by As-1 FIFFF





Denatured whey proteins studied by Centrifugal-FFF





Denatured whey proteins studied by Centrifugal-FFF

RI/MALS/ Rg

20

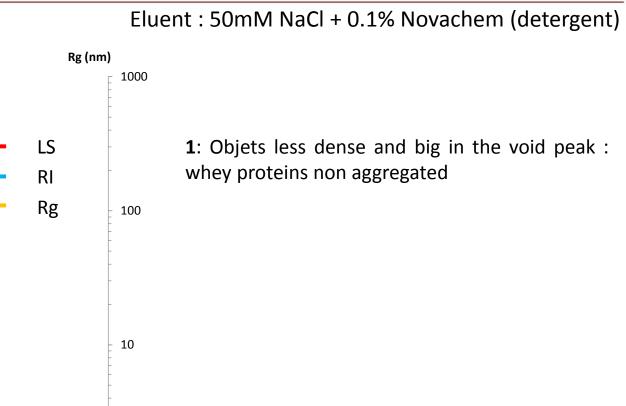
30

Time (min)

40

50

60



Mass recovery of 92% : No membrane : less interaction Detergent in the eluent

2

Relative scale

1

10

1.20

1.00

0.80

0.60

0.40

0.20

0.00

0



Denatured whey proteins studied by Centrifugal-FFF

Rg (nm)

10

60

LS

RI

Rg

50

40

RI/MALS/ Rg

2

1000 1: Objets less dense and big in the void peak 2: Rg from 25 to 850nm 100 Mass recovery of 92% : No membrane : less interaction

Eluent : 50mM NaCl + 0.1% Novachem (detergent)

Detergent in the eluent



Mass fractions 51%

10

20

30

23%

Time (min)

2

Relative scale

1.20

1.00

0.80

0.60

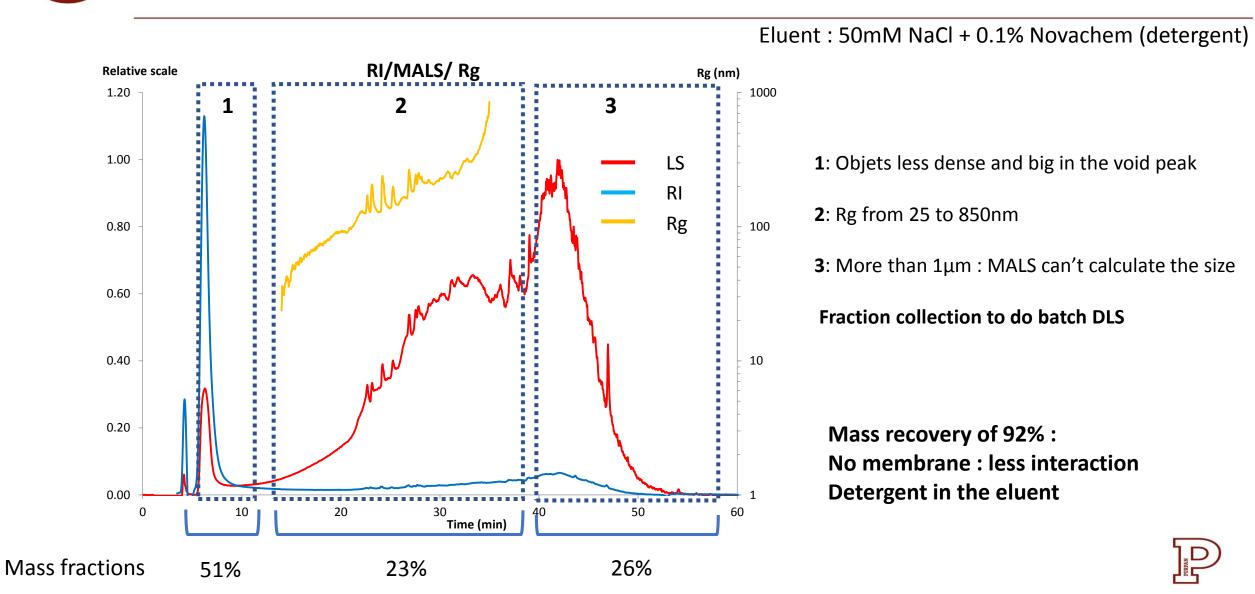
0.40

0.20

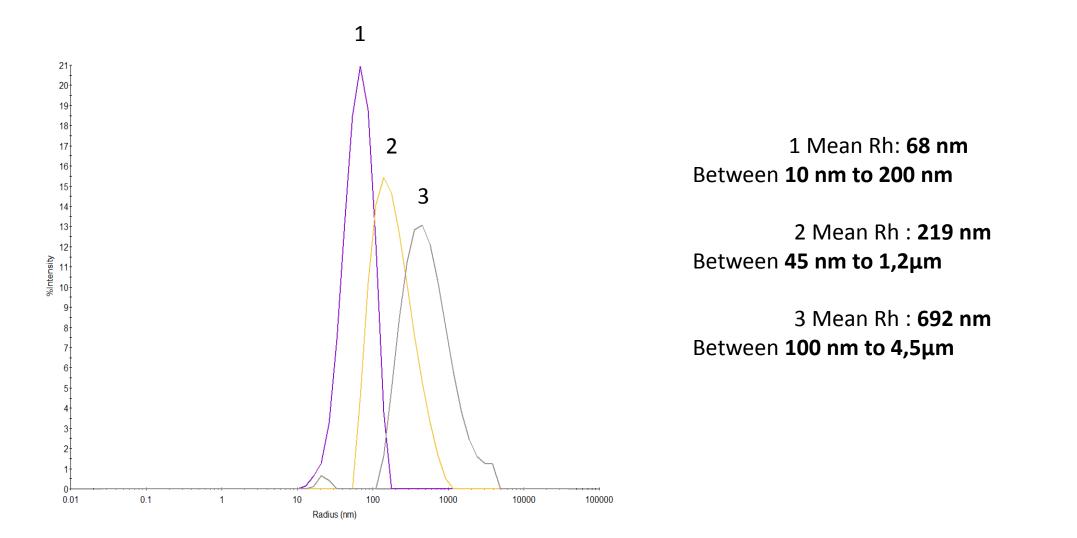
0.00

0

Denatured whey proteins studied by Centrifugal-FFF









Conclusion and outlook

• 1^{rst} study : casein-WP pasteurized structures are different in function of the WP composition in IMF and heat temperature :

- Addition of lactoferrin in IMF destabilize the casein micelles

– Addition of Lactoferrin form sturctures with ams around the micelle after heat treatment at $80\,^\circ\text{C}$

– Addition of Lactoferrin + α -lac form smaller structures with WP aggregates inside the micelles at $80\,^\circ\text{C}$

Digestibility of this structures? Scale-up for formulation and pasteurization : Stuctural difference?



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Quantitative experiment with calibration curve and industrial mix study

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for

 3rd study : Centrifugal-FFF allow to have a better mass recover denatured WP

Acknowledgmen

Amira Halabi, Amélie Deglaire, Thomas Croguennec, Audrey Romelard, Alain Baniel, Frédéric Violleau

<u>The TFFFC team</u>: Estelle N'Tsiba, Audrey Ric, Mireille Gaucher, Kévin Pascotto







Projet cofinancé par le Fonds Européen de Développement Régional





Thank you for your attention!

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TFFFC

Annexes

IMFs	Heat treatment	Elution time ranges	$\overline{M_w}$	$\overline{R_g}$	$\overline{R_h}$	Shape	df_{app}
	parameters	(min)	(10 ⁸ g.mol ⁻¹)	(nm)	(nm)	factors	
Control	Unheated	20-28	6.4	138	96	1.4	3.0
	67.5℃	22-28	5.8	121	80	1.5	2.6
	80°C	22-28	6.8	126	82	1.5	2.7
LF+	Unheated	20-28	4.7	123	91	1.4	2.6
	67.5°C	20-28	6.0	128	85	1.5	2.8
	80°C	20-28	7.3	157	87	1.8	2.3
LF+α-LA+	Unheated	20-28	4.7	142	91	1.6	2.5
	67.5°C	20-28	6.0	153	82	1.9	2.7
	80°C	20-28	6.2	123	78	1.6	2.9