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Asymmetrical Flow Field-Flow Fractionation coupled with Multi-Angle Light Scattering and Differential Refractometer (AF4-MALS-DRi): An analysis tool to characterize milk protein aggregates produced in semi-industrial way

**M.Hennetier^{1*}, D.De-Guibert², Y.Gu³, V.Solé¹, C.Garnier¹, G.Delaplace³,
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Context



**Thickening
or gelling
agents**



Introduction

Methods

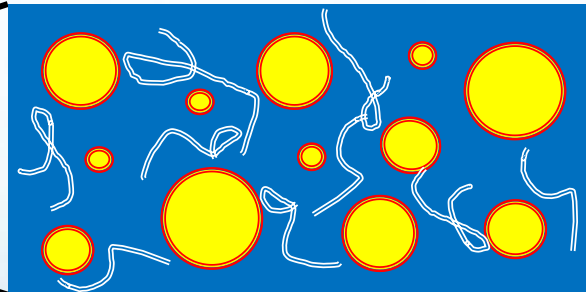
Results

Conclusion and
perspectives

Context



Thickening
or gelling
agents



Oil/Water emulsion

Gelling agents texture
the continuous phase
and entrap the oil
droplets

Introduction

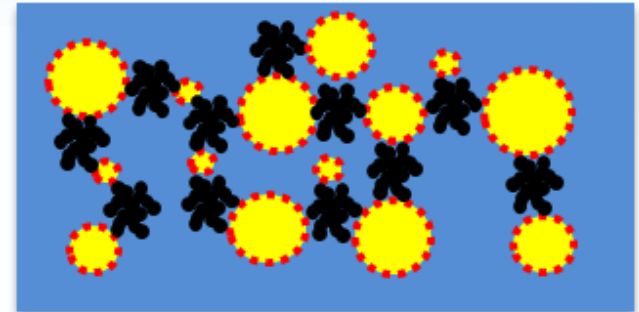
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Whey protein aggregates properties

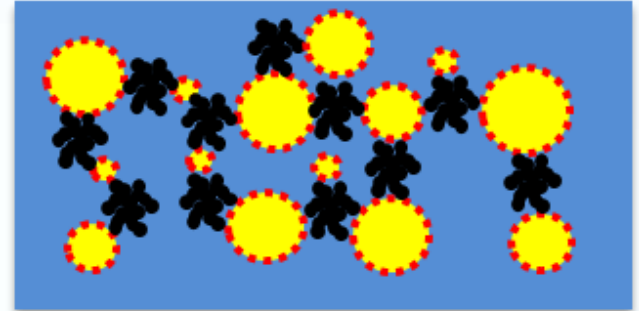
- ✓ Use oil droplets to texture emulsion without gelling agents
- ✓ Connect oil droplets through whey protein aggregates
- ✓ Heating can form aggregates with different shapes and sizes (Nicolai, 2011)



C. Surel, J. Fouquier, N. Perrot, A. Mackie, C. Garnier, A. Riaublanc, and M. Anton, "Composition and structure of interface impacts texture of O/W emulsions," *Food Hydrocoll.*, vol. 34, pp. 3–9, 2014.

Whey protein aggregates properties

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Applications

Replace food additives like thickening or gelling agents in neutral dairy products

Texture emulsions even at low fat content



Fractal aggregates production from Whey Protein Isolate (WPI) at semi-industrial scale

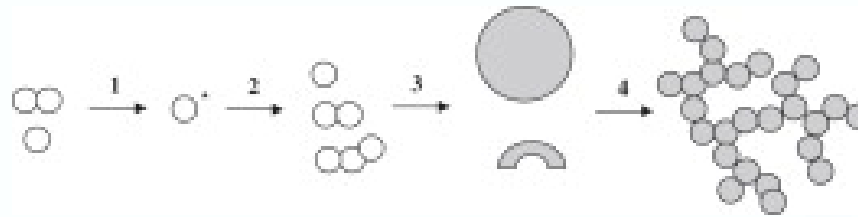
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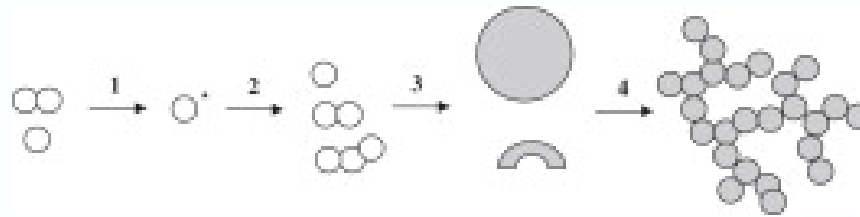
Fractal aggregate formation



(Nicolai, 2011)

- 1: Dimer/trimer of whey proteins (β -lactoglobulin) are heated at 80°C
→ Denaturation, unfolding and thiol/hydrophobic group exposition

Fractal aggregate formation

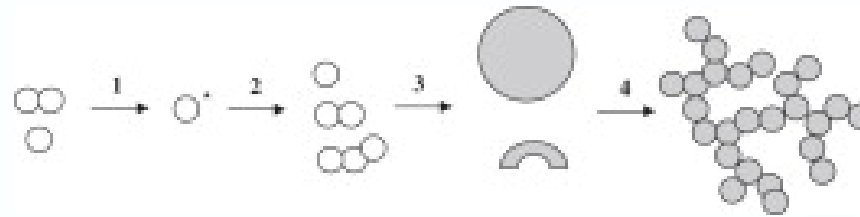


(Nicolai, 2011)

1: Dimer/trimer of whey proteins (β -lactoglobulin) are heated at 80°C
→ Denaturation, unfolding and thiol/hydrophobic group exposition

2: Denatured proteins form oligomers (irreversible)

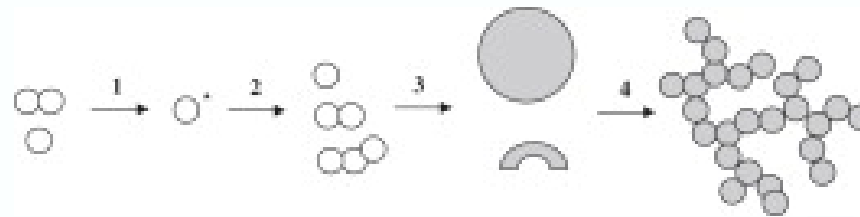
Fractal aggregate formation



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- 3: Critical concentration in oligomer : Primary aggregates : curved if pH=7

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→ Denaturation, unfolding and thiol/hydrophobic group exposition
- 2: Denatured proteins form oligomers (irreversible)
- 3: Critical concentration in oligomer : Primary aggregates : curved if pH=7
- 4: When Salt concentration is sufficient: Branched aggregate formation (disulfides bonds, hydrogen and hydrophobic interactions)

Objective:

Characterization of aggregates produced at semi-industrial scale

Dynamic light scattering (DLS) in batch



- Large objects are exacerbated
- No differentiation in large particle populations

Size exclusion chromatography (SEC)



- Aggregates obstruct the column

Objective:

Characterization of aggregates produced at semi-industrial scale

Dynamic light scattering (DLS) in batch



As-FIFFF

- Large objects are exacerbated
- No differentiation in large particle populations

- Injection of entire product without filtration
- Fractionation of large particles



Size exclusion chromatography (SEC)



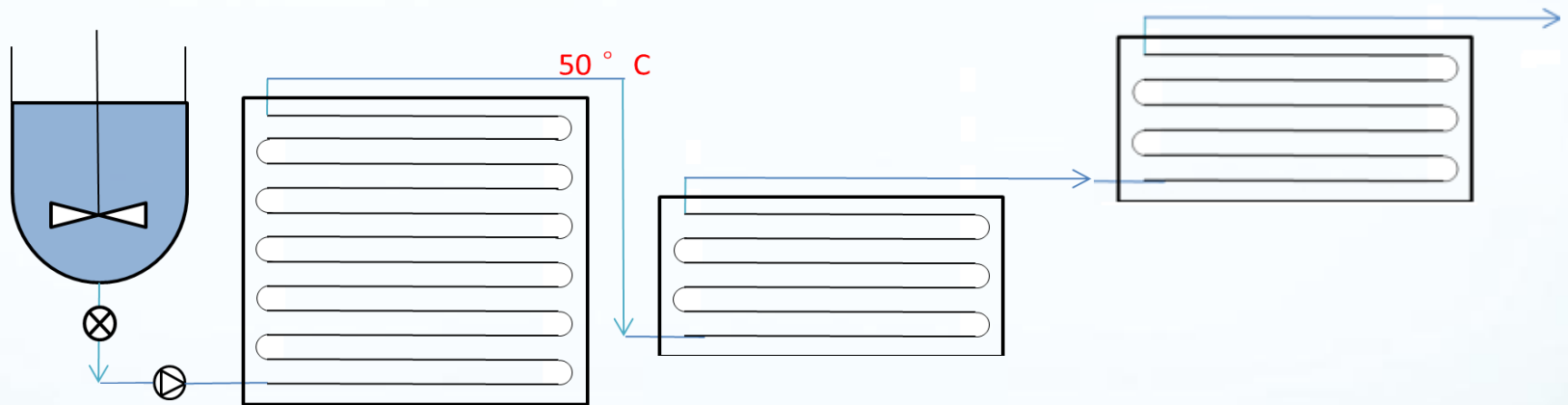
Transmission Electronic Microscopy (TEM)

- Aggregates obstruct the column

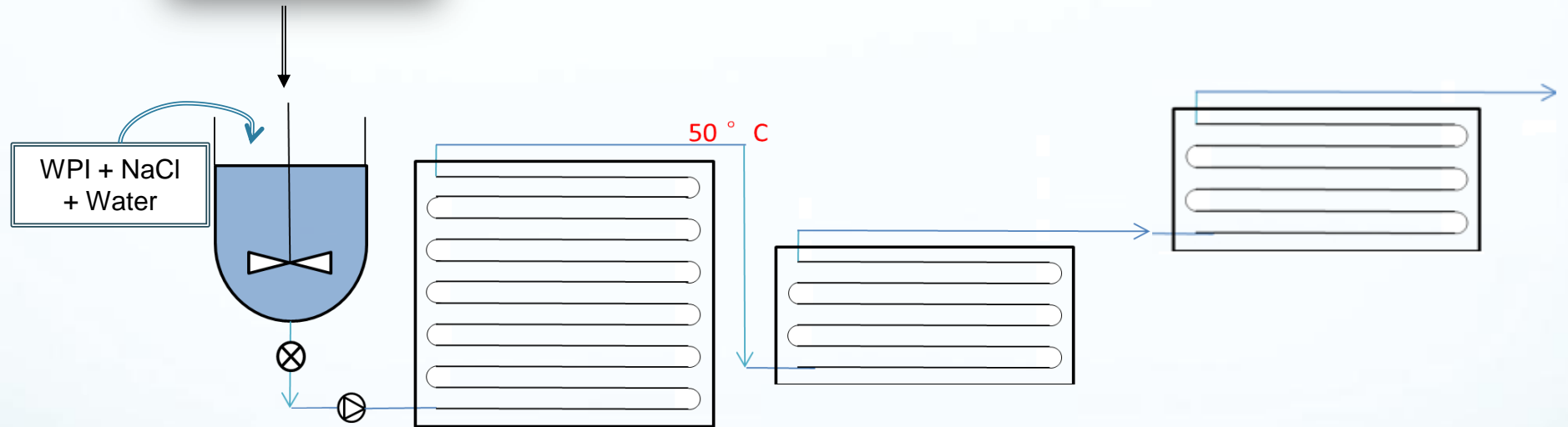
- Correlation between size/form factor (As-FIFFF) and microscopic visual



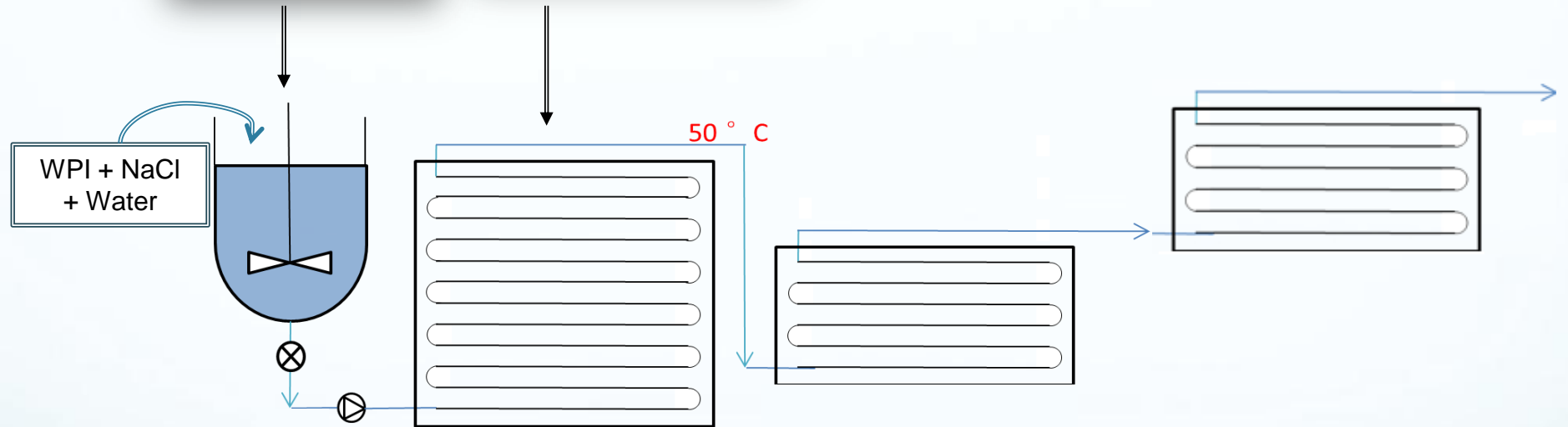
Aggregates production by semi-industrial way



Aggregates production by semi-industrial way



Aggregates production by semi-industrial way



Preheating
20°C → 50°C

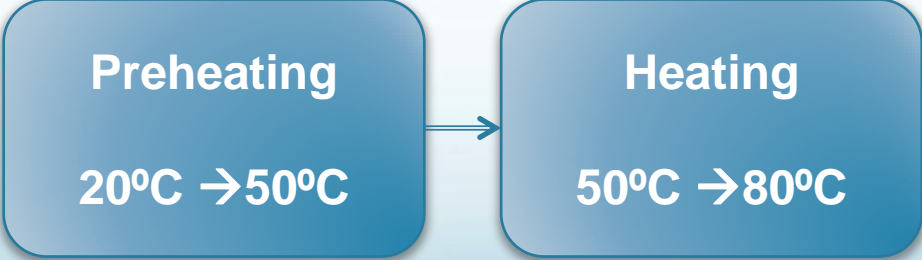
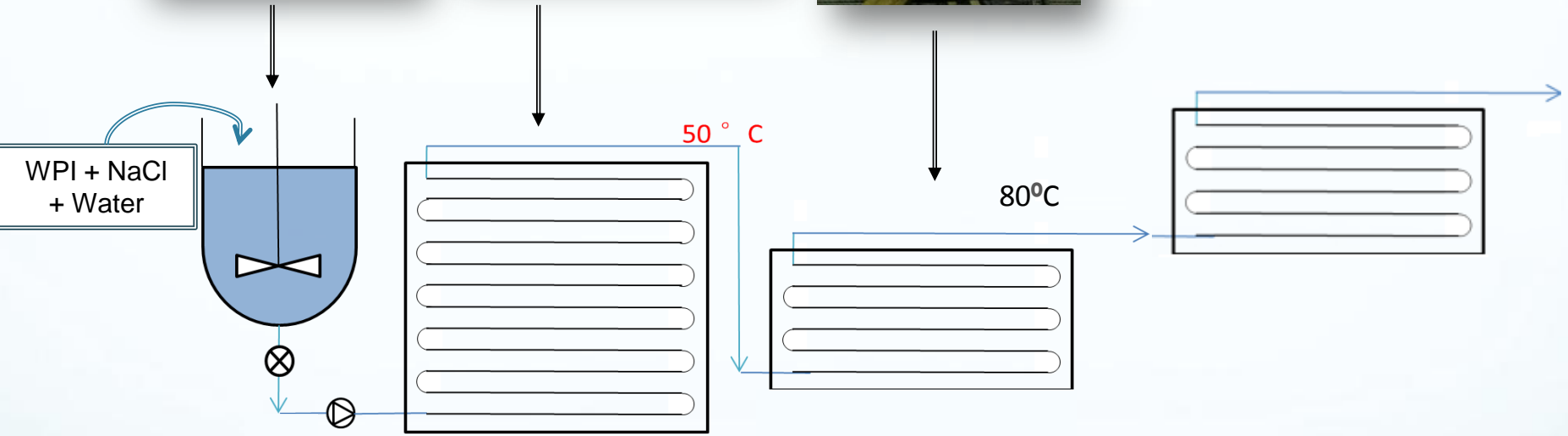
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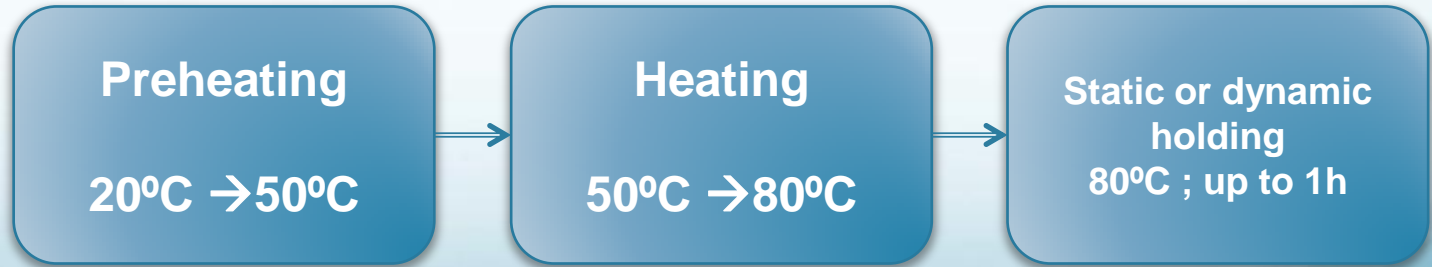
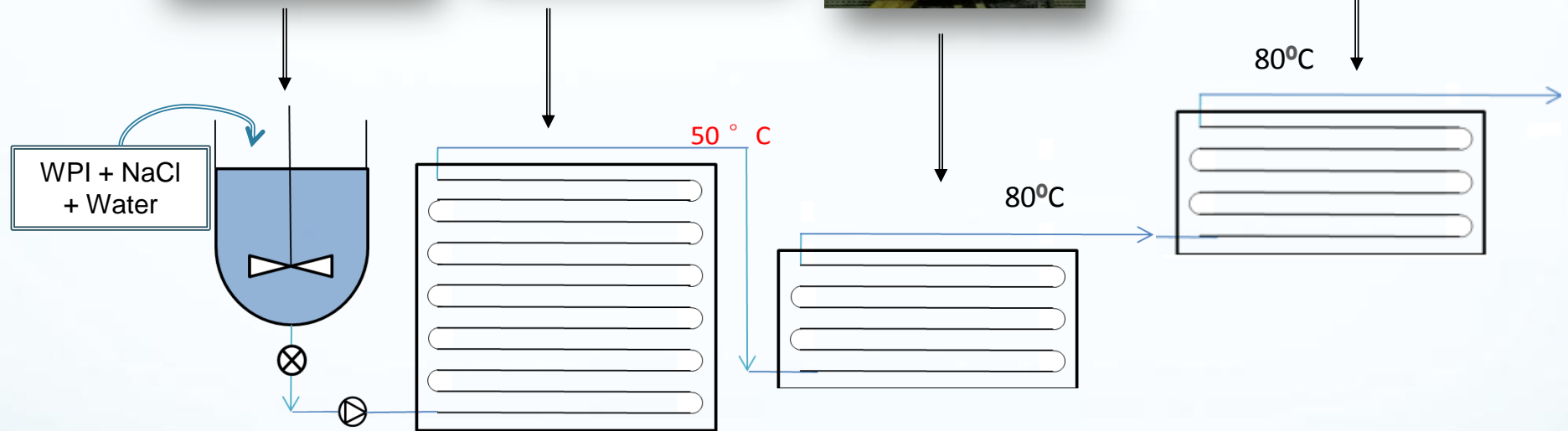
Conclusion and perspectives

Aggregates production by semi-industrial way



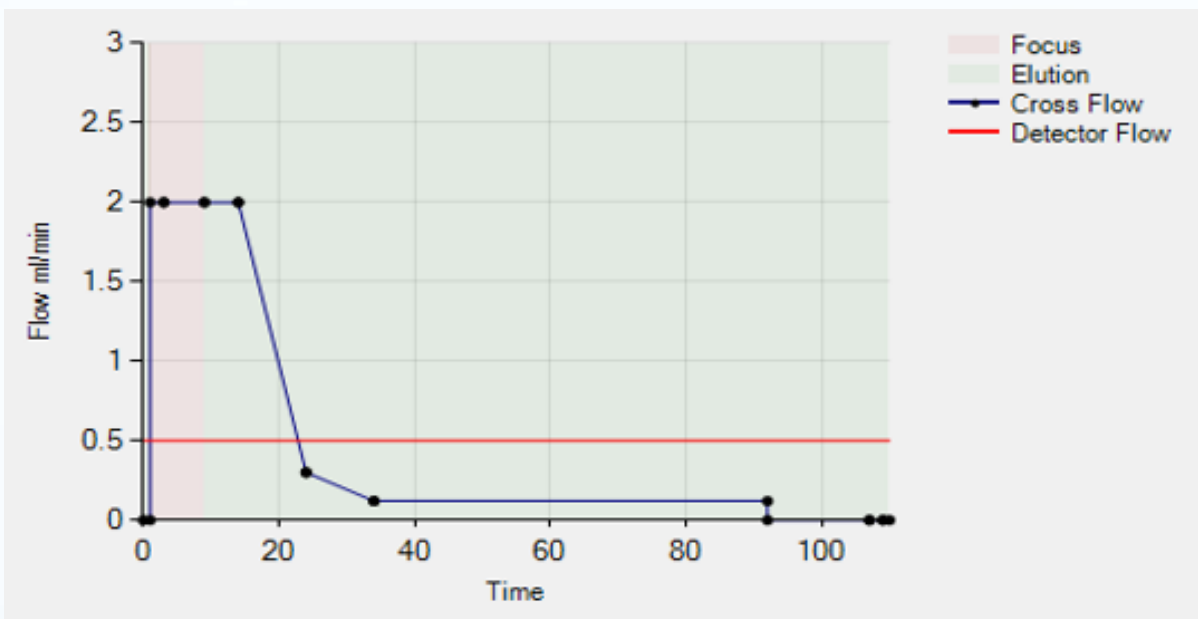
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Aggregates production by semi-industrial way



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As-FIFFF method



✓ Wyatt Dualtec System

✓ MALS

✓ Differential Refractometer

Elution : 1 min

Focus : 2 min

Focus + inject : 6 min

Eluent	Ultrapure water + 0.02% NaN_3 + 45mM NaCl
Membrane	Regenerated cellulose, cut-off 10kDa
Spacer (μm)	350
Canal	Short Canal
Injected mass (μg)	20

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Interest parameters

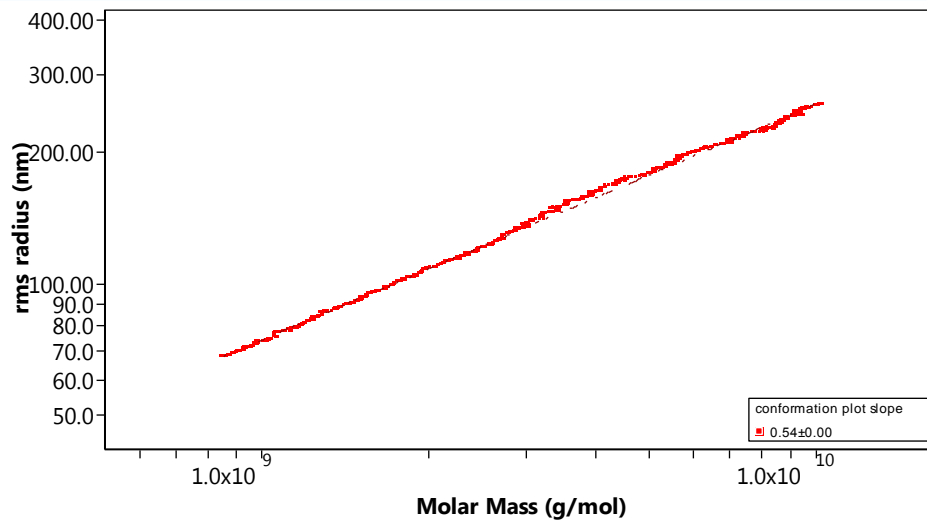
- Radius of gyration (R_g) distribution
- Molar mass (M) distribution

Interest parameters

- Radius of gyration (Rg) distribution
- Molar mass (M) distribution
- Normalized concentration: Concentration from RI divided by Calculated Mass (=Mass recovery) from refractometer ; $dn/dc=0,1850$ (it allows to compare samples)

Interest parameters

- Radius of gyration (R_g) distribution
- Molar mass (M) distribution
- Normalized concentration: Concentration from RI divided by Calculated Mass (=Mass recovery) from refractometer ; $dn/dc=0.1850$ (it allows to compare samples)
- Fractal dimension (D_f): $\log(R_g) \cdot D_f = f(\log(M))$



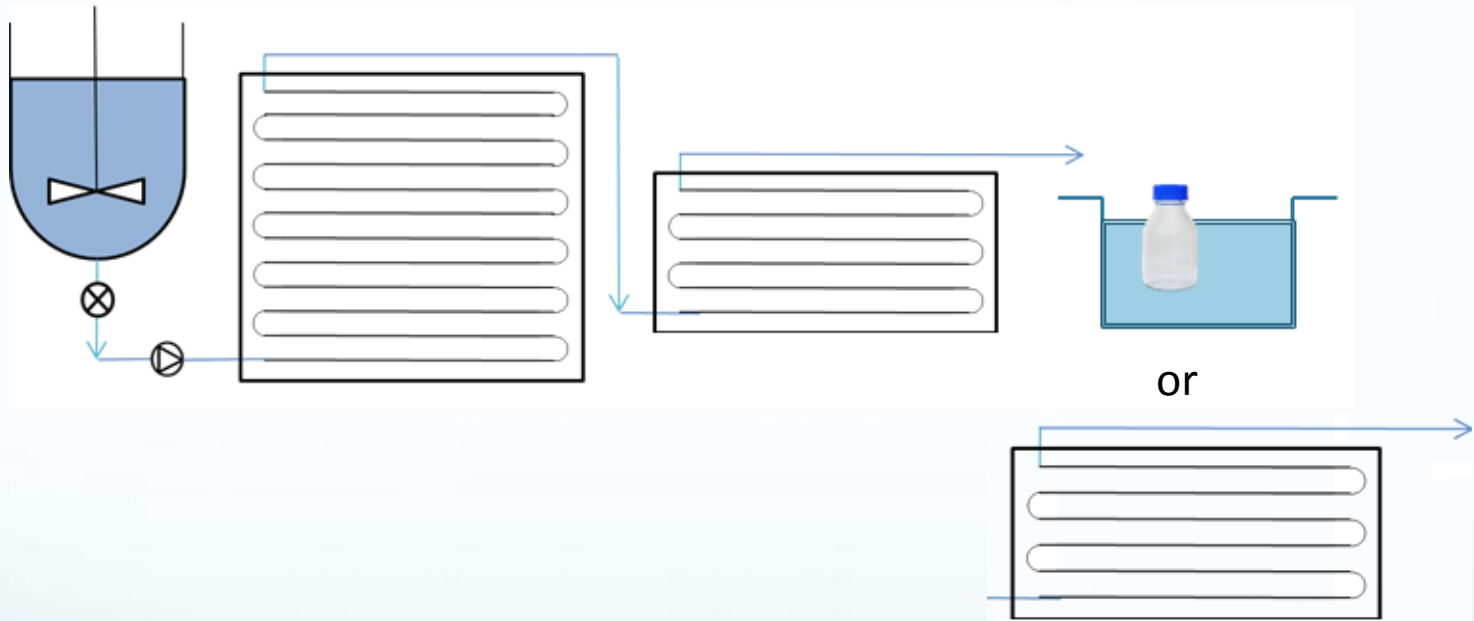
Molecular conformation model :

$D_f = 1$: Rod shape

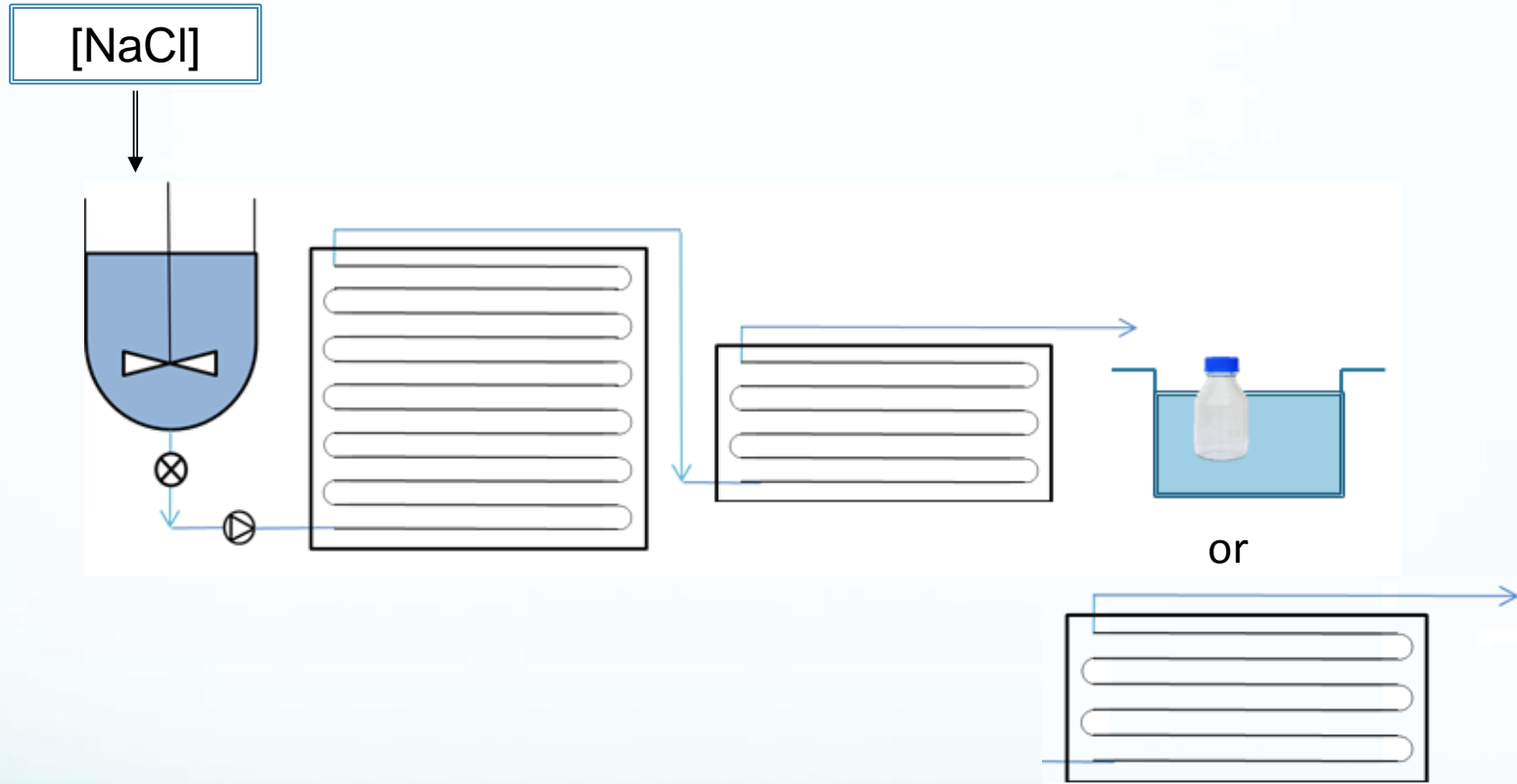
$D_f \sim 2$: Coil

$D_f = 3$: Spherical shape

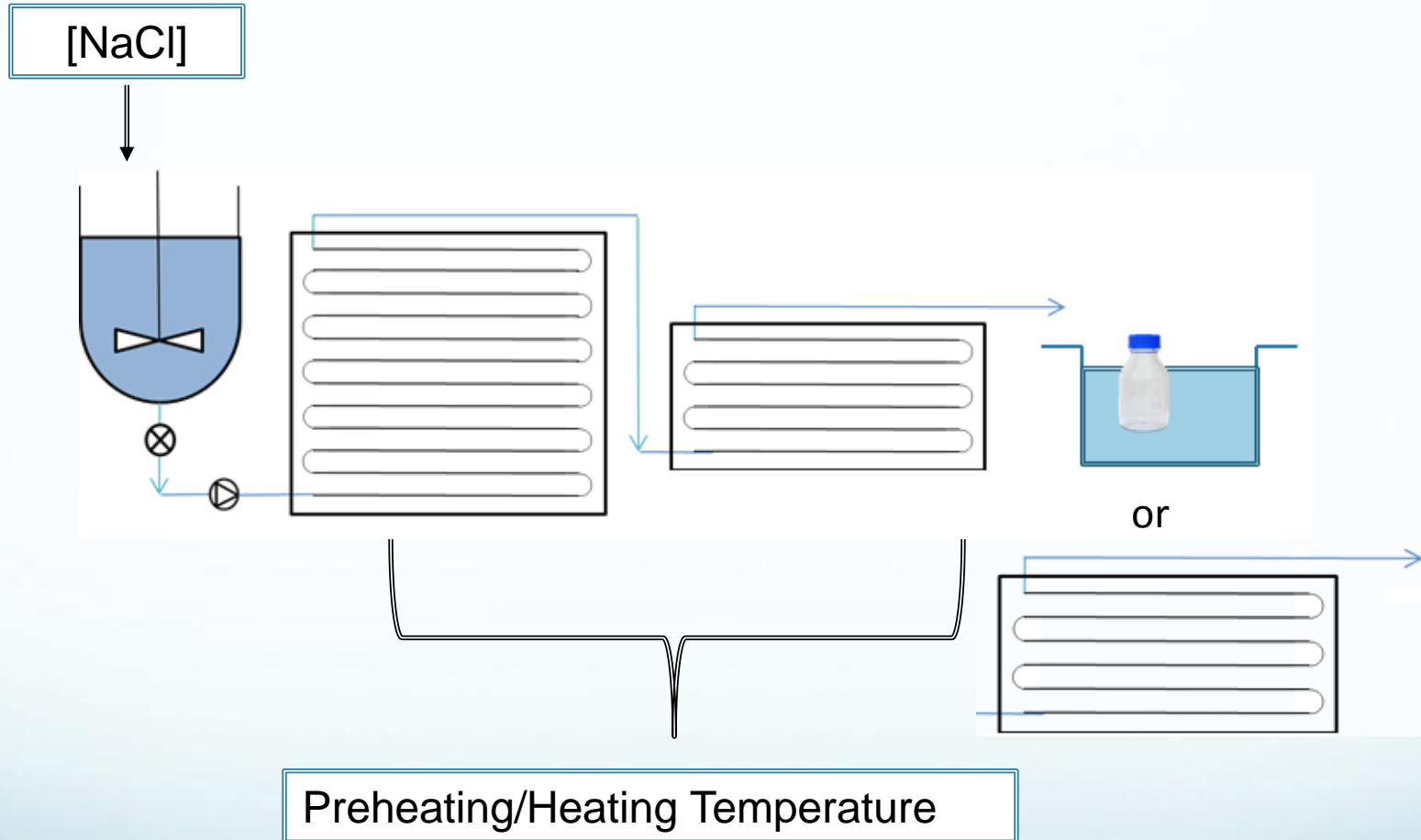
Size/shape aggregate control : process parameter variations



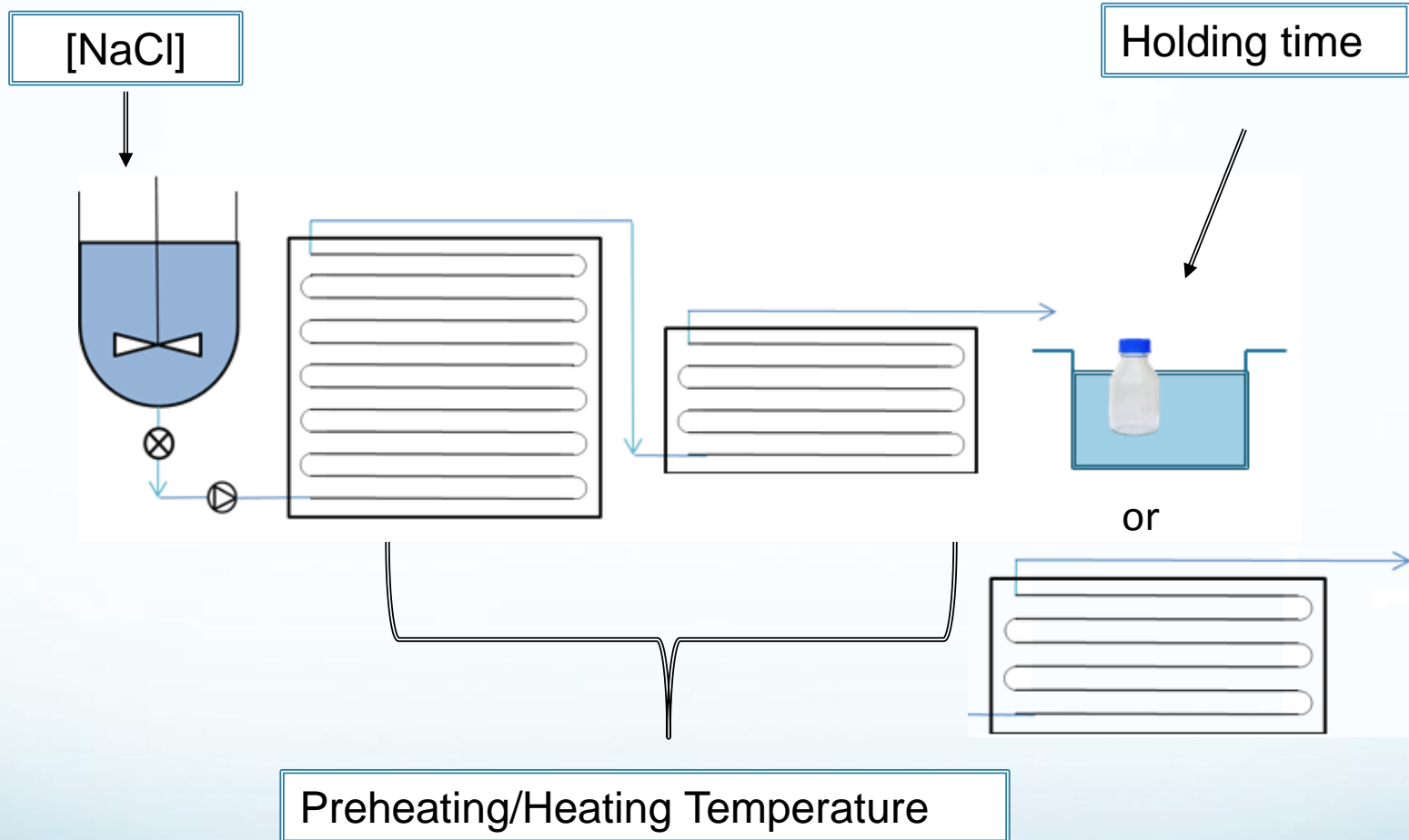
Size/shape aggregate control : process parameter variations



Size/shape aggregate control : process parameter variations



Size/shape aggregate control : process parameter variations



Preheating/Heating Temperature

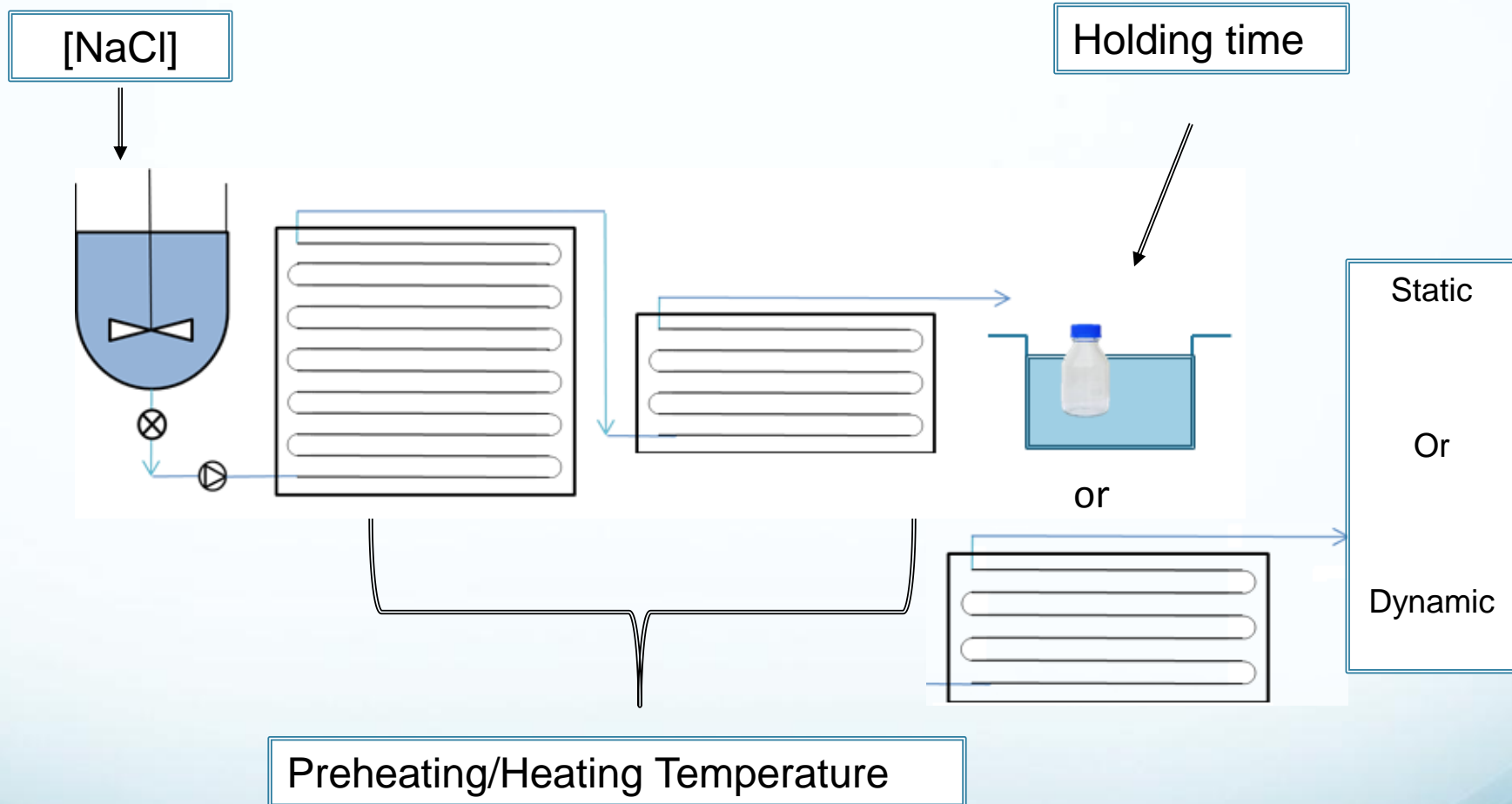
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Size/shape aggregate control : process parameter variations



First production results

5% proteins ; **10mM** NaCl ; **Static** holding ; **15min** of holding time ; **50/80°C** Preheating/Heating

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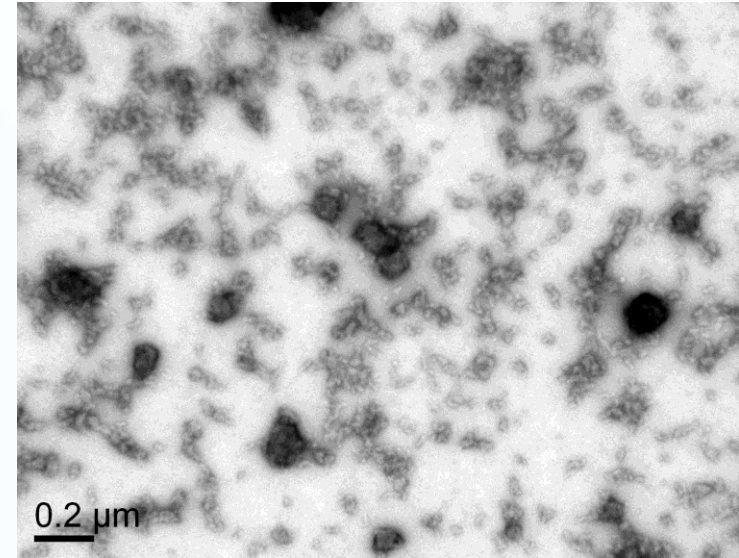
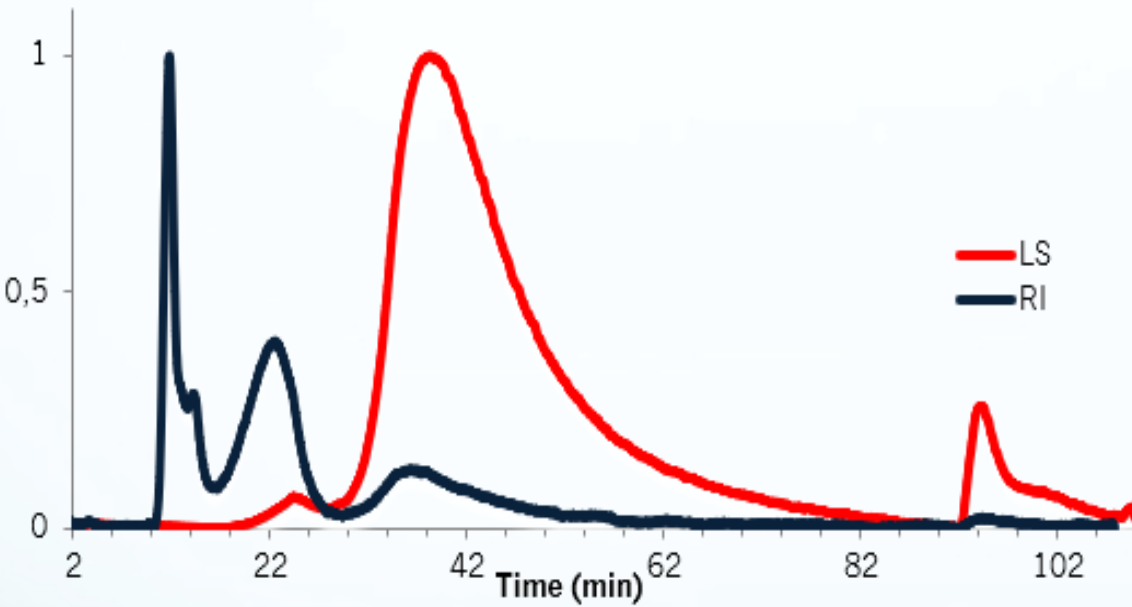
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Relative Scale



Transmission Electronic microscopy
(negative coloration)

Introduction

Methods

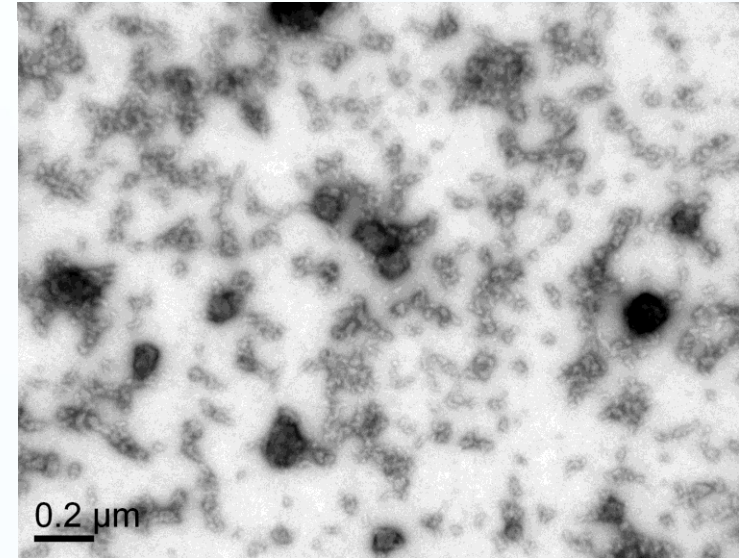
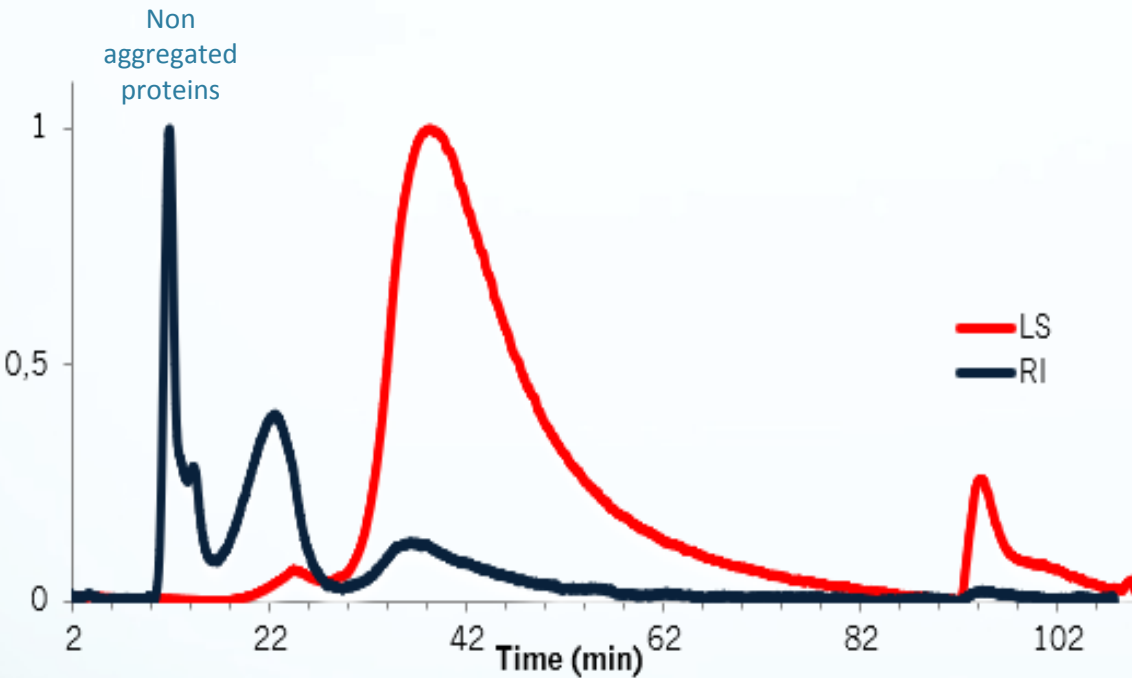
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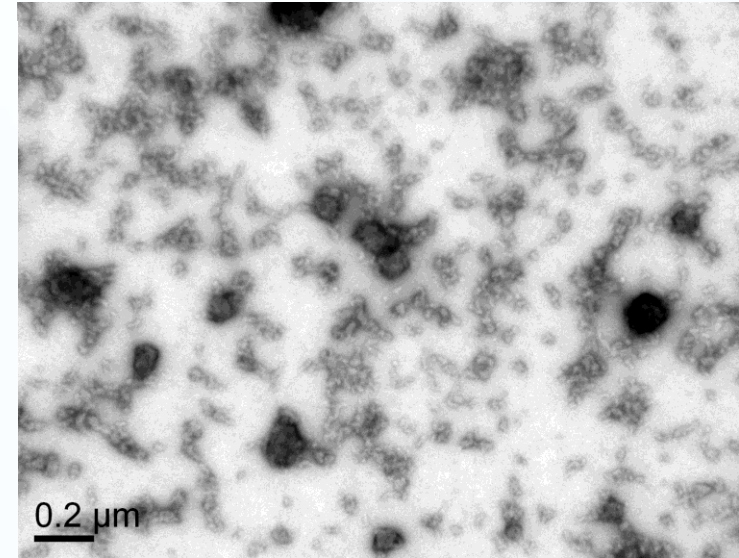
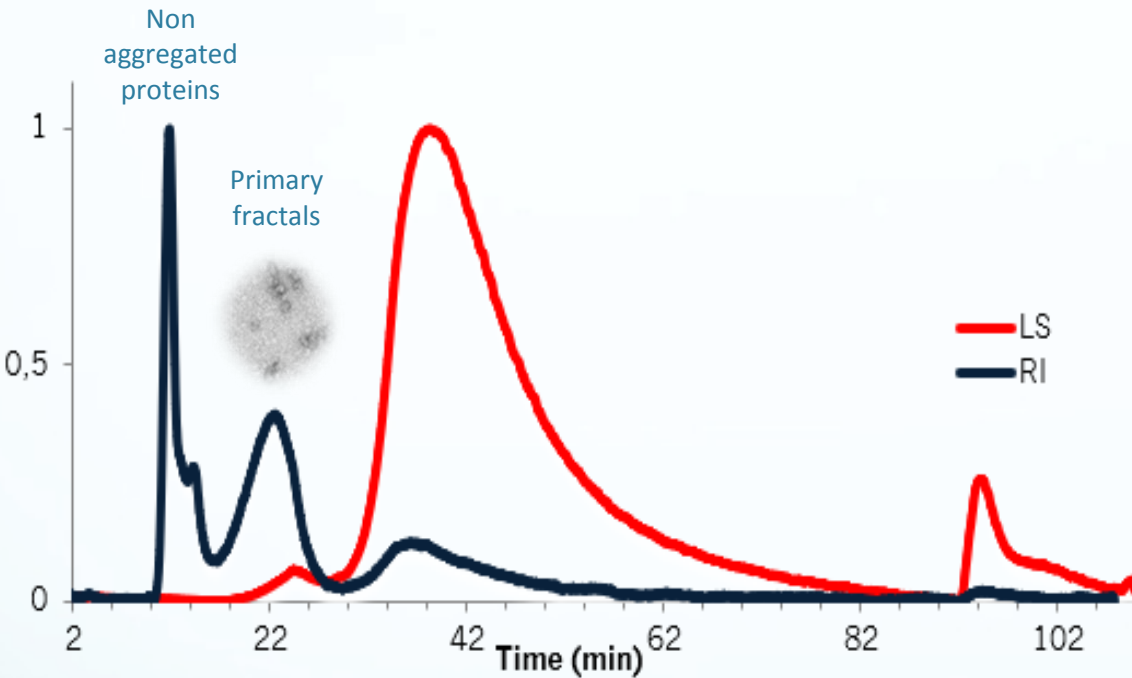
Transmission Electronic microscopy
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- 1st population (10-15min): non aggregated whey proteins

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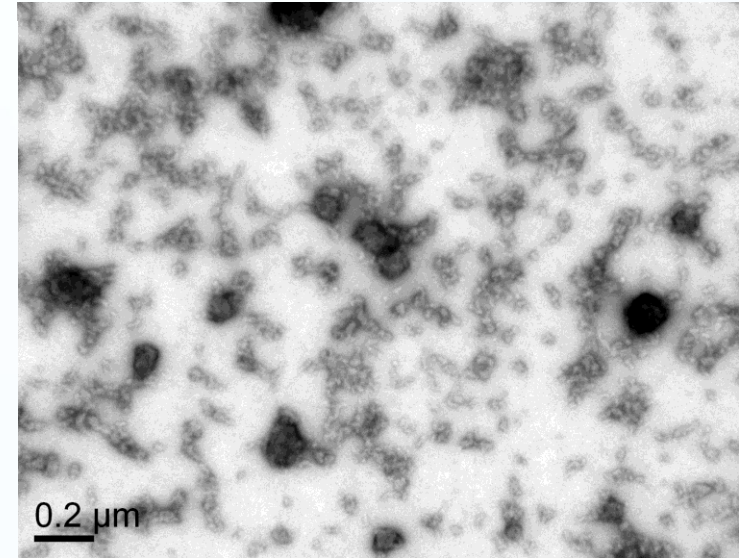
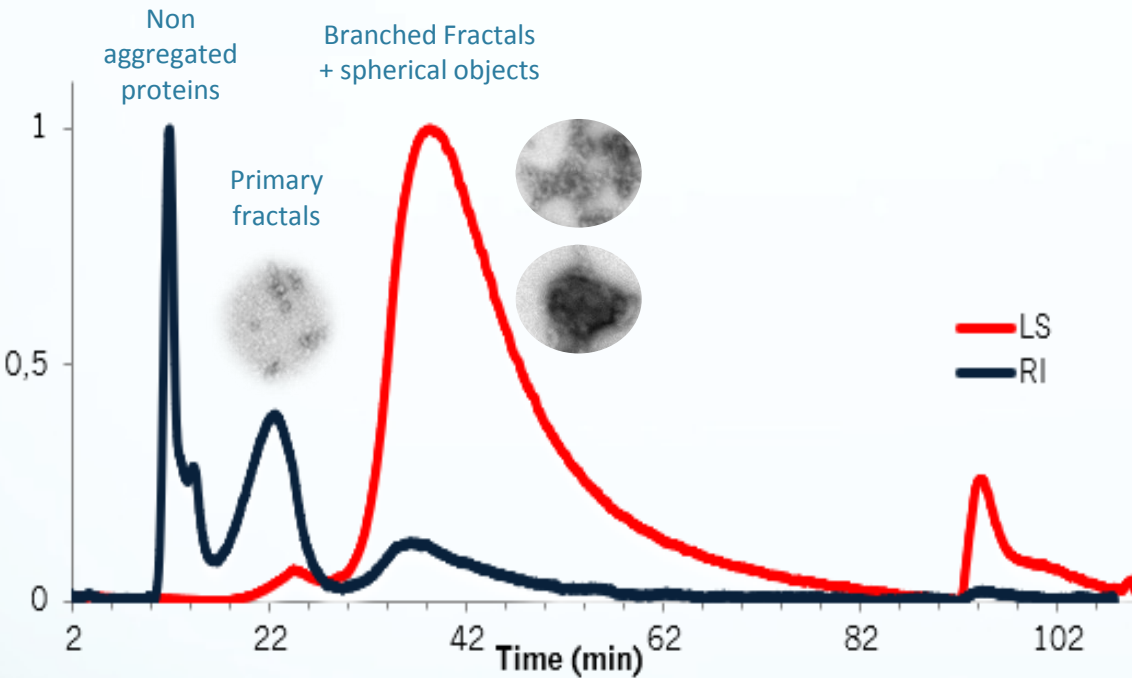
Transmission Electronic microscopy
(negative coloration)

- 1st population (10-15min): non aggregated whey proteins
- 2nd population (15-30min): primary fractals

First production results

5% proteins ; 10mM NaCl ; Static holding ; 15min of holding time ; 50/80° C Preheating/Heating

Relative Scale



Transmission Electronic microscopy
(negative coloration)

- 1st population (10-15min): non aggregated whey proteins
- 2nd population (15-30min): primary fractals
- 3rd population (after 30min): mix between large fractals and spherical objects

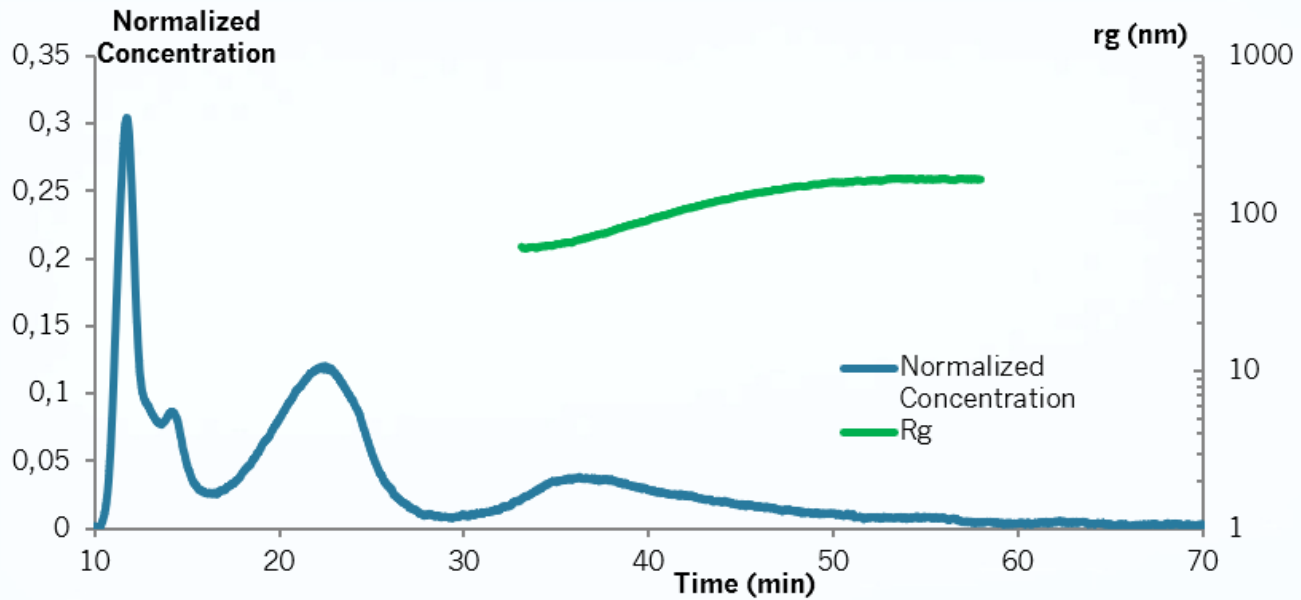
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First production results



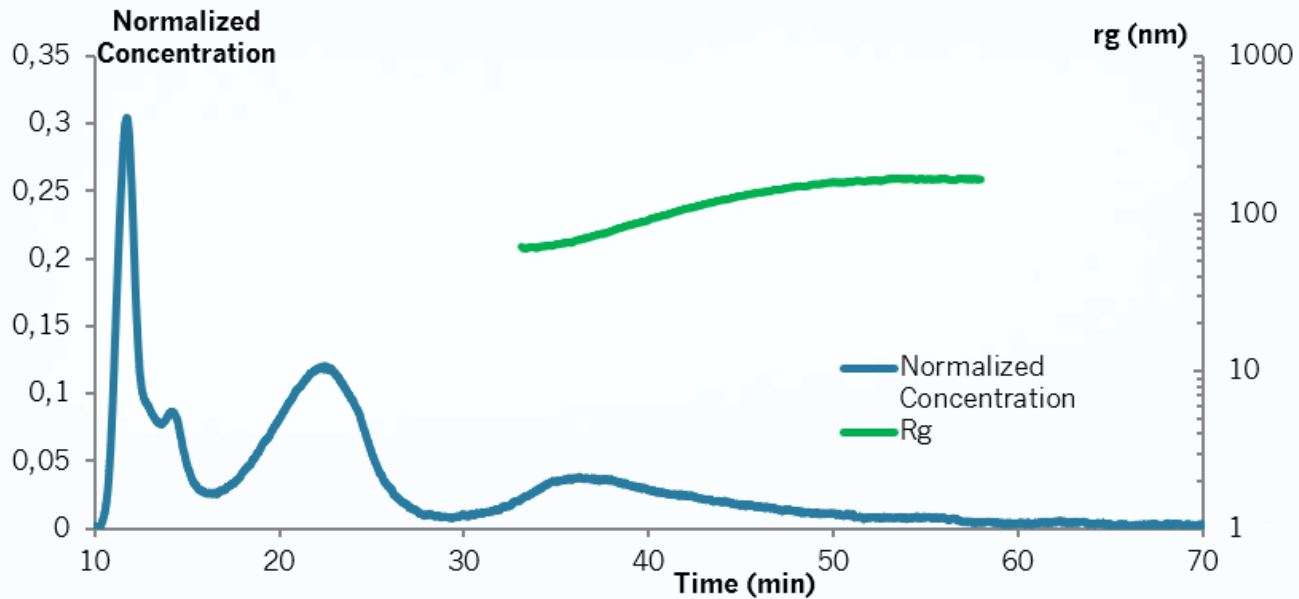
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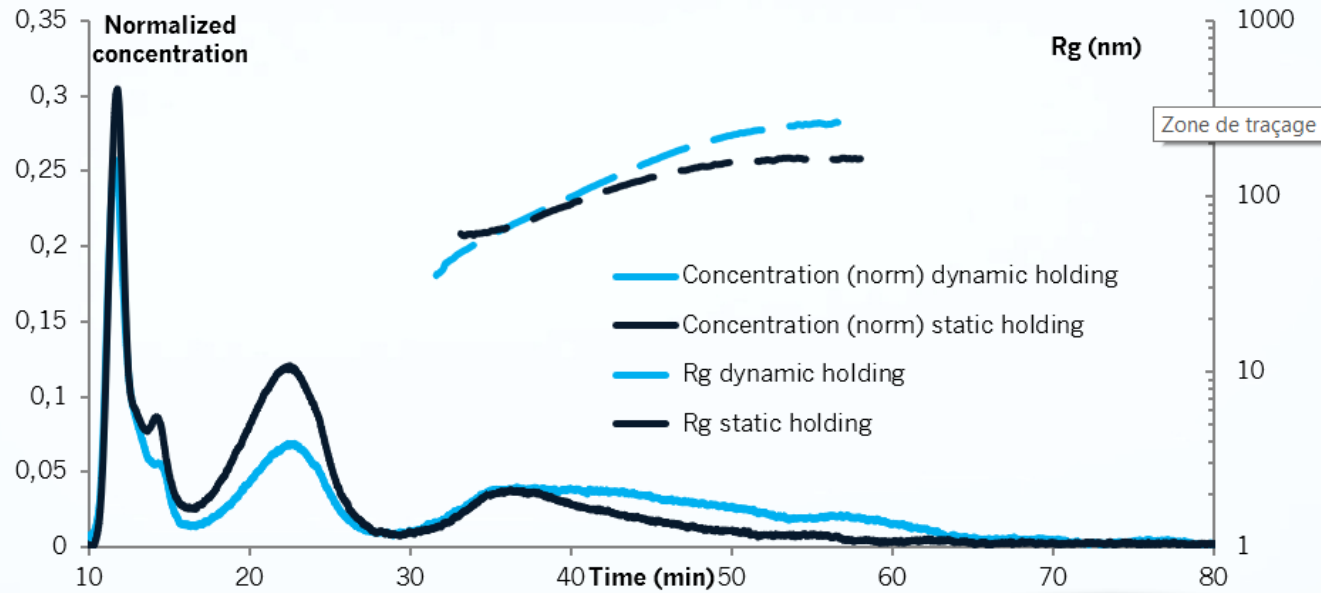
First production results



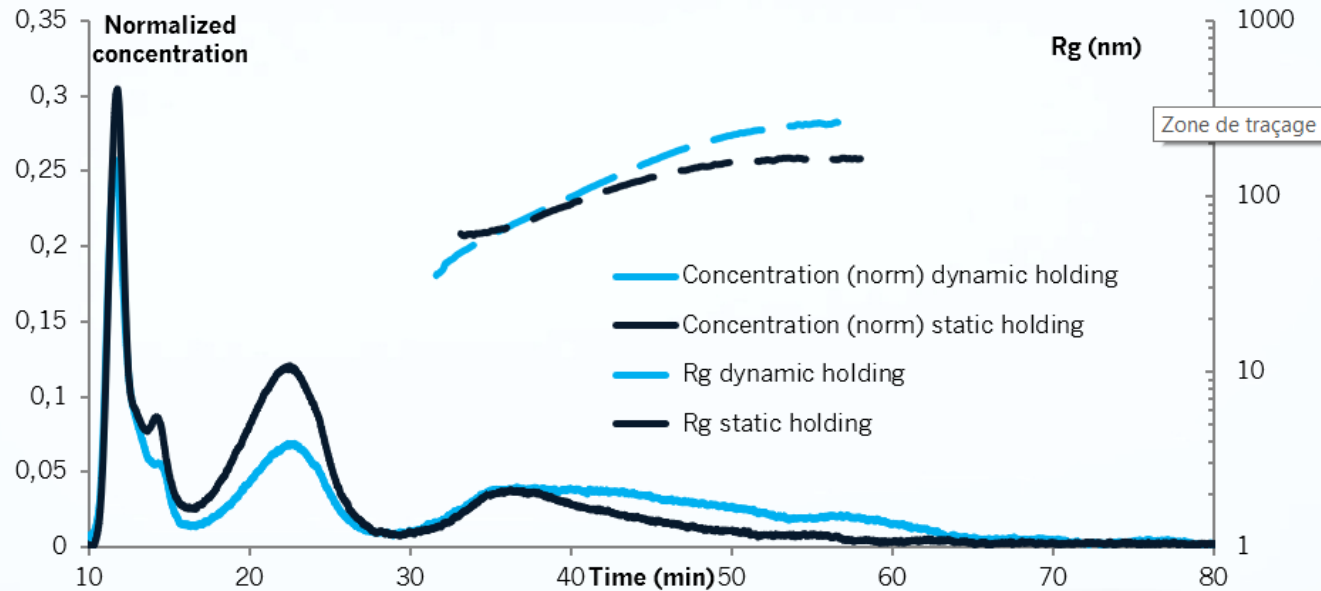
	\overline{Mw} (g/mol)	\overline{Rg} (nm)	Mass fraction (%)
1st population	/	/	30
2nd population	$8,5 \cdot 10^6 \pm 14\%$	/	40
3rd population	$8,3 \cdot 10^8 \pm 1,3\%$	$79 \pm 3\%$	30

Objective : Vary process parameters to obtain largest aggregates

Dynamic/static holding

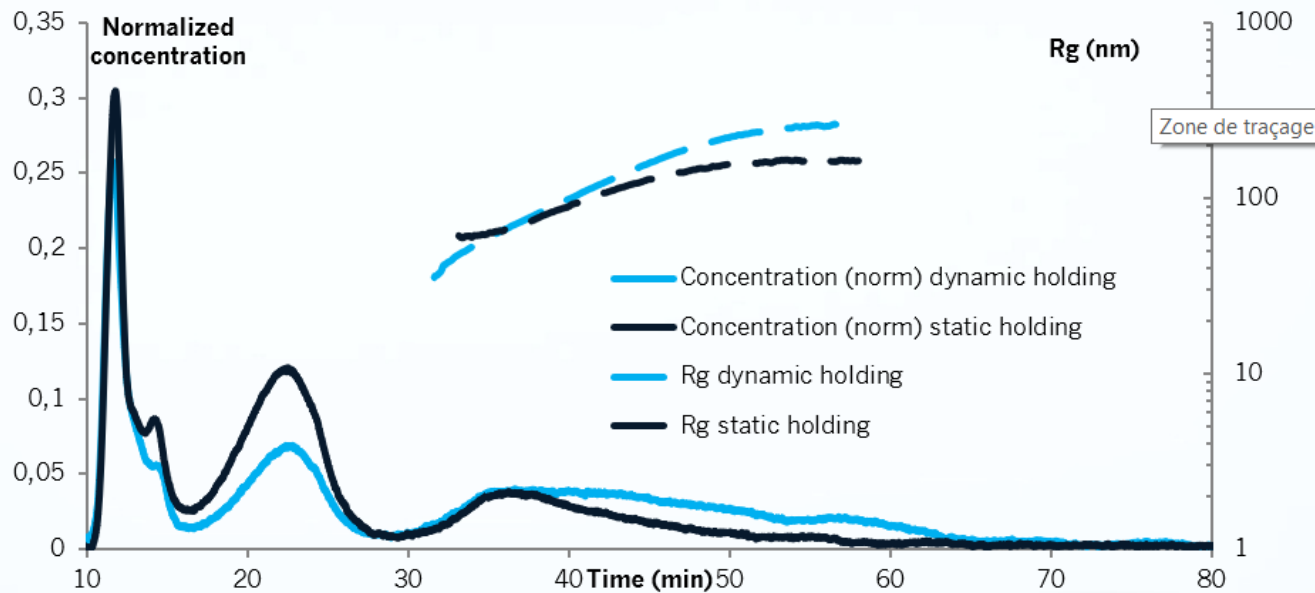


Dynamic/static holding



Primary fractals aggregate together
to form larger objects
 $\overline{R_g} = 95\text{nm} \pm 3\%$

Dynamic/static holding



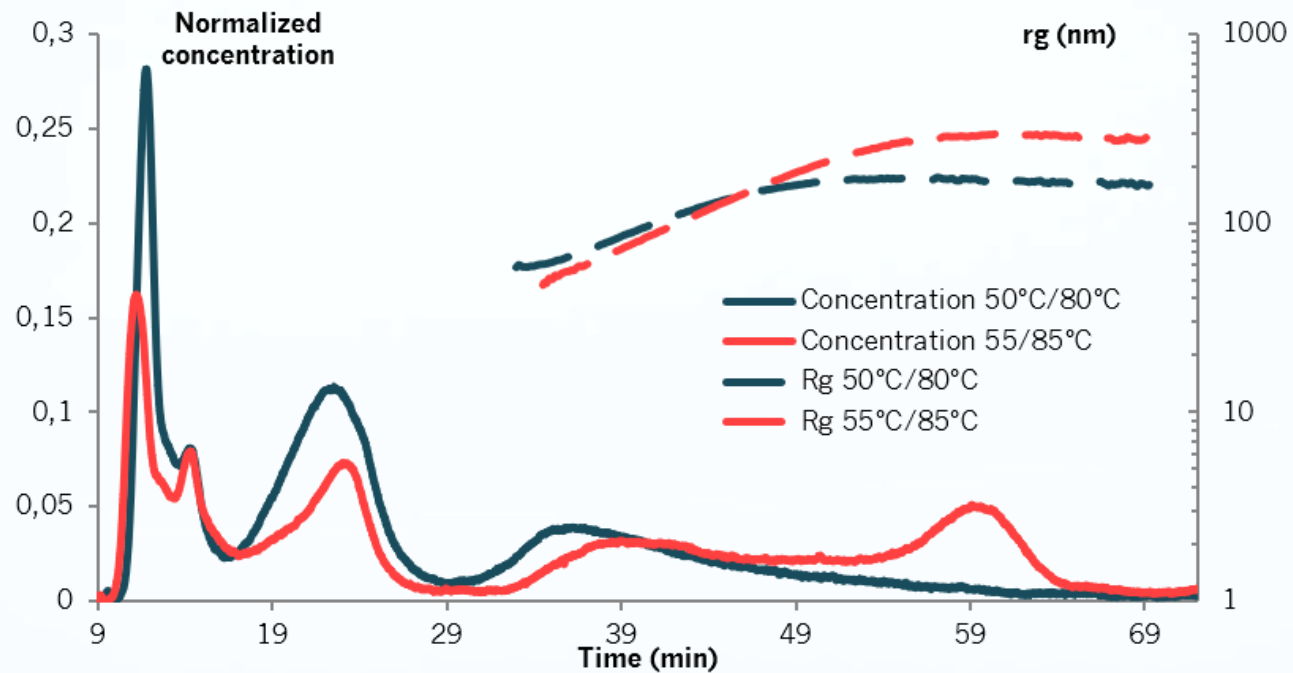
	<i>Static</i> Mass fraction (%)	<i>Dynamic</i> Mass fraction (%)
1st population	30	30
2nd population	40	26
3rd population	30	44

Primary fractals aggregate together to form larger objects

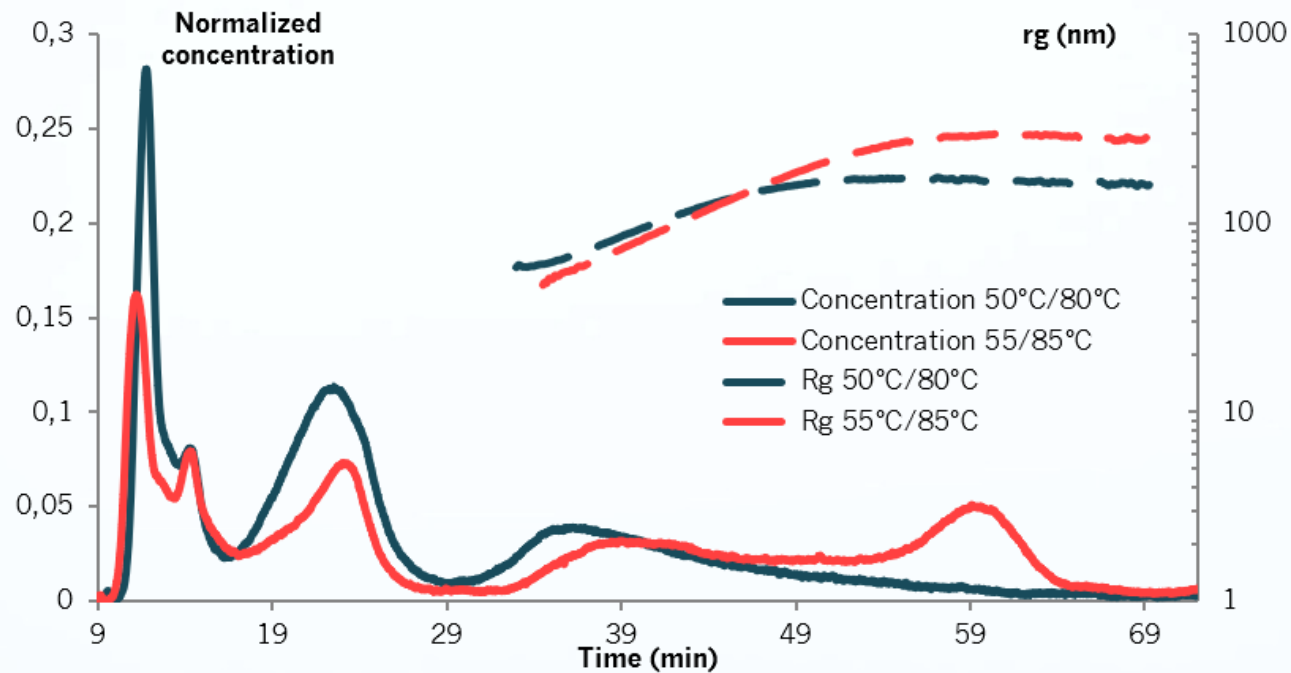
$\overline{Rg} = 95\text{nm} \pm 3\%$

Interaction probability between fractals is more important in dynamic system

Preheating/Heating Temperature Variation 50/80° C \implies 55/85° C



Preheating/Heating Temperature Variation 50/80° C \longrightarrow 55/85° C

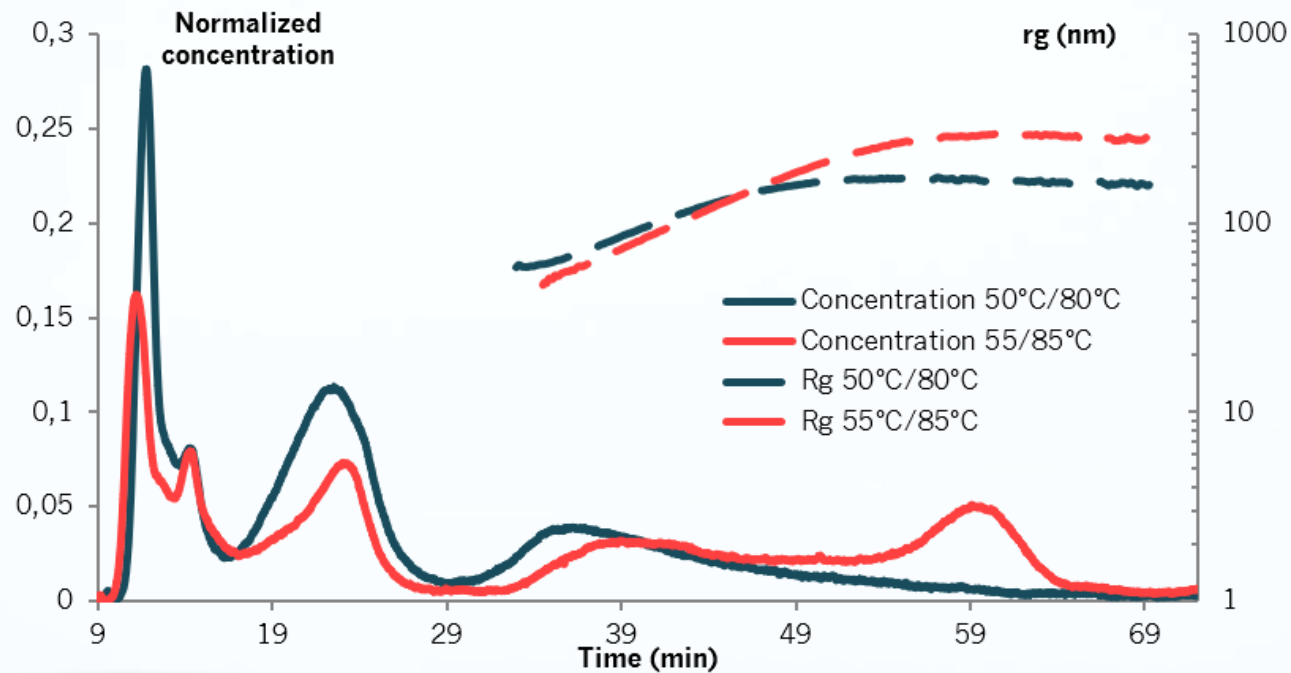


New population formation (55-65min)
by increasing preheating/heating
temperature

$$\overline{Rg} = 285\text{nm} \pm 0,7\%$$

Preheating/Heating Temperature Variation

50/80° C \longrightarrow 55/85° C

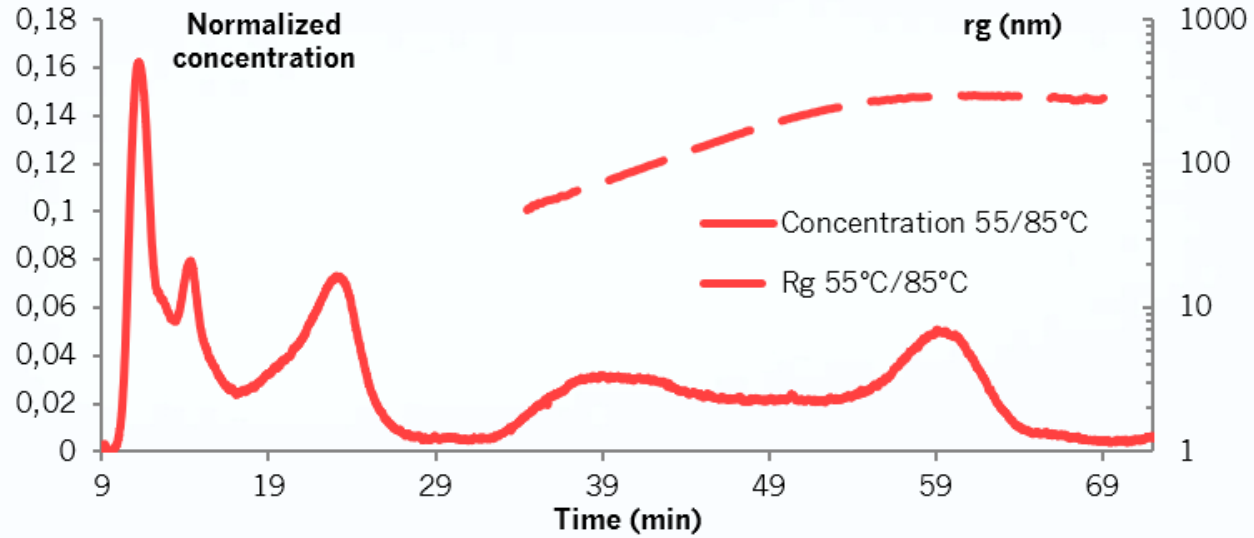


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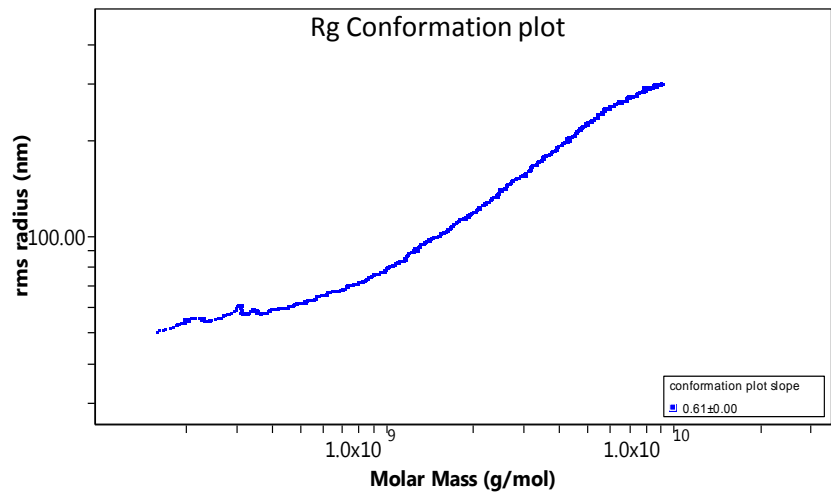
$$\overline{Rg} = 285\text{nm} \pm 0,7\%$$

→ Whey proteins and primary fractals
concentration reduces to form a 4th
largest population

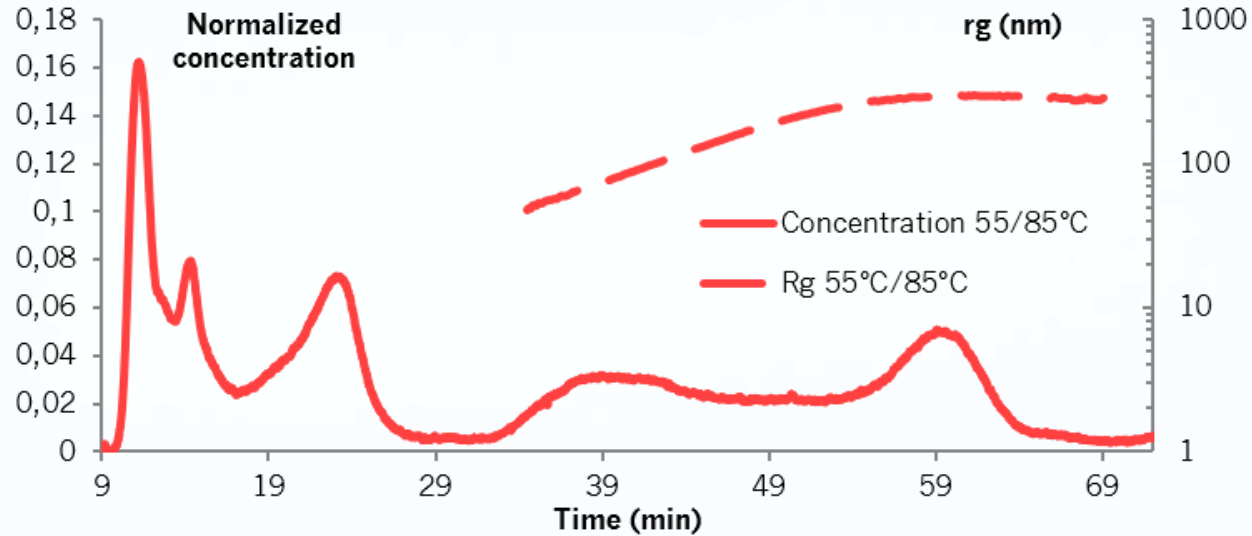
Fractal Dimension



$$\log(Rg) = \frac{1}{Df} \cdot f(\log(Mw)) \text{ between 35-65min}$$

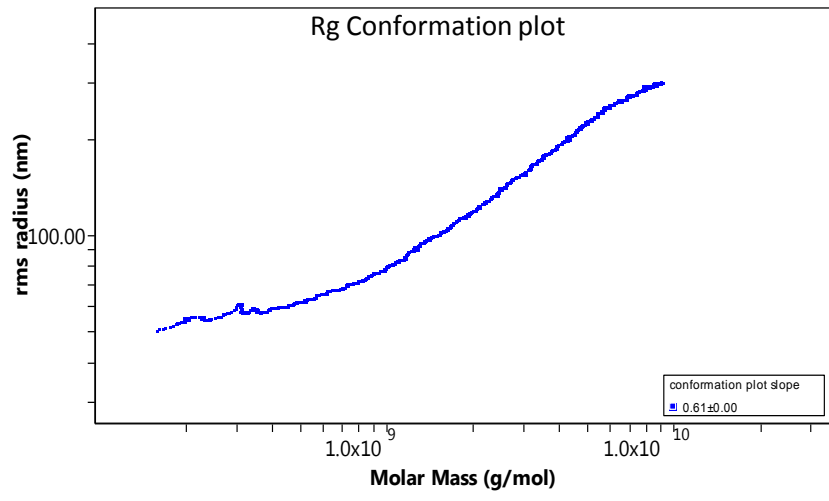


Fractal Dimension

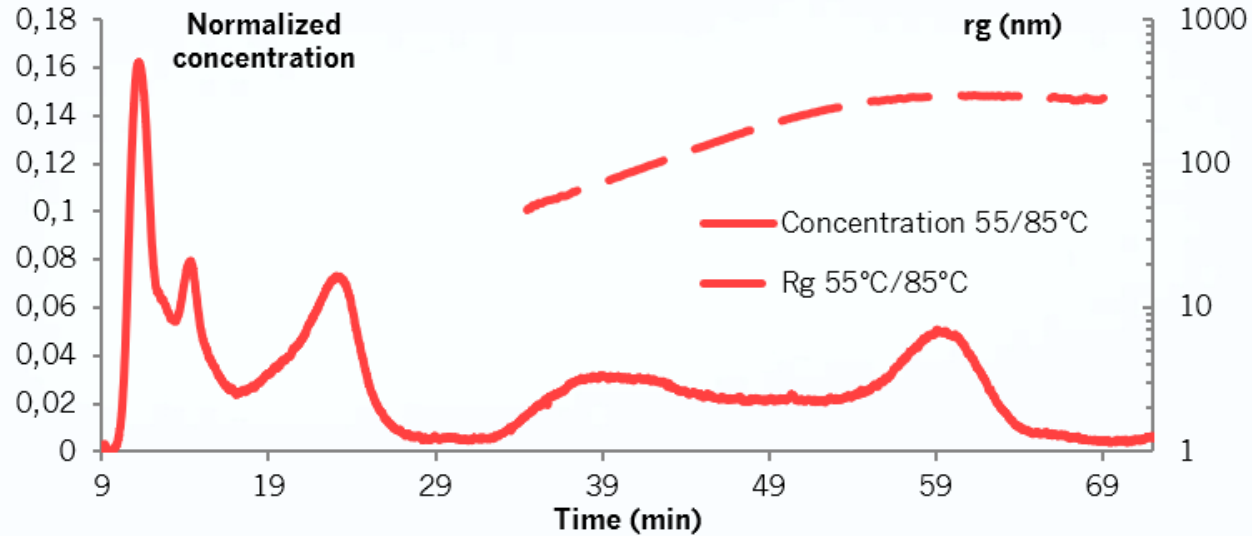


$$\log(Rg) = \frac{1}{Df} \cdot f(\log(Mw)) \text{ between 35-65min}$$

The slope increases over time (Df decreases)
 → 4th population is less dense than 3rd one
 Maybe there are less spherical objects?



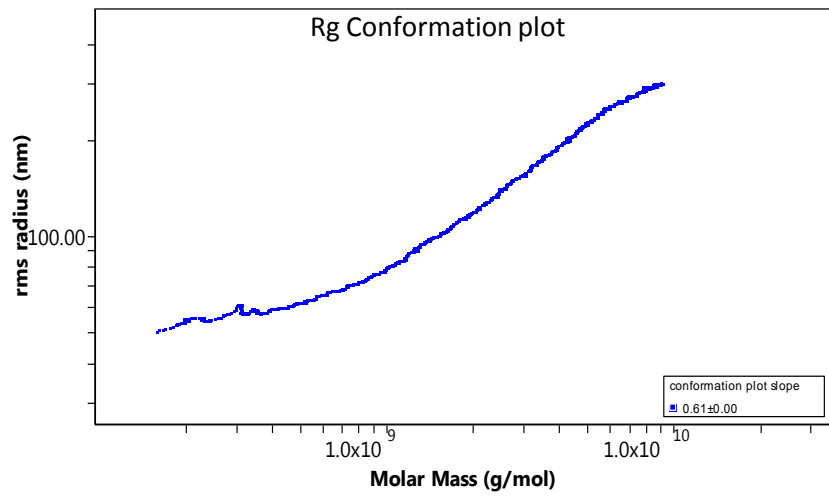
Fractal Dimension



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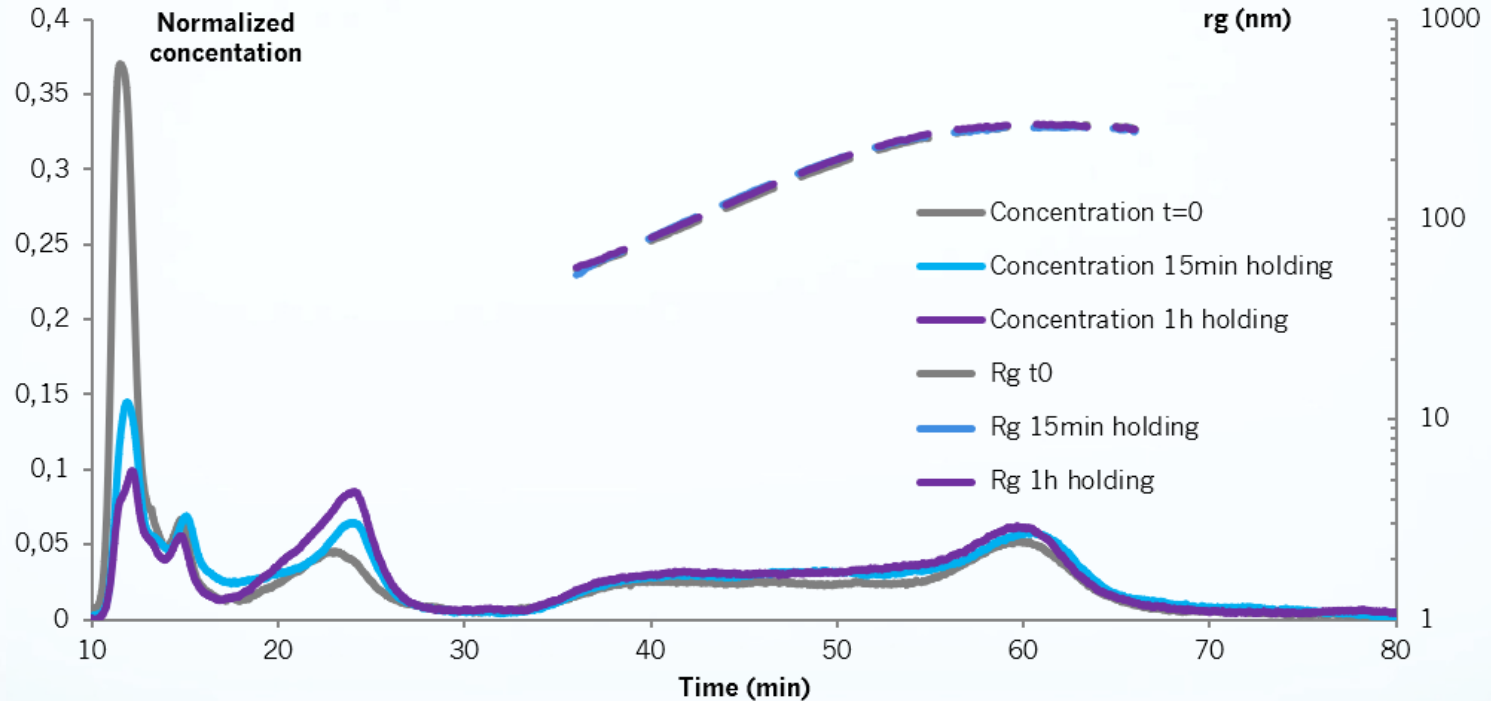
The slope increases over time (Df decreases)
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→ Fraction collection + TEM



Holding time variation

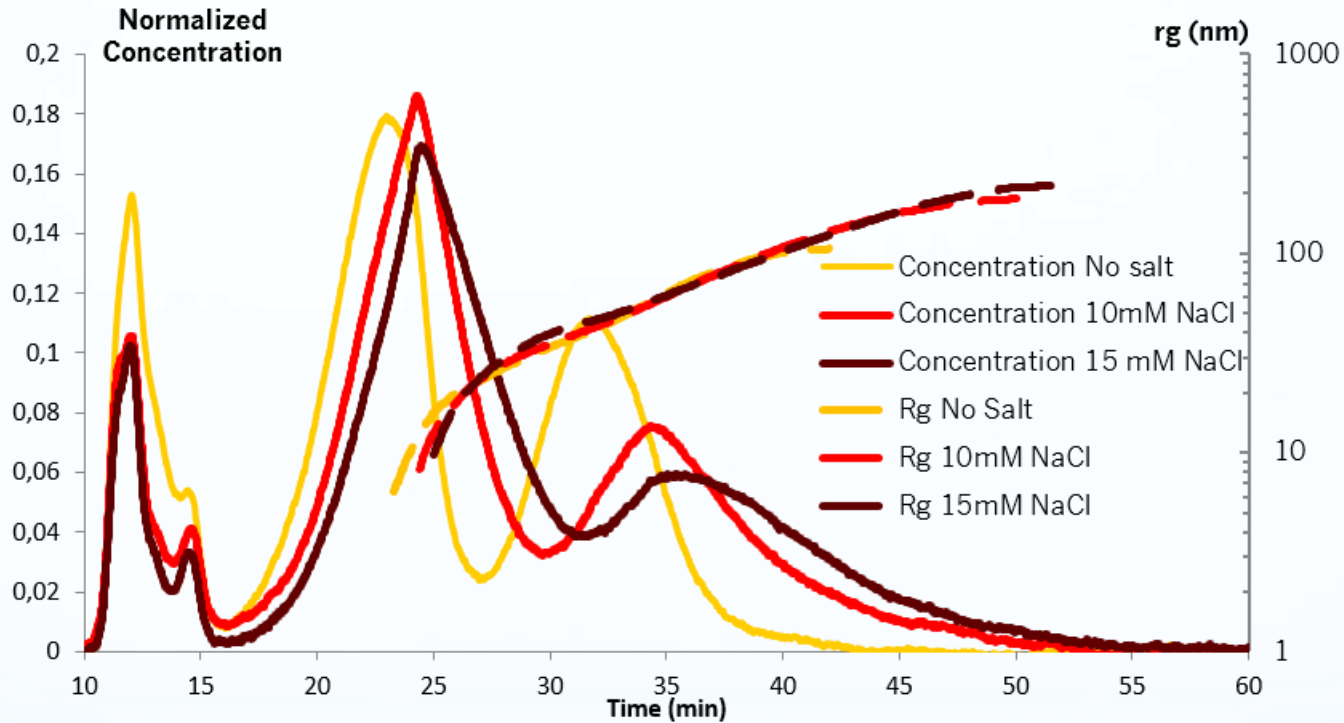
Holding time : 0/15min/1h at 80° C



- Last population size does not change with holding time
 - Large aggregates are present after heating without holding (t=0)
 - During holding, non-aggregated proteins form primary fractals
- Work on pre-heating and heating parameters

Salt Variations

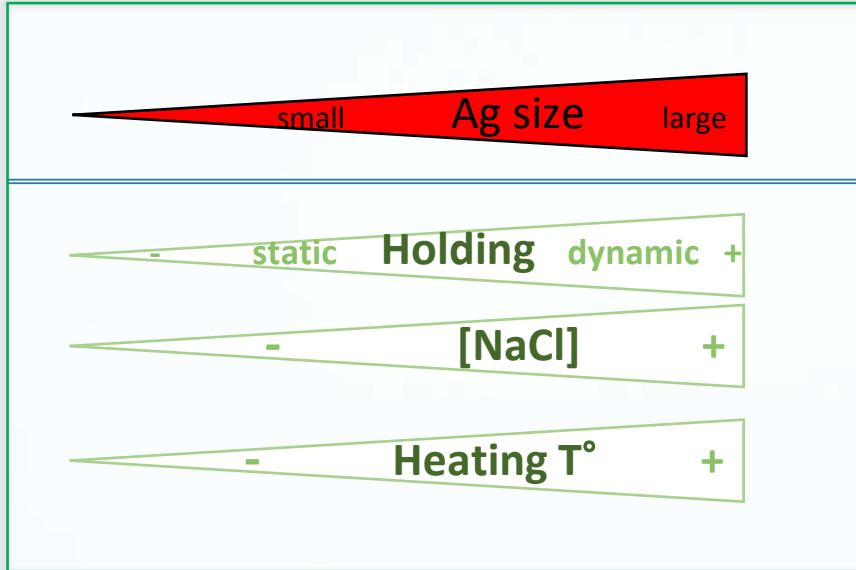
No Salt/10mM/15mM NaCl



The more salt concentration increases, the larger aggregates are formed

→ Salt decreases interaction hiding charges: aggregates are produced easier

Conclusions and Perspectives



Conclusions and Perspectives



small **Ag size** large

- static **Holding** dynamic +

- **[NaCl]** +

- **Heating T°** +

As-FIFFF

- Fractionate whey protein aggregates
- Filtration-free injection
- Informations: Radius of gyration / Molar mass / Conformation
- Tool to analyze and compare semi-industrial productions

Conclusions and Perspectives



small **Ag size** large

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As-FIFFF

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Perspectives :

- Flow rate variations
- Tube dimension variations
- Heating rate variations
- Fraction collection + TEM

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Thank you for your attention

