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Cicely Warne, Sami Fadlallah, Florent Allais, Georg Guebitz, Alessandro Pellis. Controlled Enzymatic Synthesis of Polyesters Based on a Cellulose-Derived Triol Monomer: A Design of Experiment Approach. ChemSusChem, 2024, 17 (11), pp.e202301841. 10.1002/cssc.202301841. hal-04652003

HAL Id: hal-04652003 https://agroparistech.hal.science/hal-04652003v1

Submitted on 17 Jul2024

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Controlled Enzymatic Synthesis of Polyesters Based on a Cellulose-Derived Triol Monomer: A Design of Experiment Approach

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18 Abstract

Regioselective enzymatic polycondensation of the bio-based cellulose derived polyol, 19 Triol-citro, and dimethyl adipate using Candida antarctica Lipase B (CaLB) was 20 investigated. A Design of Experiment approach with MODDE® Pro 13 was used to 21 determine important factors in the branching behavior of this polymer, and reactant 22 ratio, temperature, reaction time and enzyme wt% were the studied factors. 23 24 Multifunctional polyesters with pendant hydroxy groups were synthesized and fully characterized using 2D NMR techniques to determine degree of branching. 25 Branching was minimal, with a maximum of 16% observed, and monomer ratio, 26 temperature and reaction time were all determined to be significant factors. In this 27 28 work, M_n of up to 13 kDa were achieved, while maintaining degree of branching below 15%, resulting in a linear polyester with the potential to be further 29 30 functionalized.

31 Introduction

Since the concept first emerged in the 1990s, the drive to incorporate Green Chemistry into 32 all aspects of the chemical industry has steadily increased. The need for sustainability 33 requires researchers to look away from fossil resources and their growing problematic 34 utilization, and instead turn to biomass feedstock. One platform molecule that has risen to 35 prominence in recent years is levoglucosenone (LGO). It can be synthesized from waste 36 cellulose, and there are many publications focusing on more sustainable production of this 37 chemical.^[1-4] This chiral molecule containing both a cyclic acetal and an α , β -unsaturated 38 ketone has increasingly been used as a starting feedstock for solvents,^[5,6] specialty drugs,^[7,8] 39 and polymers.^[9-12] A recent example involves the development of a green one-pot pathway 40 to synthesise a multifunctional triol monomer (Triol-citro) from LGO.^[13] This monomer 41 features a photocrosslinkable citronellol side group and has been employed in polymer 42 synthesis (Scheme 1.).^[13–15] 43





Triol compounds are particularly interesting as there is the potential to synthesise polymers 47 with pendant hydroxy groups. Such polymers often have increased hydrophilicity,^[16] which 48 can be useful in drug delivery^[17] or membrane applications.^[18] The presence of reactive 49 hydroxy groups also opens the polymer to functionalisation for further tailoring of their properties. Quaternary ammonium compounds,^[19] proteins^[20] and antibodies,^[21] have all 50 51 been grafted to a polymer backbone through reaction with an available functional group. 52 53 Linear polyesters can be converted to crosslinked structures to improve mechanical properties.^[22,23] and these crosslinks can even be reversible to maintain processability.^[24] 54 While functional groups can be introduced at the post-polymerisation stage through surface 55 modification techniques such as enzymatic^[25] or chemical^[26] hydrolysis, as well as physical methods such as plasma treatment,^[27] it is important to note that these methods may lead to 56 57 a reduced molecular weight polymer. Directly polymerizing a trifunctional monomer 58 59 eliminates this drawback, but to avoid branching resulting from the reaction of all three hydroxy groups, protecting groups or a selective mode of catalysis must be employed. 60

Lipases are an example of such catalysts, as they catalyze (trans)esterification reactions and often display regioselectivity in their choice of substrate.^[28] Biocatalysts are considered more sustainable compared to more traditional organic catalysts; they are safe, energy efficient and minimize the use of toxic reagents. The enzyme *Candida antarctica* Lipase B (CaLB), due to its *sn*-1,3 regioselectivity, has been used to synthesise polymers based on glycerol,^[29– 43] sorbitol^[36,43–49] and other bio-based triols.^[45,48] By changing different factors and controlling the degree of branching; crosslinked, linear and dendritic polymers can be produced enzymatically.

Overall, the controlled synthesis of linear polyesters using triol monomers, with a focus on directing the reaction towards two of the three hydroxy groups to prevent or minimize branching, is of great interest, but most research focuses on glycerol and sugar-derived compounds. The advantage of a monomer such as Triol-citro is twofold. Firstly, the presence of a second (alkene) functional group allows for more than one functionalisation of the polymer. Alternatively, this citronellol side chain can also be photocrosslinked, as shown in Triol-citro based polycarbonates.^[14] Secondly, the synthesis of this molecule is theoretically extremely versatile. The oxa-Michael addition could proceed with a wide range of alkene
 alcohols, not only citronellol. There are therefore many potential triol based compounds that
 can be synthesized from LGO using this method.

Triol-citro based polyesters have been synthesized.^[13] but use of diacyl chloride reagents in 79 approaches resulted in insoluble polymers, due to the formation of 80 previous branched/crosslinked structures. In addition, from a green chemistry perspective, diacyl 81 chloride poses notable hazards due to its highly reactive and corrosive nature. In contrast, 82 exploring enzymatic approaches, such as utilizing lipase CaLB with diesters, provides a 83 greener and safer alternative for polymer synthesis. Here, the aim is to use CaLB to 84 synthesise linear polyesters from the monomers Triol-citro and dimethyl adipate (DMA) 85 (Scheme 2). A Design of Experiment (DoE) approach was used to investigate the influence 86 of several factors on polymer molecular weight, conversion, and the degree of branching, 87 with the intent of producing high molecular weight and preferentially linear polyesters. 88



89

90 **Scheme 2.** CaLB-catalyzed polycondensation reaction of Triol-citro and dimethyl adipate to produce 91 linear polyesters.

92

93 **Results and Discussion**

94 **Polymer synthesis and characterization**

In order to identify the factors that affect branching in the enzymatic polycondensation of polyols, a literature search was conducted. Numerous publications that synthesized polyesters from polyol monomers were found.^[29,31–44,46–49] Subsequently, temperature, enzyme wt%, reaction time and monomer ratio were identified as pertinent factors for investigation in this work (Table 1.). To streamline and optimize the experimental process, a DoE approach was used.

Table 1. Factors that have been shown to affect branching and the observed effect in the enzymatic
 polycondensation of polyols in previous literature.

Factor	Effect on branching
Temperature	Increased or decreased branching depending on solvent. ^[29] Increased branching with increasing temperature. ^[32]
Enzyme wt%	Increased enzyme wt% results in increased branching. ^[29,31]
Reaction time	Increased reaction time results in increased branching. ^[31,39] Increased reaction time increases molecular weight and has no effect on branching. ^[32]
Monomer ratio	Increasing the amount of polyol monomer results in both increased branching ^[31] , and decreased branching. ^[29] Excess (other) monomer results in increased branching. ^[29,35,40]

103 A central composite design model generated 27 experiments (N1 – N27), seen in Table S1. 104 The polymers produced pre-washing were liquids of varying viscosity and degrees of 105 precipitation (see table S2). Conversions and molecular weights were obtained directly from 106 unwashed sample and were determined using ¹H-NMR and GPC analysis, respectively, with 107 Equation (1) used to calculate conversion.





Figure 1. Monomers with labels pertaining to equations (1) and (2).

Determining the degree of branching proved to be challenging. ¹H-NMR analysis clearly showed the presence of unreacted monomer in the obtained products, as along with other impurities (Figures S1 – S27). Attempts were made to wash the crude polymer with MeOH, however a significant portion of the polymer mass consisted of short chain oligomers that were also soluble in MeOH at this concentration. For example, N6 was washed with 2 x 5 mL MeOH and the mass of the polymer obtained was insufficient for ¹³C-NMR and any 2D techniques.

Subsequently, three experiments (N28 - N30) were performed in a longer timeframe in an 117 attempt to increase molecular weight and induce some branching of the polymer, as it has 118 done for other polyol based polyesters in the literature.^[31,39] The conditions of experiment 119 N15 (1:1 monomer ratio, 20 wt% enzyme, 90 °C, 24 h) were selected as this was the highest 120 molecular weight polymer produced, with the reaction time extended from 24 h to 96 h. 121 122 Immediately after the workup, it was evident that the polymer had branched/crosslinked structures to some extent as experiment N28 was mostly insoluble in tetrahydrofuran (THF). 123 Chloroform (CHCl₃) dissolved a larger proportion, leaving a small amount of insoluble residue 124 in the glassware. N29 and N30 were soluble in THF and dimethyl sulfoxide (DMSO) which is 125 attributed to the fact that these polymers were synthesized using a different batch of Triol-126 127 citro monomer which was flash purified using a slightly different method (see ESI). NMR analysis for unwashed N28 was performed in CDCl₃, which produced spectra of sufficient 128 resolution so as to assign peaks. The remaining amount of N28 was washed (2 x 1 mL), and 129 130 an attempt was made to dissolve it in DMSO- d_6 and CDCI₃ for analysis, which was unsuccessful. This is likely due to the washing step having removed a lot of the short linear 131 132 chains and increasing the proportion of branched/crosslinked structures present.

Observing the ¹H-NMR spectrum for unwashed N28 (Figure 2.), there is a considerable 133 decrease in signal intensity from 3.2 - 3.6 ppm and the formation of new signals from 3.7 -134 5.0 ppm are observed when compared to that of the Triol-citro monomer. These new signals 135 relate to the formation of ester groups, and the corresponding shift of the signals. To 136 definitively assign signals in the ¹H-NMR spectrum, ¹³C-NMR analysis was also performed, 137 as well as HSQC, HMBC and COSY (Figure 2. and Figures S28 - 29). From the HSQC, 138 HMBC and COSY, the majority of peaks could be definitively assigned to protons in the 139 polymer structure. It was determined that Triol-citro protons 2, 4 and 7 (Figure 2.A.), if 140 reacted, will show a shift towards a higher ppm on the ¹H-NMR spectra, but not necessarily 141 in the ¹³C-NMR one. Reacted forms of 2, 4 and 7 are noted as 2', 4' and 7'. When reacted, 142 proton 2 shows a shift from 4.07 ppm to 4.22 ppm on ¹H, while the ¹³C peak stays at 66 ppm. 143 Proton 7 shows a shift from 3.76 ppm to 4.22 ppm on ¹H, and the ¹³C stays at 61 ppm. 144 However, proton 4 shows a shift from 3.76 ppm to 4.94 ppm on ¹H spectra, and also shows a 145

shift in the ¹³C spectra, from 71 ppm to around 77 ppm. It is difficult to determine exactly where the ¹³C peak falls for 4', as peaks in this region overlap with the CDCl₃ signals; both signals for carbons 4' and 9 are expected to fall in this area. DMSO- d_6 was therefore used as an NMR solvent for samples N29 and N30.

All ¹H spectra have the peak characteristic of the methyl groups of DMA (3.56 ppm in DMSO-150 d_6), meaning unreacted Triol-citro will also likely be present. Furthermore, many experiments 151 152 were run with an excess of Triol-citro (Table S1), and this unreacted monomer is expected to contribute significantly to the signals for protons 2 and 7. While substituted terminal units 153 may also be responsible for 2', 4' and 7' proton signals, the presence of the 4 substituted 154 155 terminal unit (as shown in Figure 2.A.) was deemed improbable due to enzyme regioselectivity. In previous works investigating CaLB synthesized glycerol-based polyesters, 156 glycerol units that had reacted only on the secondary hydroxy were a very small proportion of 157 polymer^[29,31], and observed all.^[31] 158 the in some cases not at



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Figure 2.A. Labelled Triol-citro and possible linear, dendritic and terminal structures of the Triol-citro unit within the polyester, with atom labelling of the 2,7 linear unit, **B.** ¹H-NMR spectra of N28 in CDCl₃ with peaks fully assigned, **C.** ¹³C-NMR spectra of N28 in CDCl₃ with peaks fully assigned, and **D.** HSQC spectra of N28 in CDCl₃ with peaks for 2, 2', 4, 4', 7 and 7' highlighted and labelled in red. Peaks labelled with ' are from the reacted form of that atom.

Unwashed N29 and N30 were not characterized, instead they were washed with MeOH, 163 which produced sufficient polymer for comprehensive characterization using NMR. DMSO- d_6 164 was used, resulting in spectra with noticeably fewer impurity peaks and the absence of a 165 sharp peak at 3.56 ppm attributed to the methyl ester group of DMA. Although terminal group 166 analysis of this polymer would be of interest due to its post-polymerisation functionalisation 167 168 potential, this is not possible due to overlapping peaks in this region. Similar to the ¹H spectra of N28, a peak was observed at 4.93 ppm, attributed to 4' (Figure S32). The ¹³C-NMR 169 spectra no longer had obscured peaks in the range of 75 - 80 ppm (Figure S31). The 170 171 positioning of peaks in DMSO-d₆ spectra differed from CDCl₃ spectra slightly, which can be seen when comparing Figure 2.B. and Figure S32. DMSO- d_6 was therefore used as a 172 173 solvent for ¹H-NMR analysis to determine branching of N1 – N27. All 27 spectra can be found in the ESI. 174

Upon observing N1 – N27, a peak at 4.93 - 4.94 ppm is noticeable in the majority of ¹H 175 176 spectra. In many spectra, this peak clearly overlaps with an impurity peak (see Figures S1 -27), consequently, selected samples underwent washing in an attempt to eliminate this 177 issue. Polymers with several different molecular weights were selected (N13, N9 and N27) 178 179 and approx. 100 mg were washed with 2 x 1 mL MeOH. The solubility of the short chain 180 oligomers in MeOH is evident, as these 2 washes resulted in up to 96% mass loss (Table S3). After washing, a white powder was obtained; a contrast to the samples before washing, 181 which were viscous liquids. These samples proved insoluble in DMSO- d_6 , suggesting that 182 183 this solid fraction is the part of the polymer that has crosslinked or branched. This is supported by the fact that N2, a polymer lacking a peak at 4.93 ppm, was fully soluble in 1 184 185 mL of MeOH. Degree of branching was therefore calculated with the spectra of the unwashed polymer using equation (2); with ' denoting the reacted form of the atom (Figure 186 187 1.), and integrals normalized at the CH_3 peak (0.82 ppm).

Branching (%) =
$$\left(\frac{(I_{4'}/N_{4'})}{(I_{4'}/N_{4'}) + (I_{2'+7'}/N_{2'+7'})}\right) \times 100$$
 (2)

188

The results for those polymers discussed in the text are summarized in Table 2., and a more 189 complete table can be found in the ESI (Table S1). Overall, conversion was consistently high 190 (always >68%), and degree of branching never exceeded 16%, but there was significant 191 192 variability in molecular weight. Higher molecular weight polymers also exhibited considerably greater dispersity (D), with a highest observed D of 37.3 (polymer N30). Such high 193 dispersities are unusual for polycondensations conducted in bulk, [50] especially for reactions 194 195 N29 and N30, which were run for 96h. This can likely be attributed to a combination of molecular weight and branching behavior of these polymers, as branched structures exhibit a 196 much larger diversity in shape compared to linear structures. Once a structure has branched, 197 198 it has more sites where polycondensation can occur, further increasing molecular weight in the branched chains. There are many examples in the literature of highly disperse branched 199 polymers to the extent that many researchers focus on limiting D in these structures.^[51,52] 200

201	Table 2. Effect of several reaction conditions (highlighted in grey) on the conversion, molecular weight
202	and degree of branching (DB) observed in the synthesis of poly(Triol-citro adipate).

Reaction	Triol/DMA [moles]	Enzyme [wt%]	Temp. [°C]	Reaction time [h]	Conv. [%] ^[a]	М _n [Da] ^[b]	М_w [Da] ^[b]	Đ ^[b]	DP [b]	DB [%] ^[a]
N2	2	5	40	6	68	2,058	4,792	2.33	5.8	0
N9	1	5	40	24	88	735	869	1.18	2.1	13
N13	1	5	90	24	92	960	1,335	1.39	2.7	12
N15	1	20	90	24	92	2,789	8,044	2.88	7.8	16

N27	1.5	12.5	65	18	93	1,161	2,130	1.84	3.3	9	
N28	1	20	90	96	92	-	-	-	-	13	
N29	1	20	90	96	100 ^[c]	13,300	157,400	11.82	37.4	13	
N30	1	20	90	96	100 ^[c]	13,000	484,500	37.3	36.4	12	
^[a] Determined through ¹ H NMR											

^[b] Determined through GPC analysis (All chromatograms available in ESI, Figures S35 – 63). N28 was not analyzed due to lack of sample so is marked with -

^[c] Calculated after washing with MeOH to remove DMA.

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204 To gain a better understanding of factors that significantly affected these parameters, as well as any interaction effects, the results were analyzed using DoE software. 205

206 **DoE** analysis

The raw data was analyzed using MODDE®. Data for N28 – N30 were not included as N28 207 208 had a considerable amount of insoluble residue which could not be tested and N29 and N30 were washed with methanol, removing a significant amount of short chain oligomers. This 209 would affect the degree of branching and molecular weight respectively. Data for molecular 210 weight and conversion required a transformation to obtain a normal data distribution; a 211 logarithm (Log(10Log(Y)) and negative logarithm (-10Log(1-Y)) transformation respectively. 212 Selection of significant terms for the coefficients plot was performed using the auto tune 213 214 function.

The summary of fits for each factor tested can be seen in Figure 3.A., where a value of 1 is 215 216 optimal. The model for the branching factor is statistically significant, with a $R^2 > 0.5$, a $Q^2 > 0.5$ 0.1 and a difference of < 0.3 between the two parameters. The conversion and molecular 217 weight factors, however, have a low validity and a large difference between the R² and Q² 218 parameters. R² is a measure of how well the model fits to the original data, while Q² shows 219 how well this model can be used to predict future datasets. Although Q² for molecular weight 220 and conversion is above 0.1, the rather low model validity indicates statistically significant 221 issues for both conversion and molecular weight. Nevertheless, the main objective of the 222 DoE was to maximize the limitation of the polymer branching, and the model for this 223 224 response was both valid and reproducible.



Figure 3.A. Summary of fit for all responses showing the degree of fit (R²), prediction power (Q²)
model validity and reproducibility, and the coefficients plot for **B**. conversion, **C**. molecular weight and
D. branching.

229 From the graph (Figure 3.C.), although there are many factors that look to have an effect on the molecular weight, the p values (Table S4) show that only temperature is significant (at a 230 confidence value of 0.95). Temperature is well known to affect molecular weight in enzymatic 231 232 polycondensations; by increasing reaction temperature higher molecular weights are achieved. This has been observed for many different polymer systems, [53,54] although 233 234 temperature is limited by the thermostability of the enzyme. Interaction effects for temperature are also significant; namely Enz*Temp and Temp*Time. Mahapatro et al. also 235 observed some interaction between reaction time and temperature, as no significant effect of 236 temperature on M_n was observed up to 24h, but at 48h there was a significant difference in 237 the Mn between 65 and 90 °C.^[53] This should be further investigated, as GPC analysis of N29 238 and N30, which were run for 96h, gave much higher molecular weights. 239

240 At a 0.95 confidence, temperature and reaction time are significant factors for the conversion, which agrees with established literature. The monomer ratio is significant only 241 242 when it interacts with either temperature or time (Ratio*Temp and Ratio*Time). Conversion did not differ greatly between conditions, ranging from >68% up to 99%, with the lowest 243 conversion obtained at 40 °C run for 6h. Enzyme wt% seems to have little effect on the 244 conversion, indeed, with all other conditions kept the same, an increase from 1 wt% to 25 245 wt% of enzyme actually resulted in a decrease in conversion (from 93% to 90%) seen in 246 247 Table S1.



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Figure 4. 4D contour response plot showing the effect of monomer ratio, enzyme wt%, temperature and reaction time on the degree of polymer branching.

251 Ratio, temperature, reaction time and Time*Time are all significant factors (p = 0.95) that determine the degree of branching (Figure 3.D.). The predictive contour plot in Figure 4. 252 gives an overview of this, and it can clearly be seen that a higher temperature, lower ratio 253 and a longer reaction time are associated with a higher degree of branching, while enzyme 254 wt% is much less significant. Increasing amounts of Triol-citro, increasing temperature and 255 256 increasing reaction time all result in more branched structures. The most branched structure obtained (N15) had a 16% degree of branching, was run at the highest temperature (90 °C), 257 had the longest reaction time (24h) and had a 1:1 ratio of triol to diester. 258

The fact that enzyme wt% is an insignificant parameter in predicting degree of branching 259 supports the hypothesis that branching is mainly due to acyl migration rather than enzymatic 260 coupling on the secondary hydroxy group. This phenomenon is well studied in 261 carbohydrates, and is an intramolecular reaction that allows the exchange of ester groups 262 among adjacent hydroxy groups,^[55] independent of the enzyme.^[56] This reaction has been 263 shown to be affected by temperature, reaction time, reaction solvent, material used to 264 immobilize the enzyme, pH and acyl species.^[57] Temperature is particularly significant, both 265 in literature and in this work, which is to be expected as this reaction is a thermodynamic 266 process.^[58] 267

Perin et al., determined that in the enzymatic synthesis of esterification of poly(glycerol 268 sebacate), the esterification of primary hydroxy groups was mainly due to acyl migration. 269 When compared to our work, they observed much higher degrees of branching, which the 270 authors hypothesized could be due to the use of solvent or carboxylic acids as reagents.^[29] 271 Solvent affects acyl migration in different ways. Polarity has been shown to have an effect, 272 with low polarity solvents dispersing the charge of the transition state, lowering its energy 273 state and decreasing the rate of the migration reaction.^[57] However when compared to 274 solventless systems, Wang et al. found that all solvents tested had an inhibitory effect on 275 acyl migration (albeit to different degrees)^[59] The use of solvent may also have a beneficial 276 277 effect on molecular weight, as solventless polycondensations often lead to a highly viscous reaction matrix. This was most obvious in reactions N28 - N30, where polymers were 278 viscous and gel-like, and had visibly hindered stirring. Use of diesters as an alternative to 279 dicarboxylic acids would also reduce the rate of acyl migration, due to the different leaving 280 groups. Although both water and methanol have been found to increase the rate of acyl 281 migration,^[57,60] methanol is easier to remove, and is done so via reduced pressure in this 282 283 work.

Aside from this, there are quite a number of factors affecting acyl migration that were not 284 285 investigated here. The CaLB used was immobilized on an acrylic resin, which is a polar carrier, and more polar materials have been known to increase acyl migration.[57] 286 287 Furthermore, the structure of Triol-citro may also have an influence. Unlike glycerol or other 288 sugars such as sorbitol, only two hydroxy groups are adjacent in this monomer; instead of an α - β -y there is an α - β - ϵ triol structure. A previous work attempted to induce acyl migration 289 over the same number of C atoms in mannopyranoside structures, but were unsuccessful, 290 suggesting that two hydroxy groups must be spatially close for this reaction to occur.^[61] 291

292 **Conclusions**

Overall the desired outcome must be carefully considered. For example, reaction N15 gave the highest molecular weight, but also resulted in the most branched structure, at 16%. It is worth performing several more experiments to obtain a valid model for both molecular weight and conversion. In this way, all three responses can be predicted, and parameters can be carefully selected based on the desired properties of the final polymer. With respect to obtaining a high molecular weight, the design space with regard to reaction time should be extended to 96 hours, and the use of solvent should be considered.

The utility of CaLB in selectively polymerizing bulky triol monomers has been clearly shown. 300 301 This work investigated the use of MODDE to determine significant factors in the branching 302 behavior of poly(Triol-citro adipate), and determined that monomer ratio, temperature and 303 reaction time are all significant factors. It was clearly demonstrated that branching was minimal, with a maximum of 16% observed, and this was attributed to acyl migration rather 304 305 than enzyme catalyzed transesterification. Extending the reaction time to 96h resulted in a polymer with a number average molecular weight of up to 13 kDa and only 13% of 306 307 branching. CaLB has been shown to be a suitable biocatalytic alternative to traditional metal catalysts in this reaction. Polyesters produced using this method can be further 308 309 functionalized through reactions with pendant hydroxy groups for more specialized purposes.

310 **Experimental Section**

311 Materials

Tetrahydrofuran (inhibitor-free, for HPLC, \geq 99.9%), Methanol (for HPLC, gradient grade, \geq 99.9%), Chloroform (HiPerSolv CHROMANORM), and *Candida antarctica* lipase B immobilized on acrylic resin (CaLB, code: L4777, \geq 5,000 U/g) were purchased from Sigma Aldrich. Dimethyl adipate (99%) was purchased from Alfa Aesar. All deuterated solvents were purchased from Eurisotop.

317 Trio-citro was synthesised according to a previously published procedure.^[13]

318 **Design of Experiment (DoE)**

The software MODDE® Pro 13 (Sartorius) was used to optimize polycondensation conditions. Reactant ratio, enzyme wt%, temperature and reaction time were selected as quantitative factors. Conversion, number average molecular weight (M_n) and degree of branching were the responses measured, with target for conversion set to 99%, M_n target set high and degree of branching set at 1%. A central composite orthogonal (CCO) design (star distance: 1.54671) was chosen with a quadratic model, to generate a set of experiments for optimisation.

326 Enzymatic Polycondensations

Triol-citro (233.2 mg, 8.5 x 10^{-4} mol) and the appropriate amount of dimethyl adipate (6.4 x 10^{-4} mol – 1.9 x 10^{-3} mol) were added to a 25-mL round bottom flask with CaLB (1 – 25% by weight of monomers). The flask was heated (25 – 100 °C) and stirred at 400 rpm for 6 hours, at which point the system was placed under vacuum (20 mbar) in order to remove the

methanol by-product. Experiments were run from 1 - 30 hrs, and experiments that were run for ≤ 6 hrs were not placed under vacuum.

Upon completion of the reaction, the polymers were washed with approx. 2 mL THF and filtered through cotton packed into a glass Pasteur pipette to remove the insoluble enzyme pellet. The flask was washed with 1 mL THF a further two times, and all fractions were filtered and collected. THF was removed via a rotary evaporator, and the resulting polymer was characterized by NMR and GPC analysis.

If a wash step was performed, the polymer was transferred into a 1.5 mL or 5 mL Eppendorf, precipitated through addition of ice-cold methanol (amount as specified), and centrifuged (10 min at 4 °C, 3700 rpm for 5 mL Eppendorfs and 12700 rpm for 1.5 mL Eppendorfs). The supernatant was removed and the washing step repeated as specified. The resulting polymer was characterized by NMR analysis.

343 Characterization

344 Gel permeation Chromatography (GPC)

Polymers were dissolved in CHCl₃ or THF to a concentration between 2.0 and 2.5 mg/mL and filtered through cotton. The analysis was performed at 30 °C on an Agilent Technologies HPLC System (Agilent Technologies 1260 Infinity) and an Agilent Technologies G1362A refractive index detector was employed for detection. Linear polystyrene calibration standards (250 – 70,000 Da) purchased from Sigma-Aldrich were used to calculate the molecular weights of the polymers.

For those polymers dissolved in $CHCl_3$, a 17,369 6.0 mm ID × 40 mm LHHR-H, 5 µm Guard column and a 18,055 7.8 mm ID × 300 mm L GMHHR-N, 5 µm TSK gel liquid chromatography column (Tosoh Bioscience, Tessenderlo, Belgium) was used, with $CHCl_3$ as an eluent (at a flow rate of 1 mL min⁻¹ for 20 min). For those polymers dissolved in THF, a guard column and chromatography column of the same specification was used with THF as the eluent (at a flow rate of 1 mL min⁻¹ for 20 min).

357 Nuclear Magnetic Resonance (NMR)

¹H-NMR, ¹³Č-NMR, COSY, HMBC and HSQC spectroscopy was performed using a JEOL ECZ400R/S3 at a frequency of 400 MHz, with some ¹H-NMR performed on a Bruker Avance II 400 (resonance frequencies 400.13 MHz for ¹H) equipped with a 5 mm N₂-cooled cryo probe (Prodigy) with z–gradients at room temperature with standard Bruker pulse programs. Samples were dissolved in either CDCl₃ or DMSO- d_6 as a solvent.

363 Acknowledgements

Dr Louis M. M. Mouterde (URD ABI) is acknowledged for his contribution to the original synthesis of Triol-citro (reference [13]). This work is supported by European Union's Horizon 2020 research and innovation programme under grant agreement No 953073, project UPLIFT (sUstainable PLastIcs for the Food and drink packaging indusTry). S.F. and F.A. thank Grand Reims, Département de la Marne and Grand Est Region for their financial support.

370 Data availability statement

All data not present in the main text or the ESI are available from the authors upon request.

372 Conflicts of Interest

- 373 The authors declare no conflicts of interest.
- **Keywords:** Bio-based, Enzyme catalysis, Green Chemistry, NMR spectroscopy, Polyesters
- 375 **References**

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- 476 Polyesters were enzymatically synthesised from a bio-based triol, and a Design of
- 477 Experiment approach was used to investigate factors that affect the branching during

478 synthesis of this polymer.