

Bio-catalytic hydrolysis of paper pulp using in- and ex-situ multi-physical approaches: Focus on semidilute conditions to progress towards concentrated suspensions

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1	BIO-CATALYTIC HYDROLYSIS OF PAPER PULP USING IN- AND
2	EX-SITU MULTI-PHYSICAL APPROACHES: FOCUS ON
3	SEMIDILUTE CONDITIONS TO PROGRESS TOWARDS
4	CONCENTRATED SUSPENSIONS
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In order to make 2nd-generation biofuels more competitive, high solid-matter content 20 21 has to be reached. To progress towards this target, the mechanism for destructuring 22 lignocellulose fibres in semidilute conditions has to be well understood, as this 23 configuration shows the basic mechanism which limits transfers and efficiency. This 24 study aims to delve deeply into the biophysical and transfer limitations occurring during 25 enzymatic hydrolysis. A specific experimental set-up associating in-situ and ex-situ 26 physical (rheometry, chord length analysis) and biochemical analysis was used to 27 expand the knowledge of hydrolysis of extruded softwood paper pulp over 24 h under 28 different substrate concentrations (1% to 3%) and enzyme doses (Accellerase 1500, 5 29 and 25 FPU/g cellulose). Non-Newtonian behaviour associated with pronounced yield 30 stress stand as the major factors limiting process efficiency. A critical time was deduced 31 from viscosity evolution, and the existence of a unique, dimensionless viscosity-time 32 curve was established, suggesting similar mechanisms for fibre degradation. In addition, 33 chord length distribution allowed for the description of population evolution and was 34 discussed in the light of in-situ viscosity and hydrolysis yield. Physical (viscosity, 35 particle size) and biochemical (substrate) kinetics were modelled (second-order) and 36 coefficients identified. A chronology of the encountered phenomenological limitations 37 demonstrates the necessity of optimising bioprocesses by considering physical 38 parameters. A reference feed rate is proposed in order to reach high solid loading under 39 fed-batch strategy.

40

41 Keywords: biorefinery; enzymatic hydrolysis; kinetics modelling; paper pulp;
42 rheometry; yield stress.

43

44 Nomenclature

С	Torque (mixing system)	(N.m)
Cm	Mass concentration	(gdm.L ⁻¹)
Cellu	Cellulose content	(%)
DM	Dry matter content	(% g.L ⁻¹)
DP	Degree of polymerisation	(/)
E/S	Enzyme/substrate ratio	(mL.g ⁻¹)
[Glc]	Glucose concentration	$(g.L^{-1})$
G'	Elastic modulus	(Pa)
G"	Viscous modulus	(Pa)
Кр	Power constant	(/)
Ks	Metzner-Otto constant	(/)
k_{μ}	Rheological kinetic coefficient	$(Pa^{-1}.s^{-2})$
k _{lc}	Granulometric kinetic coefficient	$(\mu m^{-1}.s^{-1})$
ks	Biochemical kinetic coefficient	((gdm/L) ⁻¹ .s ⁻¹)
lc	Chord length	μm
lcm	Mean chord length	μm
ms	Quantity of substrate	(g humid matter)
N	Rotation speed	(revolutions per second)
Np	Power number	(/)
Р	Power consumption	(W)
Re	Reynolds number	(/)
R ²	Correlation coefficient	(/)
$V_{\rm w}$	Water volume	(L)
Ϋ́	Shear rate	(s ⁻¹)
ρ	Density	(kg.m ⁻³)

ρ_s	Substrate density	(kg.m ⁻³)
μ	Viscosity	(Pa.s)
μo	Initial viscosity	(Pa.s)
μ_∞	Final viscosity	(Pa.s)
μ*	Dimensionless viscosity	(/)
τ	Shear stress	(Pa)
τ_0	Yield stress	(Pa)



46 **1** Introduction

47 The latest global agreement, the Paris agreement, was established in 2015 to reduce the 48 scale of the fossil-fuel-based economy. Greater use of renewable resources with low 49 carbon footprints is being advocated to substitute the usage of fossil fuels and to achieve 50 the goal of a decarbonised economy. Biofuel stands as one potential alternative, and is 51 being scrutinised in many countries [1]. This development is not only justified by 52 economic reasons, but also by societal demand and environmental constraints that 53 necessitate the move towards renewable energies. Cellulosic biomass provides a low-54 cost, renewable, and abundant resource that has the potential to support large-scale 55 production of fuels and chemicals via biotechnological routes [2]. Among the major 56 users of lignocellulose resources generated through forestry and agricultural practice, 57 the pulp and paper industry holds a strategic position. Currently, the promotion of 58 biorefineries producing multiple products, including higher-value chemicals as well as 59 fuels and power, is a major objective of numerous consolidated programs in the world. 60 In order to achieve economic viability, the biorefinement of lignocellulosic resources 61 must be operated at very high feedstock dry matter content. This strict prerequisite 62 imposes a considerable constraint, particularly on the physicochemical and biocatalytic 63 steps, which, overall, aim to produce high-quality, fermentable sugar syrups. Moreover, 64 industrial criteria regarding maximum reactor volumes, energy and water consumption, 65 and wastewater management must also be respected. The pulp and paper industry is able 66 to provide a tried and tested industrial model for processing lignocellulose biomass into 67 pre-treated cellulose pulps. The pulp product of this industry is appropriate for modern 68 biorefining because it displays low lignin content, is free of inhibitory compounds that 69 can perturb fermentations, and is devoid of microbial contaminants [3]. The process of 70 biofuel production can thus be coupled with the pulp and paper industry. Two main 71 advantages are highlighted here. The first is the perfect control of paper pulp quality due

to the efficiency of woody substrate pre-treatment and to the elimination of the lignin
fraction in the initial biomass. The second is that the pulp and paper industry produces
energy by valorising residues and co-products, thus reducing final biofuel costs [4].

In order to produce 2nd-generation biofuels, pre-treated paper pulp would be hydrolysed 75 76 and then fermented to convert the simple sugars (hexoses and also pentoses) into 77 molecules of interest. Conventionally, this process involves two separate steps: 78 hydrolysis and fermentation (SHF). In addition, it can also use simultaneous 79 saccharification and fermentation processes or, more recently, consolidated 80 bioprocessing, which integrates enzyme production, saccharification, and fermentation into a single process. In the SHF process, cellulose is enzymatically hydrolysed by 81 82 cellulases during the first step to form simple sugars (e.g. glucose), which are consumed 83 in the second step by Saccharomyces, Zymomonas, or other microorganisms to obtain the desired product [5-7]. The main advantage of SHF is the ability to carry out each 84 85 step in its optimum conditions: temperature, pH, etc.

86 The key steps are always the pre-treatment techniques and the conversion into 87 fermentable sugars [8]. A better scientific understanding, and ultimately exact technical 88 control, of these critical biocatalytic reactions, which involve complex matrices at high 89 solid content, currently pose a major challenge that must be overcome to facilitate the 90 intensification of biorefining operations. Among the main parameters to be studied, the 91 rheological behaviour of the hydrolysis suspension and the fibre particle size stand out as major determinants of process efficiency, and are responsible for the choice of 92 93 equipment to be used and the strategies to be applied [9]. Nevertheless, only a limited 94 number of studies cover the rheological behaviour of lignocellulose matrices in highly 95 concentrated suspensions during biocatalytic degradation [10].

96 The purpose of this study was to investigate the destructuring of fibre during enzyme 97 attack under a multi-scale approach using different analytical techniques. Semidilute 98 conditions were chosen, as they introduce the complexity of particle-particle

99 interactions, which are strongly involved in the transfer limitations observed in high dry 100 matter content, without inhibiting the bioreactions. Figure 1 illustrates the three blocks corresponding to the three levels of observation: macro-scale with viscometry and 101 102 rheometry, micro-scale with chord length distribution (CLD) and molecular scale with 103 biochemistry (chemical analyses of soluble fraction and solid fraction). This three-104 membered framework permits the analysis and comparison of in- and ex-situ methods 105 (excluding biochemical analysis). Experiments were conducted using three substrate 106 concentrations in a semidilute regime and two enzyme concentrations to explore the 107 impact of substrate properties and enzyme ratio on fibre destructuring kinetics 108 (bioconversion rate). Then, phenomenological models can be established and 109 considered as a whole to provide a full overview of the mechanism. In this regard, the 110 phenomenological models should fulfill certain criteria, such as reliability, simplicity, 111 and congruity with the experimental information. It is hoped that a global result from 112 these three sub-blocks will provide a "knowledge block" to explain certain scientific 113 limitations and lead to the implementation and intensification of considered 114 bioprocesses.

- 115 **2** Materials and Methods
- 116 2.1 Experimental set-up

117 An experimental set-up was specifically developed which consisted of a bioreactor and 118 a home-designed impeller system associated with several in-situ sensors (temperature, 119 pH, rotation speed, torque, FBRM sensor). The bioreactor was a homemade glass tank 120 (diameter: 130 mm, H_{max}: 244 mm, V: 2.0 L) with a water jacket for thermal regulation. 121 A specific agitator included a double impeller to minimise the difficulty in substrate 122 mixing and ensure suspension homogeneity (Figure 2). The first impeller consisted of 123 three inclined blades (diameter: 73.5 mm, angle: 45° , h = 38 mm) located 75 mm above 124 the bottom to ensure mixing. The second mixer, which had 2 large blades (diameter:

125 120 mm, h = 22 mm), was set close to the bottom to avoid substrate decantation. The 126 impeller shaft was connected to a viscometer working at a set speed (Viscotester Haake VT550, Thermo Fisher Scientific). This allowed for on-line torque measurements. The 127 128 rotational speed ranged between 0.5 and 800 rpm, and torque ranged between 0.1 and 129 30 mN.m (< 400 rpm) and 20 mN.m (> 400 rpm) (accuracy $\pm 0.5\%$). Temperature was 130 controlled by circulation (Haake DC30 cryostat, -50 to 200 °C ±0.01, Thermo 131 Scientific) through the water jacket. Suspension pH was monitored by a pH meter 132 (Mettler Toledo Seveneasy S20, 0-14±0.01, -5 to 105 °C) and the pH adjusted with 0.5 133 N NaOH or 0.5 N H₂SO₄. The viscometer and the cryostat were controlled by 134 HaakeRheoWin Job Manager software (Thermo Fisher Scientific) which also ensured 135 data recording (temperature, torque, mixing rate). A focused beam reflectance sensor 136 (FBRM-G400-Mettler Toledo, range of 0.1 to 1000 µm) was located in the reactor in 137 order to measure the distribution of particle chords.

138 **2.2** Substrates and enzyme

139 Paper pulp from coniferous wood (Softwood, obtained via the Kraft process, with pulp 140 extracted before bleaching, Tembec Co., Saint-Gaudens, France, type FPP27) after 141 extrusion (Extruder Eurolab 16, 400 mm failure, extrusion line: 25 L/D 18/25 142 conveying, 7/25 shear stress) was selected for study. The humidity of this substrate was 143 72%. Per unit dry weight it contained 82% cellulose, 8% hemicellulose, 2% lignin, the 144 remainder being ashes (3%) and extractive fractions (2-4%) whose composition has not been determined. The substrate density is 1034±9 (kg.m⁻³) and the mean volume 145 146 dimeter of particles is 497 ± 77 (µm). This pulp is favourable for enzymatic hydrolysis 147 because of its low lignin content (2%). It also contains a low proportion of 148 hemicellulose (xylan, mannan, etc.).

An enzyme cocktail (ACCELLERASE[®] 1500 Genencor, USA, ref. 3015155108)
containing exoglucanases, endoglucanases (2200 to 2800 CMC U/g), hemicellulases

and β -glucosidases (525 to 775 pNPG U/g) was used. Its optimal temperature and pH were 50 °C (range 50 to 65 °C) and 4.8 (range 4 to 5), respectively. An ACCELLERASE® 1500 dosage rate of 0.1 to 0.5 mL per gram of cellulose or roughly 0.05 to 0.25 mL per gram of biomass (depending on the biomass composition) was recommended by the suppliers. Inactivation may occur at temperatures higher than 70 °C and for pH < 4 or pH > 7. Enzyme activities were characterised in the range of 50 to 60 FPU/mL as reported [11, 12].

158 2.3 Physical analysis

Lignocellulose suspensions demonstrate complex rheological behaviour, and there is no standard method for studying their flow behaviour. To characterise their rheological properties as finely as possible, two measurement strategies were combined: (i) ex-situ rheometry (oscillation mode), which provided yield stress and elasticity information, and (ii) in-situ viscometry, which followed suspension viscosity in real time during enzyme attack. Particle size was analysed by in-situ chord length measurement (FBRM).

166 2.3.1 Ex-situ rheometry

167 Classic rheometry based on continuous permanent shear rate appears irrelevant for 168 lignocellulosic substrates because of the rapid aggregation of fibres [10]. Thus, 169 oscillatory measurements (including strain sweeps and frequency sweeps) were 170 performed. These measurements offer several advantages: firstly, they prevent fibre 171 aggregation caused by constant unidirectional shear flow; secondly, they provide 172 additional information on the rheological behaviour of the suspensions (yield stress in 173 this case). The storage modulus G' and the loss modulus G'' were measured using a 174 Mars III rheometer (Thermo Scientific). Dynamic measurements were performed with 175 serrated plates (60 mm, roughness 400 µm, gap size: 1.5 mm) on samples taken during hydrolysis. First, an oscillatory shear flow was set up with increasing shear stress 176

177 amplitude from 0.1 to 20 Pa and at a fixed frequency (1 Hz). This first measurement 178 was used to determine the linear domain. Then a scan was carried out in the linear 179 domain for frequencies from 0.5 Hz to 20 Hz and fixed shear stress amplitude. The 180 analysis was performed at 20 °C. Several methods can be used for yield stress 181 determination. It can be interpreted as the stress amplitude at which the elastic modulus 182 G' becomes smaller than the shear modulus G", or as the stress amplitude at which the 183 loss modulus G" reaches a maximum. It can also be identified as the maximum elastic 184 stress verifying a linear relation $\tau = G' \cdot \gamma$ where γ is the strain amplitude [13, 14]. 185 Especially for lignocellulose substrates, which cannot stand long measurement times, it 186 was defined as the first departure from the linear viscoelastic region [9, 15]. In the 187 present study, a 20% reduction in G' was chosen.

188 2.3.2 In-situ rheometry

189 Ex-situ measurement was limited by the number of samples and the substrate properties, 190 predominately decantation and flocculation of material. To overcome these difficulties, 191 in-situ viscometry was conducted throughout hydrolysis. It was based on the 192 determination of consumption power (or power number $Np = \frac{P}{d^5 \cdot \rho \cdot N^3}; P = 2\pi \cdot N \cdot C$) versus the Reynolds number (Re = $\frac{\rho \cdot N \cdot d^2}{\mu}$) during 193

suspension mixing (see [16] for details). The viscosity was calculated from the power
consumption curve of the mixing system under consideration using a semi-empirical
model including laminar and transition regions for the reference curve with a one-to-one
relationship between Np and Re:

198
$$Np = \left(\left(\frac{Kp}{Re} \right)^q + N_{p0}^q \right)^{\frac{1}{q}}$$
 with : $N_{p0} = 0.128; q = 0.782$ (Eq. 1)

199 The Kp constant for the mixing system was 97.9, while the Metzner-Otto constant used 200 to estimate an equivalent shear rate was Ks = 32. All the geometric constants were determined from measurements using Newtonian (water, Marcol oil, and glycerol) and non-Newtonian fluids (xanthan-sucrose solutions). Once the experimental set-up was characterised by its power consumption curve Np(Re) and its Ks value, in-situ viscometry of the suspension was performed before the addition of enzymes and then throughout the biocatalytic reaction.

206

2.3.3 In-situ particle size analysis

207 Focus beam reflectance measurement (FBRM) enabled in-situ quantification and 208 characterisation of chord length distribution (CLD). The FBRM sensor (FBRM G400, 209 Mettler-Toledo, range: 0.1 to 1000 µm) was set up in the bioreactor to detect and 210 monitor the changes of particle dimensions during enzymatic hydrolysis in real time. FBRM measurement is a laser-based technique. A solid-state laser light source ($\lambda = 795$ 211 212 nm) provides a continuous beam of monochromatic light that is sent down the FBRM 213 probe. A precision motor—pneumatic or electric—is used to rotate the precision optics at a constant speed. The scan speed is fixed at 2 m.s⁻¹. As the scanning-focused beam 214 215 sweeps across the face of the probe window, individual particles or particle structures 216 (agglomerated or floc) backscatter the laser light towards the probe. Particles and 217 droplets closest to the probe window are located in the scanning focused spot and 218 backscatter distinct pulses of reflected light, which are detected by the probe and 219 translated into chord lengths based on the simple calculation of the scan speed (velocity) 220 multiplied by the pulse width (time). A chord length is simply defined as the straight-221 line distance from one edge of a particle or particle structure to another edge. Typically, 222 thousands of individual chord lengths are measured each second to produce the chord 223 length distribution, which is the fundamental measurement provided by FBRM.

224 **2.4 Biochemical analysis**

225 **2.4.1** Dry matter content

The water content of substrates and hydrolysed suspensions was determined by drying at low temperature and pressure. Volumes of samples (\approx 1 mL) were put in Eppendorf tubes (known mass, m_{epp}). These Eppendorfs (m_{ini}) were then placed in an oven at 60 °C, 200 mbar for 5 days, and afterwards, weighed (m_{fin}). Water content (W) and dry matter content (DM) were calculated using **Eq. 2** (accuracy ±0.5%):

231
$$W(\%) = \frac{m_{ini} - m_{fin}}{m_{ini} - m_{epp}} \cdot 100; DM(\%) = 100 - W (Eq. 2)$$

232 2.4.2 Monomers and oligomers

233 Samples were centrifuged at 13000 rpm for 5 min. Glucose and soluble cello-234 oligosaccharides, (i.e. with a degree of polymerization lower than 6) in the supernatants 235 were quantified by an AMINEX HPX-87P carbohydrate analysis column (Bio-Rad 236 Laboratories, Richmond, CA) using a high-performance liquid chromatography system 237 (separations module: Waters Alliance 2690; refractometer detector: Waters 2414, 238 Milford, MA). Conditions were optimised to detect and quantify the different cello-239 saccharides. Analysis was performed at 60 °C with deionised water and 0.1M Pb(NO₂)₂ 240 as mobile-phase (ratio 80/20 v/v) at a flow rate of 0.5 mL/min for 30 min. Glucose and 241 cello-oligosaccharide standards with a DP ranging from 2 to 5 were used (Sigma 242 Chemical Co., St. Louis, MO).

243 **2.5 Experimental strategy**

Enzyme hydrolysis was carried out at 40 °C due to enzyme heat stability (activity reduction at high temperature), energy conservation, and taking into account the optimal conditions for the cell cultivation step. The pH of the medium was adjusted to 4.8. To prevent microbial contamination, 0.2 mL of a solution of chloramphenicol (5 g.L⁻¹) was 248 added. Enzymes were added when the suspension reached homogeneity (visual 249 monitoring and stabilised torque, reached within 30 min). Hydrolysis was carried out 250 over 24 h with a mixing rate of 100 rpm (corresponding approximately to a shear rate of 251 50 s⁻¹), reactor volume 1300 mL and using three substrate concentrations, 1% (10.1 and 252 9.7 gdm/L formally), 2% (only for in-situ viscometry analysis, 19.6 and 20.0 gdm/L 253 formally) and 3% w/v (29.2 and 28.7 gdm/L formally), two enzyme/substrate ratios, 0.1 254 and 0.5 mL enzyme/g cellulose respectively (corresponding to 5 and 25 FPU/g 255 cellulose). Suspensions were sampled with a 7 mm diameter flexible tube connected to a 50 mL syringe. Each sample was about 15 mL at 0 h, 15min, 1h, 2h, 3h, 5 h, 7h, 10 h 256 257 and 24 h of hydrolysis time. The enzyme reaction was stopped by adding 0.1 mL 10 N 258 NaOH. The total volume of samples removed was approximately 130 mL (< 10% of 259 initial volume). Samples were analysed in terms of rheological, granulo-metric and 260 biochemical properties during enzyme degradation.

261 **3 Results and Discussions**

262 **3.1 Bioconversion rate**

The production of monomers (C6, C5) and water-soluble oligosaccharides is crucial for the development of new intensified bioprocesses. First of all, the bioconversion rate was calculated from the glucose produced using **Eq. 3**.

$$Bioconversion (\%) = \frac{[Glc]_{measured}}{[Glc]_{theory}}$$
$$[Glc]_{theory} = \left(m_s \cdot \frac{DM}{100} \cdot \frac{Cellu}{100 \cdot 0.9}\right) / \left(V_w + \frac{m_s}{\rho_s}\right) (Eq. 3)$$

267 The theoretical conversion coefficient from cellulose to glucose is 1/0.9.

268 Under enzyme activity, the polymeric cellulose chain is broken down to produce 269 monomers. Oligomers (DP > 2) were not detected for any of the experiments. Xylose 270 was quantifiable only in the case of 3% w/v - 0.5mL enzyme/g cellulose. 271 As expected, the concentration of monomers (glucose, xylose) increases with hydrolysis 272 time. For example, in the 3% w/v cases, the monomer concentration increased regularly 273 during hydrolysis, so that enzyme loading had an impact on the quantity of the glucose released: 5.1 to 19.1 g.L⁻¹, which corresponds to 19 and 73% bioconversion for 0.1 and 274 275 0.5 mL/g cellulose respectively (**Table 1**). Few articles have examined the evolution of 276 water-soluble cello-oligosaccharides. Sun and Cheng [17] analysed the hydrolysis of 277 microcrystalline cellulose (10 g.L⁻¹) by cellulase produced by *Cellulomonas fimi*. They 278 did not detect soluble oligomers with DP>4, but the cellotriose concentration varied 279 between 0.2 and 0.6 g.L⁻¹ depending on the enzyme used-endo-glucanase or 280 cellobiohydrolase. Solubilisation reached 61% and 50% respectively for each enzyme 281 separately. In contrast, the intermediates of cellulose hydrolysis were not found because the Accellerase 1500 cocktail contains all the types of activity required to rapidly 282 283 degrade these intermediates during hydrolysis.

The bioconversion yields are comparable to data reported in the literature. For a 2% (w/w) suspension of oven-dried corncob, after 24 h hydrolysis the glucose conversion varied between 30 and 82% for 6 and 30 FPU/g [18]. With the hydrolysis of a pretreated wheat straw suspension at 1% (w/v), a doubled bioconversion (from 45.9 to 87%) after 18 h with enzyme loading between 9.6 and 57.6 FPU/g was reported [19].

289 Changes in dry matter content were monitored by determining the water content in the 290 sample at different hydrolysis times. This technique involves the error inherent to 291 sampling heterogeneous suspensions. Beside this technique, the dry matter content was 292 also determined from dissolved and undissolved substrate measurements. According to 293 the hypothesis of the conservation of substrate mass before and after hydrolysis, the dry 294 matter in suspension can be deduced from the initial quantity and the hydrolysed 295 quantity (which released soluble components-monomers and cello-oligosaccharides). 296 As previously reported, the water-soluble cello-oligosaccharides with DP > 2 were not 297 quantifiable, so only glucose, xylose, and cellobiose were used to calculate the quantity

of hydrolysed substrate. Mass balance was conserved around 95% (**Table 1**).

- 299 **3.2** Rheological behaviour during hydrolysis
- 300 3.2.1 Viscosity and yield stress

301 In-situ viscometry showed that substrate suspensions were non-Newtonian shear-302 thinning fluids. The viscosity of suspensions at 100 rpm (corresponding to an equivalent shear rate of 50 s⁻¹) as a function of hydrolysis progress is illustrated in **Figure 3-A**. As 303 304 expected, slurry viscosity decreased during hydrolysis. Excluding the very beginning of 305 the reaction (t < 0.2 h), viscosity changed drastically at the beginning of hydrolysis. 306 This sharp reduction in viscosity was observed whatever the concentration or 307 enzyme/substrate ratio. For example, at 3% dm (w/v), 0.5 mL E/g cellulose, the 308 viscosity decreased from 68 to 3 mPa.s. This change in the physical appearance of the 309 slurry is associated with the biochemical changes and particle size changes occurring in 310 the fibres [20, 21]. The drop in viscosity is a combination of the decrease of solid 311 concentration (solubilisation) and of the fragmentation of cellulose fibres [22, 23]. 312 Under the effects of enzymes, the cellulose chains are broken up to produce smaller 313 particles and non-dissolved cellulose is converted into soluble compounds such as 314 monomers and water-soluble oligomers. In addition, the viscosity drop is suggested to 315 be strongly connected with the degradation and decrease in water binding capacity of 316 the lignocellulose matrices during enzyme-based hydrolysis [24].

The initial viscosities of the 3% (w/v) suspensions were 14-fold higher than those of the 1% (w/v) suspension (~70 and ~5 mPa.s, respectively). The nonlinear dependence of viscosity on the concentration is expected and can be explained by an increase in particle interactions, less free water, and hydrogen bonding between cellulose chains. As the solid concentration increases, the average distance between particles in the slurry decreases, leading to enhanced contact between particles, especially if there is an 323 entanglement of amorphous fibres between particles [25]. The high increase in initial 324 viscosity of between 1% (w/v) and 3% (w/v) signifies that the increase of substrate concentration becomes the major stress for enzymatic hydrolysis at high (and very high) 325 326 dry matter content levels. That is why a special strategy to reach the condition of high 327 substrate concentration is absolutely necessary. During hydrolysis, a significant drop in 328 slurry viscosity is observed within the first 10 h of the hydrolysis reaction. These results 329 are supported by the literature for a wide range of matrices, particle sizes, and 330 enzyme/cellulose ratios [26-28], although a large heterogeneity of viscosity data is 331 reported. This can be explained by different substrates, pre-treatments, morpho-332 granulometry and enzymes, and operating conditions.

333 The effect of enzyme concentration on viscosity is clearly demonstrated even for the 334 lowest concentration in **Figure 3-A**. For the same substrate concentration, the higher the 335 enzyme/g cellulose ratio (E/C), the faster the viscosities decrease. Within 5 h, the 336 viscosity of a 3% (w/v) suspension was reduced by 85% for 0.5 mL E/g cellulose; 337 whereas its reduction was limited to 30% for 0.1 mL E/g cellulose. This observation is 338 in agreement with other authors [22, 26, 29]. Furthermore, two tendencies in viscosity 339 evolution were found for suspensions containing 0.1 mL/g cellulose. Firstly, the 340 suspension viscosity increased during the first hour consecutive to swelling and 341 unbinding effects (clearly observed for 2% w/v and 3% w/v, not significant for 1% 342 w/v); after that, the viscosity decreased because of depolymerisation by the enzymes.

343 Shear stress sweep and frequency sweep experiments were performed in oscillation 344 mode (see §2.3.1). The frequency sweep revealed that the elastic modulus G' and the 345 viscous modulus G'' do not depend on frequency (**Figure 3-B**). In addition, G' is 346 always greater than G'', with a ratio G'/G'' which can be identified in Figure 3-B. This 347 is characteristic of viscoplastic behaviour, and shear stress sweeps were then used to 348 evaluate yield stress. All suspensions showed viscoplastic shear-thinning behaviour 349 both initially and throughout hydrolysis. Shear-thinning has already been mentioned by

350 other authors [9, 27, 29]. Before the introduction of enzyme, the rheological behaviour 351 for small deformations is mainly elastic, with a high value of G' and a ratio G'/G'' close to 5. For a 3% (w/v) suspension at t = 0 h and in the linear domain, G' and G'' reached 352 353 about 1000 and 200 Pa. This is supported by the literature for different matrices. For 354 corn stover, at 12% dm, G' and G" were reported as 2000 and 300 Pa [15]; for acid pretreated softwood, these values were 100 and 50 Pa respectively [9]. When increasing the 355 356 shear stress, all the samples exhibited two zones: a first where G' and G" did not depend 357 on shear stress (linear domain) and a second where both moduli decreased. In this 358 second zone, G' deviated sharply from a stable curve and crossed (or tended to cross) 359 G". During hydrolysis, a regular decrease in both moduli was observed. For example, in 360 the 3%-0.5 case, a 1000-fold reduction was observed for both G' and G" after 24 h of 361 enzyme attack. Moreover, the elastic character is noticeably preserved. One also notes 362 that the higher the enzyme activity, the faster the decrease in elasticity (in agreement 363 with the viscosity results above). This discussion highlights, yet again, the strong impact 364 of enzyme concentration on how rheological behaviour varies during hydrolysis.

365 As viscometric yield stress measurements were not possible for these suspensions, this 366 yield stress τ_0 was deduced from the elastic modulus (G') (see Figure 3-B). Yield stress 367 can be regarded as the stress required to initiate flow. Yield stress values between 1 and 368 20 Pa were determined at 1% and 3% dm (w/v) before hydrolysis. With an increase in 369 substrate concentration from 1% to 3% dm (w/v), yield stress increased 20-fold (Figure 370 4). This confirmed the critical substrate concentration ($\approx 3\%$ dm (w/v) determined by in-371 situ measurements) beyond which the viscosity increased exponentially [16]. The 372 results presented were slightly higher than those obtained from pre-treated softwood: 0-373 28 Pa for 4-12% substrate concentration [9] and from pre-treated corn stover: 0.26-22.9 374 Pa for 5-17% dm [30]. This difference could be due to the nature and the physical 375 characteristics of the matrices and to the method used to determine yield stress.

376 During hydrolysis, yield stress decreased sharply as enzyme attack progressed (Figure. 377 4). This can be explained by the drop in solid content in suspension and by a reduction 378 in fibre-fibre interactions due to hydrolysis progress. The decrease of yield stress was 379 greater for experiments with higher enzyme concentrations. The final yield stress (at 24 380 h hydrolysis) of 3%-0.1 was 10 times higher than that of 3%-0.5. This observation is 381 directly correlated to the impact of enzyme activities on fibre degradation. In addition, 382 the yield stress of the hydrolysed samples (3%-0.5 at 24 h, DM = 1.14%) exhibited 383 lower values for similar DM content compared to the original material (yield stress at 0 384 h of 1% - 0.5, DM = 0.97%). This can be explained by the modification of fibre 385 structure, diameter, and shape. Decreasing yield stress during enzyme hydrolysis was 386 previously reported for corn stover [21] and pre-treated softwood [9]. These two studies 387 also reported a lower yield stress value for hydrolysed slurries when compared to non-388 hydrolysed slurries at the same solid content.

389 Figure 4 shows the dependence of yield stress during hydrolysis presented in terms of 390 dry matter content. Taking the criterion that the fluid behaves as a pourable liquid at a 391 yield stress below 1 Pa (the values of 1% are negligible), interestingly, results for 3% 392 show that the two yield stress curves collapse onto a single curve when plotted against 393 dry matter content. It decreased exponentially with the decrease in dry matter content 394 during hydrolysis. Yield stress becomes negligible when solid matter < 2%. This has 395 real significance for the choice of substrate flow rate for the cumulative feed strategy or 396 if the slurry has to be pumped into another bioreactor.

397 **3.2.2** Uniqueness of viscosity-time curve

In order to explore the change in viscosity during enzyme hydrolysis and to compare its
kinetics between different experimental conditions, a critical time is estimated through a
normalised viscosity defined as follows:

401
$$\mu^* = \frac{\mu_t - \mu_{fin}}{\mu_0 - \mu_{fin}} (Eq. 4)$$

402 μ^* is a dimensionless viscosity; and μ_0 , μ_t , μ_{fin} are the viscosities at t = 0 h, t_i and the 403 final viscosity (the viscosity of a suspension containing the whole dry matter as soluble 404 fractions – hypothesizing total conversion).

405 μ^* describes the reduction of viscosity during enzyme hydrolysis. It starts from 1 (t = 0 h) and tends to 0 (t = 24 h). From this quantity, a critical time noted $t(\mu^* = x)$ can be 406 407 defined as the time at which the viscosity was x% of the initial suspension viscosity. 408 The relationship between these critical times $t(\mu^* = x)$ and dimensionless viscosity, μ^* 409 only depends on the enzyme concentration used regardless of the substrate 410 concentration. These critical times decreased exponentially with the dimensionless 411 viscosity. The ratio of these critical times between both conditions was around 4-fold. 412 Thus the hydrolysis time, t, can be normalised using a selected critical time $t(\mu^* = x_0)$ 413 (Eq. 5):

414
$$t^* = \frac{t}{t(\mu^* = x_0)}$$
 (Eq. 5)

The dimensionless time-viscosity curves are plotted in Figure 5 ($\mu^* = 0.25$). 415 416 Interestingly, a single curve is exhibited for $t^* > 0.05$, i.e. when the possible swelling 417 and unbinding effects can be neglected. This uniqueness of dimensionless time-418 viscosity curves was observed whatever the hydrolysis conditions in the studied cases 419 (substrate concentrations, enzyme ratio). These results suggest that during the period 420 and for a 75% reduction of initial viscosity, a similar degradation mechanism could be 421 assumed in semidilute regime ($\leq 3\%$ dm w/v), although rheological behaviour strongly 422 differs between 1% and 3% w/v suspensions.

The uniqueness of viscosity-time curves at macroscopic observation suggests a similar
mechanism of fibre degradation. Considering these in-situ and ex-situ rheological

results, macro scales will be explored with chord length distribution analysis, and thenkinetics will be modelled.

427 **3.3 Particle size evolution**

The time evolution of the mean chord length and chord number for the four different conditions is shown in **Figure 6**. Considering chord length distribution (CLD) for the initial suspension, we define four classes (I: < 60 μ m, II: 60 to 90, III: 90 to 160 μ m, IV: 160 to 600 μ m) which correspond to 25% of this population. **Figure 7** illustrates the evolution of each class during hydrolysis for the case 1% w/w - 0.5 mL enzymes/g cellulose.

434 A sharp decrease in mean chord length from 120 to 60 µm (Figure 6) is observed: 435 within two hours for E/S = 0.5 mL/g and six hours for 0.1 mL/g. Considering chord 436 number, initial values are almost proportional to concentration and during hydrolysis, 437 two successive trends are observed. During the first step, the total chord number 438 increased for all experiments, although the durations of this step differed strongly: 439 around 8 to 10 hours for E/S = 0.1 mL/g, and 1 to 3 hours for E/S = 0.5 mL/g. 440 Meantime, the relative increase of chord number is lower for the largest E/S ratio. 441 Considering now Figure 7, coarse population (class IV) decreased regularly while the 442 finest population (Class I) increased.

The initial augmentation of chord number can be explained by the fragmentation of cellulose fibres. Coarse particles are attacked and divided into several fine particles. Beyond that, enzyme activity occurs and the finest particles are converted into dissolved compounds, generating a reduction in chord number. As demonstrated in **Figure 6**, the fragmentation mechanisms appear to be the dominant effect in the strong viscosity reduction observed during the first 5 h, while solubilisation increased to 31%. In contrast, after 5 h, fragmentation is negligible (class IV almost constant) while

450 solubilisation increased to 76% (at 24 h). Correlatively, the viscosity evolution was
451 limited and the finest populations still increased from 38% to 50%.

452 **3.4** Modelling kinetics at macro-, micro-, and biochemical scales

453 Hydrolysis induces a reduction in particle size, dry matter content, and viscosity, which 454 promotes mixing and fibre accessibility. The kinetics of viscosity (macro), chord length 455 (micro) and substrate and product concentrations (biochemical) stand as key indicators 456 for understanding and controlling bioprocess performance. The kinetics of biochemical 457 enzymatic reactions have been extensively reported in the literature [31], contrary to 458 physical parameters such as viscosity and granulometry. A lot of enzyme reactions (e.g. 459 hydrolysis, oxidation, and reduction or cofactor mechanisms) are second- or higher 460 order reversible reactions. Many are limited by diffusion and physical accessibility. 461 Adsorption (reaction) in high molecular weight structures (for example, proteins, 462 polynucleotides, polysaccharides, or heterogeneous protein-phospholipid, protein-463 nucleotide, and protein-polysaccharide structures) is more complex. Among the 464 numerous models reported in the literature, one of the best known is the Michealis-465 Menten equation [32] but a first-order model was also reported by Chrastil [33].

466 The time evolution of viscosity μ , mean chord length lc_m, and residual substrate 467 concentration in dry matter S (%) are modelled by the general equation **Eq. 6**:

$$468 \qquad -\frac{dX}{dt} = k \cdot X^{\alpha} \ (Eq. \ 6)$$

469 where X is the variable being modelled (μ , lc_m and S); k is the kinetic constant, and α 470 the model/reaction order (/). As boundary conditions, X varies from X₀ to X_{∞}, 471 corresponding to initial and final values respectively.

472 A second-order model accurately describes the time dependence of the physico473 chemical parameters—viscosity, chord length, and dry matter content. For the lowest
474 enzyme ratio, an increase of the mean chord length was observed during the first hours.

In these cases, the proposed model was restricted to the time interval corresponding to the maximal value of lc_m until the end of hydrolysis (experiments with 0.1 mL enzyme/g cellulose). This lc_m increase step could be explained by fibre swelling and unwinding due to limited enzyme activity. To sum up, the most suitable kinetic models are second order and written as **Eq. 7 to 9**:

480
$$\mu = \frac{\mu_0 - \mu_\infty}{(\mu_0 - \mu_\infty).k_\mu.t + 1} + \mu_\infty (Eq. 7)$$

481
$$lc_m = \frac{lc_{m0} - lc_{m\infty}}{(lc_{m0} - lc_{m\infty}).k_{lc}.t + 1} + lc_{m\infty}$$
 (Eq. 8)

482
$$S = \frac{S_0 - S_\infty}{(S_0 - S_\infty) \cdot k_S \cdot t + 1} + S_\infty (Eq. 9)$$

483 The model was adjusted via the least squares method for each of the physico-484 biochemical parameters. The coefficients and the correlation coefficients are presented 485 in Table 2. The correlation coefficients indicated the fine agreement between the 486 models and experimental data. The enzyme cocktail (Accellerase 1500) contained only 487 cellulases and hemicellulases, so the lignin fraction can be considered as contributing to 488 the non-hydrolysable fraction S_{∞} . For our substrate, lignin represents less than 2%, and 489 thus can be neglected. The final viscosity μ_{∞} corresponds to the viscosity of a 490 suspension containing all soluble fractions (total conversion). In the present case, this 491 value would be close to the supernatant viscosity (0.7 mPa.s) and μ_{∞} is negligible 492 compared to the initial suspension viscosity. In ideal conditions, the solid fractions 493 should be converted into soluble fractions and the ultimate chord length $lc_{m\infty}$ would then 494 be null. However, experimentally, there is always some part that remains non-495 hydrolysable substrate, so $lc_{m\infty}$ cannot be neglected compared to lc_{m0} , and is assumed 496 equal to lc_{m24h} .

497 Effects of enzyme and substrate concentrations are clearly observed for all constants.
498 With k_s, the increase of the E/S ratio from 0.1 to 0.5 mL/g cellulose led to a 7- to 9-fold

499 increase. When the initial substrate concentration increased from 1 to 3% w/v, a 500 reduction of 2-3 times was observed. The same tendency was found with k_{μ} , which 501 demonstrated the strong impact of enzyme and substrate concentrations on the variation 502 of suspension viscosity during hydrolysis. Similar impacts were observed for enzyme 503 concentration and substrate concentration: 8-fold and 4-fold respectively. The final 504 value of the mean chord length, $lc_{m\infty}$, seemed identical in magnitude; however, the 505 impact of enzyme ratio can be clearly distinguished by the absolute values of k_{lc}. An 506 increase of 3.5-5.5 -fold of k_{lc} was observed when the enzyme concentration passed 507 from 0.1 to 0.5 mL/g cellulose.

508 Using equations 7 to 9, Figure 8 illustrates the successive phenomenological limitations 509 observed during hydrolysis. Rheological behaviour appears to be the primary limiting 510 factor. Transfer limitation due to high viscosity and yield stress is, however, temporary 511 (during the first 5 h) but constitutes an important phenomenon. Viscosity collapse and 512 particle fragmentation are concomitant. Beyond this, the limited reduction in size 513 indicates a threshold for biocatalytic reaction. This could be due to recalcitrant fractions 514 or inefficient enzyme activities. It is therefore necessary to study the physical and 515 biochemical structures of this recalcitrant fraction in order to identify how it could be 516 degraded.

The major challenge in 2nd-generation biofuels is to reduce costs so as to compete with 1st-generation ones. Therefore, processing at high solid content is mandatory. However, the rheological behaviour of the hydrolysis suspension stands out as the first and major determinant of process efficiency, and has to be considered a key criterion in proposing a rational strategy for reaching high dry matter content.

522 To increase transfers and bioreaction efficiency, a strategy can be built from the 523 identified parameters in a semidilute regime: the critical substrate concentration C^* 524 (meaning a drastic increase of initial suspension viscosity beyond C^*) and a targeted

525 hydrolysis time, $t(\mu^* = x_0)$ (corresponding to a chosen relative reduction of initial 526 viscosity), can be used to define a reference feed rate Qc (**Eq. 10**):

527
$$Q_c = \frac{C^*.V}{t(\mu^* = x_0)} (g.h^{-1}) (Eq. 10)$$

A cumulative feeding strategy can be defined, based on Qc, which would allow working
in a favourable regime so as to reach high hydrolysis yield by avoiding instantaneous
high substrate concentrations.

532 4 Conclusion

This study aimed to delve deeply into the biophysical and transfer limitations occurring during enzymatic hydrolysis in the cellulosic biofuels context. Considering paper pulp substrates, semidilute conditions were chosen, as they introduce the complexity of particle-particle interactions, which are strongly involved in transfer limitations, without inhibiting the bioreaction. Biochemical and physical phenomena were explored through a specific experimental set-up associating in-situ and ex-situ analyses.

539 Non-Newtonian behaviour associated with non-negligible yield stress stand as the major 540 factors limiting process efficiency and progress towards high solid loading. The 541 uniqueness of dimensionless time-viscosity curves was observed whatever the operating 542 conditions, and suggests a similar mechanism for fibre degradation. The evolution of 543 biophysical parameters during hydrolysis was observed as the results of a combination 544 of fibre fragmentation, which dominated during the first step, and solid solubilisation. 545 Kinetics modelling enabled demonstration of the successive phenomenological 546 limitations and their magnitude. All these elements lead to the proposal of a reference 547 feed rate which would be used in a cumulative feeding strategy for reaching high solid 548 loading and balancing energy consumption and process efficiency.

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- 645
- 646

Figure 1: Experimental methodology and strategy

Figure 2: Configuration of bioreactor and impellers (dimensions in mm)

Figure 3: Overview of in-situ and ex-situ rheometry. A: In-situ viscometry as a function of hydrolysis time. B: Viscous, G" (open symbols), and elastic, G' (filled symbols), as a function of shear stress (3%-0.5 signifies 3% dm and 0.5 mL enzyme/g cellulose).

Figure 4: Yield stress during hydrolysis versus dry matter content (1%-0.5 stands for: 1% dm and 0.5 mL enzyme/g cellulose)

Figure 5: Uniqueness of dimensionless viscosity-time curves (1%-0.5 signifies 1% dm and 0.5 mL enzyme/g cellulose).

Figure 6: Overview of in-situ (FBRM) particle size analysis: A: Chord number and B: Mean chord.

Figure 7: Population balance (classes defined from initial chord length distribution), in-situ viscosity, and hydrolysis yield during hydrolysis (1%-0.5 mL enzyme/g cellulose case).

Figure 8: Reduction of physico-biochemical parameters during hydrolysis using kinetic models (for 3% w/v, 0.5 mL enzyme/g cellulose)







Agitator **Ø**73 Height: 38



Agitator ϕ 120 Height: 22













Substrate concentration	Enzyme/Substrate	0h	0.25h	2h	5h	10h	24h	Mass balance
(%gdm.L ⁻¹)	(mL/g cellulose)							(24h,%)
1	0.1	0	1.0	4.7	9.6	11.6	13.2	93.7
1	0.5	0	6. <mark>0</mark>	19. <mark>6</mark>	31. <mark>1</mark>	52 .4	76. <mark>6</mark>	94.1
3	0.1	0	1.3	6. <mark>4</mark>	11 .3	14.5	19.1	94.6
3	0.5	0	6. <mark>0</mark>	20 <mark>.6</mark>	32.7	54 . 1	73. <mark>5</mark>	95.2
1 3 3	0.5 0.1 0.5	0 0 0	6.01.36.0	19.6 6.4 20.6	31.111.332.7	52.4 14.5 54.1	76.6 19.1 73.5	94.1 94.6 95.2

Table 1: Bioconversion rate (%) during enzyme hydrolysis

Coefficient	1%	w/v	3%w/v		
_	0.1mL/g	0.5mL/g	0.1mL/g	0.5mL/g	
	cellulose	cellulose	cellulose	cellulose	
k_{μ} (Pa ⁻¹ .s ⁻²)	-	61.5	2.1	15.3	
R ²	-	0.936	0.979	0.988	
$k_{lc} (\mu m^{-1}.s^{-1})$	8.1x10 ⁻³	44.7x10 ⁻³	9.0x10 ⁻³	32.1x10 ⁻³	
	(65 µm)	(66 µm)	(60 µm)	(59 µm)	
R ²	0.875	0.876	0.950	0.938	
k_{S} ((gdm/L) ⁻¹ .s ⁻¹)	7.6x10 ⁻³	71.7x10 ⁻³	3.7x10 ⁻³	26.9x10 ⁻³	
R ²	0.822	0.987	0.885	0.995	

Table 2: Kinetic coefficients of rheological, granulometric ($lc_{m\infty}$ in μm) and biochemical parameters, and associated correlation coefficients R^2 .