1 Synthesis and unambiguous NMR characterization of linear and

2 branched *N*-alkyl chitosan derivatives.

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8 Abstract

9 Chitosan, sourced from abundant chitin-rich waste streams, emerges as a promising candidate in the 10 realm of future functional materials and chemicals. While showing numerous advantageous properties, 11 chitosan sometimes falls short of competing with today's non-renewable alternatives. Chemical 12 derivatization, particularly through N-alkylation, proves promising in enhancing hydrophobic 13 functionalities. This study synthesizes fifteen chitosan derivatives (degree of substitution = 2-10 %) 14 using an improved reductive amination method. Next, selective depolymerization through acid 15 hydrolysis reduced the chain rigidity imposed by the polymer backbone. This facilitated unambiguous 16 structural characterization of the synthesized compounds using a combination of common NMR 17 techniques. Two potential side reactions are identified for the first time, emphasizing the need for 18 detailed structural information to unlock the true potential of these derivatives in future applications.

19 Hypothesis

- 20 The increase in chain mobility induced by the selective depolymerization of aliphatic *N*-alkyl chitosan
- 21 derivatives allows for an unambiguous NMR characterization.

22 1. Introduction

- 23 Chitin is a class of widely available, linear, non-toxic and biodegradable polysaccharides. Chitin is 24 naturally present in the exoskeleton of crustaceans, the fungal cell wall and the structural components 25 of insects, making it the second most abundant natural biopolymer. Chitin can easily be extracted from 26 secondary leftover waste streams, due to its wide availability, making it ever more attractive (Triunfo 27 et al., 2022). Chitin mainly consists of $\beta(1\rightarrow 4)$ linked N-acetyl-D-glucosamine units, with a few 28 percentages of $\beta(1\rightarrow 4)$ linked D-glucosamine building blocks. Chitin is generally transformed into 29 chitosan through deacetylation to improve its versatility and aqueous solubility. Chitosan is also non-30 toxic, biodegradable and biocompatible, while free amino groups afford natural antimicrobial and 31 antifungal activities. However, this activity is limited in water at physiological pH as the available amino 32 groups need to be sufficiently protonated for chitosan to dissolve ($pK_a \approx 6-8$) (Anthonsen & Smidsrød, 33 1995).
- The selective addition of aliphatic side chains onto the amino groups of chitosan produces an interesting class of chitosan derivatives called *N*-alkyl chitosans, named after their *N*-alkyl amine counterparts. These amphiphilic compounds, which are mostly obtained through reductive amination, have been applied in numerous fields as viscosity modifiers (Desbrières et al., 1996), hemostatic agents (Dowling et al., 2011), wood additives(Mati-Baouche et al., 2019), emulsifiers (Kalliola et al., 2018), edible food coatings (de Britto & de Assis, 2007), drug delivery agents (Benediktsdóttir et al., 2014) and recently even as algal flocculants (Demir-Yilmaz et al., 2023).
- Another interesting application that makes use of their amphiphilic nature is as antimicrobial agents.
 As chitosan mostly interacts with the microbial outer surface, it is hypothesized that the addition of
- 43 hydrophobic aliphatic side chains might aid in these types of interactions as well as broaden the

44 applicable pH range (Kong et al., 2010). However, most structure-activity studies focus on excessively 45 alkylated derivatives or present scattered single data points that are difficult to compare with one 46 another (Paula et al., 2020) (Ma et al., 2008). For these excessively alkylated derivatives, quaternization 47 is required to enable aqueous bio-testing, making it hard to decouple the effects of both structural 48 modifications afterward (Sahariah et al., 2015)(Kim et al., 1997)(Jia et al., 2001). Hence, structure-49 activity studies of the biological effects on microorganisms of different types of N-alkyl modifications 50 without subsequent quaternization, are scarce. This makes the exact effect on the antimicrobial 51 activity of adding hydrophobic chains rather elusive (Sahariah & Másson, 2017).

52 In the current study, the almost 30-year-old reductive amination protocol was revisited and given a 53 modern update towards a more scalable and less environmentally taxing process. The subsequent 54 protocol was applied to synthesize a library of fifteen different linear and branched N-alkyl chitosans, 55 of which several new derivatives, with an envisioned degree of substitution (DS) of 10 %. This low 56 substitution ensured a similar solubility profile as native chitosan without the need for subsequent 57 quaternization. Subsequently, these derivatives were selectively depolymerized to enable an 58 unambiguous elucidation of the structure of thirteen compounds and indications for the other two. 59 Moreover, this further led to the identification of two side reactions for the first time. These include 60 the N-isopropylation of the chitosan backbone during product isolation, as well as the acid-catalyzed 61 hydration of unsaturated side chains on the chitosan backbone. Guidelines for the NMR 62 characterization of this class of compounds are presented, aiming to serve as a welcome reference for 63 future studies, enabling an unambiguous correlation between structure and activity in the future.

64 2. Results and discussion

65 2.1. *N*-alkyl chitosans and their NMR characterization: state of the art

The degree of substitution (DS) is one of the most important structural parameters for chemically 66 67 modified chitosan. Most of the time, the DS is derived from relative ¹H NMR integration, which makes 68 it heavily reliant on the correct peak assignment thereof. However, a fact that is quite often overlooked 69 within the literature regarding N-alkyl chitosans is the employed conditions during NMR analysis, 70 especially ¹H NMR analysis. To dissolve chitosan or its hydrophobic N-alkylated counterparts in 71 (deuterated) water, acidic conditions are required. Subsequently, most of the amino groups on the 72 chitosan chain will be protonated during aqueous NMR analysis. However, this significantly influences 73 the chemical ¹H NMR shift of the neighboring 1' methylene/methine protons next to these protonated 74 amino groups. This phenomenon is well established for structurally similar N-alkyl amines, as downfield 75 shifts of about 0.5-1 ppm can be observed in ¹H NMR due to the decreased shielding of the α 76 methylene/methine proton(s) next to the protonated positively charged ammonium ion relative to 77 those of the uncharged amine (Ma & Warnhoff, 1965). This effect is relatively less pronounced in ¹³C 78 NMR (Sarneski et al., 1975). This has led to seemingly two different types of ¹H NMR peak assignments 79 for these N-alkyl chitosans (Figure 1). On one hand, all of the aliphatic proton signals are assigned to 80 the different signals observed within the 0-2 ppm region, including the 1' protons next to the 81 protonated amino groups (Bobu et al., 2011)(Rabea et al., 2005)(Sajomsang et al., 2009)(Venkataraman 82 et al., 2013). On the other hand, the 1' protons are assigned about 1-2 ppm downfield, overlapping 83 with the two distinct H₂ signals of the derivative (Desbrières et al., 1996)(Kurita & Isogai, 2010)(Sashiwa & Shigemasa, 1999). Following the reasoning above, we hypothesize the latter to be correct. 84 85 Furthermore, looking at the example depicted in Figure 1, simple integration of the signals between 0-86 2 ppm in ¹H NMR gives an integral ratio of about 3:2:1.6, which is very close to the theoretical value of 87 3:2:2 if the 1' protons are assigned downfield. A more in-depth discussion regarding this matter will 88 follow later (see Section 2.3.).



Figure 1. ¹H NMR (400 MHz) spectrum of N-butyl chitosan ($DS_{ch} = 10$ %, based upon ¹H NMR integration, see Section S4.) at 20 mg/mL in 1 vol% d-TFA in D₂O, together with the two most popular structural assignments present in literature, indicated in green and blue. Acetone as a minor impurity is indicated with an *.

89 2.2. Synthesis of *N*-alkyl amines via reductive amination

90 Within this study, we aim to synthesize fifteen different hydrophobic N-alkyl chitosans with an envisioned DS of 10 %. All of these derivatives need to be soluble in aqueous media to facilitate NMR 91 92 analysis in acidic D_2O , as well as aqueous biotesting in the future. This DS was chosen because, early 93 on, we observed, as did several other groups (Lepeltier et al., 2015)(Desbrières et al., 1996)(Sajomsang 94 et al., 2008) (Demir-Yilmaz et al., 2023), that derivatives with too high of a DS become insoluble in acidic 95 aqueous media. This phenomenon is especially prevalent when the added chains become more hydrophobic. Moreover, similar DS values are desirable as they will enable the study of the effects of 96 97 the different groups added, establishing preliminary structure-activity relationships (SAR) in the future.

98 The reductive amination developed by Hall and Yalpani (Yalpani & Hall, 1984) and refined by Rinaudo, 99 Desbières and Martinez (Desbrières et al., 1996) is by far the most utilized method to synthesize these 100 hydrophobic *N*-alkyl chitosan derivatives. However, yet effective, this method has two major 101 drawbacks: toxic NaCNBH₃ is utilized as the reducing agent and an almost 50/50 mixture of organic 102 solvent (EtOH, MeOH or DMF) and water is required for the reaction to proceed effectively (Sashiwa & 103 Shigemasa, 1999) (Sajomsang et al., 2008).

104 To try and alleviate some of the environmental burden accompanied by the production of these 105 compounds and to move towards a more scalable process while still obtaining qualitative products in 106 the end, several different reaction conditions and isolation methods were explored (see Sections S1. 107 and S2.). Afterward, the recently proposed (Cok et al., 2020)(Cosenza et al., 2011) α -picoline borane 108 complex (PICB) was selected based on the GSK reagent selection guide for reductive aminations as the 109 preferred reducing agent over the problematic NaCNBH₃ (Adams et al., 2013). Acidic water containing low amounts of cetylpyridinium chloride (CPC) was selected as an alternative solvent to the mixed 110 111 solvent systems, as water is still the preferred solvent of choice in many cases (Prat et al., 2016). Any residual PICB was quenched by acidifying and heating the sample and a pure product was obtained by 112 113 precipitation with an excess of acetone. Furthermore, utilizing this methodology, no significant 114 molecular weight reduction could be observed for the derivatives compared to the chitosan starting 115 material (see Section S3.).

- 116 Next, a relationship between the obtained DS and the added equivalents of aldehyde (see Figure 2)
- 117 was established for these newly applied conditions. As expected, as the DS increased, the aqueous 118 solubility worsened, especially for samples with a DS > 25 %.



Figure 2. The relationship between the observed DS (based upon ¹H NMR integration, see Section S4.) and the added equivalents of aldehyde. Octanal was selected as a representative long, linear and aliphatic aldehyde that is practically insoluble in water. No DS was calculated for the product obtained after the addition of 1 equivalent octanal as the product became insoluble in aqueous media. In every experiment, the same amount of equivalents of PICB were added as the equivalents of aldehyde.

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- 120 Finally, a particularly interesting ¹H-¹H COSY coupling was observed when, by accident, the quenching
- step was omitted. This coupling indicated the presence of *N*-isopropyl groups onto the chitosan chain
- 122 (Kurita & Isogai, 2010), next to the envisioned *N*-alkyl substituent (see Figure 3). This product originates
- 123 from the covalent grafting of acetone onto chitosan, which is used for the precipitation of the chitosan
- derivative (see Section 4.4.). This process occurs via a reductive amination facilitated by leftover PICB.
- 125 Therefore, to avoid unwanted side reactions during product isolation, we would like to stress the
- importance of guenching the residual PICB after the reaction has finished.



Figure 3. ¹H-¹H COSY NMR (400 MHz) spectrum of N-butyl chitosan contaminated with N-isopropyl groups ($DS_{ch, butyl} = 24$ %, $DS_{ch, isopropyl} = 9$ %, based upon ¹H NMR integration, see Section S4.) at 20 mg/mL in 1 vol% d-TFA in D₂O. The ¹H-¹H couplings of the different side chains are indicated in yellow for N-butyl and red for N-isopropyl groups.

128 2.3. Structural elucidation of the obtained derivatives

To enable a full and unambiguous structural characterization, *N*-butyl chitosan (DS = 10 %, based upon ¹H NMR integration, see Section S4.) was synthesized as a model compound. 1 vol% d-TFA in D₂O was selected as the NMR solvent of choice as this generally gave the best resolution for all the obtained 1D and 2D NMR data.

133 2.3.1.Direct one and two-dimensional NMR analysis

134 Unfortunately, the ¹H-¹H COSY spectrum of the *N*-butyl chitosan sample at 20 mg/mL showed no conclusive evidence of the position of the 1' protons in ¹H NMR either. All expected ¹H-¹H couplings 135 136 between neighboring protons on the added backbone were present, except the coupling between the 137 1' protons and their neighboring 2' protons. Most likely, the limited mobility of these 1' protons, as 138 these are the closest to the chitosan backbone, together with the low degree of substitution caused 139 this signal to dissipate in the noise. Hence, an obvious first solution would be to increase the degree of 140 substitution by increasing the equivalents of aldehyde added (see Figure 2), as can be seen in Figure 3. 141 However, for longer N-alkyl chitosan the degree of substitution has an upper solubility limit (vide infra), 142 making this a not viable solution for all fifteen derivatives. Therefore, it was hypothesized that a 143 significant reduction in polymer molecular weight would improve the overall mobility of the entire 144 molecule, including the 1' atoms of the added side chain, relatively increasing the signal strength of all 145 atoms with previously limited mobility compared to the more mobile 3' and 4' outer proton signals.

2.3.2. Molecular weight reduction of N-alkyl chitosan towards N-alkyl COS

Two well-described chemical methods for the production of chitooligosaccharides (COS) were evaluated for the direct and selective cleavage of modified *N*-alkyl chitosans towards *N*-alkyl COS. These methods are oxidative cleavage with NaNO₂ (Allan & Peyron, 1995)(Tømmeraas et al., 2001) and acidic hydrolysis with concentrated HCI (Vårum et al., 2001). To our knowledge, this approach to obtain *N*alkyl COS was not described before, as generally COS are produced first and modified afterward.

152 2.3.2.1. Oxidative NaNO₂-mediated depolymerization

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153 The oxidative HONO cleavage showed very reproducible results for non-modified chitosan, as 154 numerously described before (Chapelle et al., 2019)(Coudurier et al., 2020)(Mo et al., 2020). However, 155 despite these promising early results, an N-butyl chitosan sample yielded a very heterogeneous 156 product, which only partly dissolved after depolymerization. Moreover, several new peaks appeared 157 in the 0-2 ppm region within the ¹H NMR spectrum (see Section S5.), further indicating a complex 158 product mixture. As previously described by Allan and Peyron (Allan & Peyron, 1995), cleavage only 159 occurs at the primary amino groups of chitosan, which are subsequently deaminated, generating a 2,5-160 anhydro-D-mannose unit on one side and cleaving the glycosidic bond in the process (see Figure 4). 161 However, as secondary N-alkyl amino units are present, nitrosamines could be generated, which might 162 be one of the factors contributing to the observed complex behavior. Next to this, if the cleavage 163 selectively occurs only on the primary amino groups of chitosan, degrading them in the process, there 164 will be a relative increase in N-alkyl chains in certain COS fractions, severely limiting their further acidic 165 aqueous solubility. This could further explain the heterogeneous nature of the depolymerization 166 products. No further attempts were made utilizing this method, as the heterogeneous nature of the 167 obtained products, instead of simplifying analysis, further complicated subsequent NMR analysis.



Figure 4: The oxidative NaNO₂-mediated cleavage of chitosan.

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169 2.3.2.2. Acid hydrolysis of chitosan in concentrated DCI/HCI

170 Initially, depolymerization experiments were conducted directly in 2M DCl for 24 hours under reflux171 conditions. This had three major advantages:

- The samples could be analyzed directly afterward by NMR.
- The resonance of the residual solvent peak (HOD) does not interfere with any of the anomeric carbohydrate signals as the high acidity of the sample induced a strong downfield shift (Einbu & Vårum, 2007)(P. Wang et al., 2021).
- At this DCl concentration, the rate of deacetylation is similar to the rate of depolymerization (Einbu & Vårum, 2007). This resulted in fully deacetylated samples, yielding a more uniform copolymer with only two different monomers instead of three, further simplifying subsequent (especially ¹³C) NMR analysis.

However, no pure COS could be obtained this way as indicated by NMR (see Sections S6. and S9.). 180 Several new impurities could be observed in both ¹H NMR and especially in ¹³C NMR when comparing 181 these spectra with existing literature (Domard & Cartier, 1989)(Fukamizo et al., 1991)(Sugiyama et al., 182 183 2001). We hypothesize that most of these impurities are remnants of the remaining chitin-glucan 184 complex within the utilized chitosan of fungal origin (Aspergillus niger biomass). This aqueous insoluble complex is naturally present in fungal biomass and is prone to acidic degradation (Farinha et al., 2015). 185 186 The presence of several different types of glucans or even glucose could explain most of these signals which are situated in the 50-100 ppm region in ¹³C NMR. 187

Additionally, the automatic tuning and matching (ATMA) of our NMR probe was no longer an option as a result of the high ionic strength of the solvent (Einbu & Vårum, 2008), worsening all subsequent NMR analysis and even rendering proper two-dimensional ${}^{1}\text{H}{}^{-13}\text{C}$ HSQC analysis impossible. As these analyses are a vital part of the structural elucidation of our obtained compounds, we decided to use 1 vol% d-TFA in D₂O as our NMR solvent system instead.

193 Selective precipitation of the longer COS fraction was performed by the addition of five volumes of 194 acetone to the hydrolysate (obtained in 2M HCl after 24 hours under reflux conditions), to further 195 simplify subsequent NMR analysis. A second fraction precipitated upon the addition of excess acetone. Further ¹H NMR and especially ¹³C NMR analysis confirmed the purity of the obtained longer COS 196 197 fraction without any of the previously observed impurities (see Sections S7. and S10.). For our second 198 isolated fraction, a mixture containing a large amount of short water soluble $\beta(1,3)$ -glucans is suggested 199 based upon the comparison of our obtained ¹H NMR data with the ¹H NMR data of the isolated fraction 200 obtained in the work of Hanashima et al. (see Section S8.) (Hanashima et al., 2014).

201 2.3.3.Structural NMR analysis of N-alkyl COS

Based on the considerations above, the following workflow was used to obtain fifteen *N*-alkyl COS, which were subsequently subjected to one-dimensional ¹H, ¹³C and DEPT-135 NMR and twodimensional ¹H-¹H COSY and ¹H-¹³C HSQC NMR analysis.

- Synthesis of the aliphatic *N*-alkyl chitosan via reductive amination.
- *N*-isopropylation check via ¹H-¹H COSY NMR analysis.
- Depolymerization of the aliphatic *N*-alkyl chitosan towards its respective *N*-alkyl COS.
- Precipitation of the longer COS.HCl fraction with five reaction volumes of acetone.
- NMR analysis of the dried precipitate.
- 210 Due to the short chain length of the obtained COS compared to the original chitosan, the internal units
- could be differentiated from their corresponding reducing (α and β) and non-reducing ends (see Figure 5).



Figure 5. Schematic overview of the produced N-alkyl COS. The introduced color scheme will be adapted in all subsequent NMR analyses.

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2.3.3.1. The example of *N*-2-ethylbutyl COS

N-2-ethylbutyl chitosan (DS = 10 %, based upon ¹H NMR integration, see Section S11.10.) was selected
 as a representative example. All NMR analyses of the remaining fourteen derivatives are supplied in
 the supplementary information (see Section S11.). Similar patterns, as described below, were observed
 for all samples, except for unsaturated *N*-alkyl chitosan derivatives (*vide infra*). Additionally, tables
 containing all the observed chemical shifts (¹H and ¹³C NMR) are supplied within the supplementary
 information (see Section S13.).

First, all ¹³C signals of non-modified units were assigned based on previously published literature 221 222 (Domard & Cartier, 1989)(Fukamizo et al., 1991). Next, all new signals of modified units were assigned 223 based on their hypothesized structure (see Figure 6). Additionally, DEPT135 (Distortionless 224 Enhancement by Polarization Transfer) was utilized to further verify the nature (CH/CH₃ or CH₂) of all 225 newly assigned ¹³C signals (see Section S11.10.). For the aliphatic 2-ethylbutyl side chain, the obtained 226 ¹³C shifts of 49.3, 37.6, 22.7/22.4 and 9.8/9.3 ppm are in good agreement with previously reported 227 data (de Britto & de Assis, 2007)(Sashiwa & Shigemasa, 1999), as well as reported values for protonated 228 aqueous alkylamines (Sarneski et al., 1975). However, the chemical shifts of the carbon atoms on the 229 1-6 positions of the modified internal COS units, rather than the chemical shifts of the 1'-4' positions 230 of the attached alkyl chain, are much harder to assign, and we did not manage to find any direct data 231 to compare our results with. Yet, despite being overlooked quite often, in theory, the resonance signals 232 of neighboring carbon atoms are expected to shift if the amino group on the 2 position is modified 233 because of the covalent attachment of the added side chain. A chemical shift of 96.1, 68.7, 61.6 and 234 60.7 ppm was observed for the carbon atoms at the 1, 3, 2 and 6 positions of the modified internal COS 235 units. When comparing these results with our previously obtained data for lowly modified (DS = 20%) N-sulfopropyl chitosan (Van Poucke et al., 2023), very similar values of 96.5 and 61.5 ppm were
 obtained for the carbons situated on the 1 and 2 positions, respectively, of the modified chitosan units.



Figure 6. ¹³C NMR (100.6 MHz) spectrum of N-2-ethylbutyl COS ($DS_{cos} = 9 \%$, based upon ¹H NMR integration, see Section S11.10.) at 20 mg/mL in 1 vol% d-TFA in D_2O . The color scheme introduced in Figure 5 was adapted for all non-modified units. Signals of modified units are indicated in yellow. Acetone as a minor impurity is indicated with an *.

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239 Based upon the ¹³C chemical shifts, proton signals in ¹H NMR were assigned via their respective ¹H-¹³C 240 HSQC (Heteronuclear Single Quantum Coherence) NMR correlation (see Figure 7). Next, the ¹H NMR 241 chemical shifts of the non-modified internal and external units were cross-checked with existing 242 literature (Sugiyama et al., 2001). For the added aliphatic side chain, the obtained chemical shifts of 243 3.14, 1.67, 1.39 and 0.85 ppm are in good agreement with previously reported data for aliphatic N-244 alkyl chitosan derivatives (Desbrières et al., 1996) as well as protonated aqueous alkylamines (Ma & 245 Warnhoff, 1965). Notably, the data unambiguously shows the exact ¹H NMR chemical shift of 3.14 ppm 246 for the 1' protons, which undergo a significant downfield shift due to the protonation of the secondary 247 amino group. Furthermore, a similar argument can be made for ¹H NMR as for ¹³C NMR for the 1-6 248 positions of the modified internal COS units. The proton on the 1_{α} anomeric position of the modified external COS units exhibits a chemical shift of 5.54 ppm, while the protons on the 1, 3, and 2 positions 249 250 of the modified internal COS units show chemical shifts of 5.02, 4.05, and 3.24 ppm, respectively. 251 Additionally, the ¹H NMR chemical shift of the proton on the 2 position of the modified internal COS 252 unit closely aligns with our previous work (Van Poucke et al., 2023) and with findings from Kurita and 253 Isogai (Kurita & Isogai, 2010).



Figure 7. ¹H-¹³C HSQC NMR (400 MHz) spectrum of N-2-ethylbutyl COS (DS_{cos} = 9 %, based upon ¹H NMR integration, see Section S11.10.) at 20 mg/mL in 1 vol% d-TFA in D₂O. The color scheme introduced in Figure 5 was adapted for all nonmodified units. Signals of modified units are indicated in yellow. CH₂ signals are indicated in green and CH/CH₃ signals in blue. Acetone as a minor impurity is indicated with an *.

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255 Next, these ¹H NMR assignments were verified via ¹H-¹H COSY (COrrelation SpectroscopY). The most 256 important ¹H-¹H couplings are assigned in Figure 8. This leads to the following conclusions:

- 257 The presence of new 1-2 and 2-3 couplings of the protons on the internal and external reducing units, indicates that a certain percentage of the amino groups of COS are indeed covalently 258 modified. 259
- The proton coupling along the added side chain (1' to 4') indicates the covalent attachment of 260 • an N-2-ethylbutyl group.



Figure 8. ¹H-¹H COSY NMR (400 MHz) spectrum of N-2-ethylbutyl COS (DS_{cos} = 9 %, based upon ¹H NMR integration, see Section S11.10.) at 20 mg/mL in 1 vol% d-TFA in D_2O . The color scheme introduced in Figure 5 was adapted for all nonmodified units. Signals of modified units are indicated in yellow.

263 Unfortunately, no direct coupling between the 1' position of the added side chain and the 2 position 264 on the chitosan backbone could be observed via additional ${}^{1}\text{H}{}^{-13}\text{C}$ HMBC (Heteronuclear Multiple Bond 265 Correlation) analysis (see Section S12.). Nevertheless, based on the arguments above, the direct 266 covalent attachment of the *N*-2-ethylbutyl group is evident.

267 Finally, the carbons and protons at positions 3' and 4' show an interesting behavior. When solely looking 268 at the structure of the attached N-2-ethylbutyl group while neglecting the chiral COS backbone, one 269 could assume that the carbons and protons on either side of the chain are chemically equivalent. 270 However, this seems not to be the case as evidenced by the two signals at 9.3 and 9.8 ppm for the 271 carbons at the 4' position and 22.4 and 22.7 ppm for the carbons at the 3' position in Figure 6. 272 Additionally, the signal in Figure 7 at 0.85 ppm for the protons at the 4' position shows a more complex 273 multiplicity instead of the expected triplet. The observed differences are most likely due to several 274 different processes, some of which are temperature dependent as well as the inherent diastereotopic 275 relationship of the substituents due to the presence of the chiral COS backbone. To further study this 276 behavior, the signal at 0.85 ppm in ¹H NMR at 298 K was compared with the corresponding signal at 277 308, 328 and 343 K. In Figure 9, a temperature-dependent behavior can be observed, in which signal 278 broadening is reduced as the temperature increases. This is most likely due to the reduced influences 279 of several temperature-limited processes. An example might be the limited mobility of the chiral COS 280 backbone, which could impose steric constraints on the added side chain, which is reduced at higher 281 temperatures. However, in theory, at any given temperature, some small difference should remain, as 282 observed for several chiral alkyl amines in the past, due to the previously mentioned diastereotopic

relationship (Ando et al., 1979).



Figure 9. Zoom of the 0.8-0.9 ppm region of the ¹H NMR (400 MHz) spectrum of N-2-ethylbutyl COS (DS_{cos} = 9 %, based upon ¹H NMR integration, see Section S11.10.) at 20 mg/mL in 1 vol% d-TFA in D₂O at 298, 308, 328 and 343 K.

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2.3.3.2. The particular case of unsaturated *N*-alkyl chitosan derivatives

Figure 10 depicts the ${}^{1}H{}^{-13}C$ HSQC NMR spectrum obtained after acid-mediated depolymerization of *N*pent-4-enyl chitosan (DS_{ch} = 7 %, based upon ${}^{1}H$ NMR integration, see Section S11.7.). Within the spectrum (based on their expected chemical shifts and the HSQC phasing) there is a terminal methyl group at the 5' position present instead of the expected terminal methylene group, indicating a modification on the added side chain during the depolymerization reaction. Based upon the data presented in Figure 10, we hypothesized that the alkene moiety was hydrated under the thermal aqueous acidic conditions. This was further confirmed by additional ${}^{1}H$, ${}^{13}C$, ${}^{1}H{}^{-1}H$ COSY, ${}^{1}H{}^{-13}C$ HMBC and DEPT135 NMR analyses, both before and after depolymerization, confirming the structure of the original *N*-pent-4-enyl chitosan and the subsequently formed *N*-4-hydroxypentyl COS (see Sections

295 S11.7. and S11.8.).



Figure 10. ${}^{1}H{}^{-13}C$ HSQC NMR (400 MHz) spectrum of N-4-hydroxypentyl COS (DS_{COS} = 4 %, based upon ${}^{1}H$ NMR integration, see Section S11.8.) at 20 mg/mL in 1 vol% d-TFA in D₂O. Signals of modified units are indicated in yellow. CH₂ signals are depicted in green and CH/CH₃ signals in blue.

296 The addition of the hydroxyl group at the 4' position because of the higher stability of the intermediate 297 secondary carbocation is a textbook example of Markovnikov's rule. Logically, substituted alkenes 298 should react faster because of the formed intermediate tertiary carbocation. Hence, the reductive 299 amination of chitosan with natural monoterpenoid aldehydes containing prenyl groups should yield 300 substrates that are easily hydrated. To this extent, two natural aldehydes, citral and citronellal, were 301 reductively aminated to chitosan. However, the exact structure of the obtained compounds after 302 depolymerization could not be determined via NMR analysis because neither of the added sidechains 303 gave sufficiently strong signals during analysis. Only the expected covalent addition could be confirmed 304 (see Sections S11.15. and S11.16.). Yet, solely based on ¹H NMR analysis of the chitosan-citronellal 305 adduct (see Figure 11), the following structures are proposed. Before depolymerization, only a fraction 306 of the prenyl groups of the added side chain remains intact while the rest is hydrated. This is evidenced 307 by the presence of four different methyl groups while the original aldehyde has three. To explain this 308 odd observation, we propose the methyl assignment, based on their expected chemical shifts, depicted 309 in Figure 11. The two methyl groups on the original prenyl group are observed at the most downfield 310 positions, specifically at 1.67 and 1.60 ppm, while the remaining methyl group is situated at the most 311 shielded position, resonating at 0.89 ppm. The methyl groups on the hydrated prenyl group overlap 312 and resonate at an intermediate value of 1.17 ppm. Additionally, there is a small signal of the methyne proton of the prenyl group present at 5.18 ppm. Unsurprisingly, after depolymerization, the obtained 313 product is fully saturated (see Figure 11 and Section S11.15.). Based upon the disappearance of the 314 315 two methyne protons at 5.27 and 5.16 ppm after the depolymerization of the chitosan-citral adduct 316 (See Figure 11), a similar behavior is expected. We hypothesize that the initial hydration took place 317 during the quenching of residual PICB (see Section 4.4.) as the mixture was acidified and heated. A similar selective acid-catalyzed hydration was observed by Ishino and Kumanotani for several 318 citronellal-amine adducts towards hydroxycitronellal (Ishino & Kumanotani, 1974). The observations 319 320 above could also be of particular interest when essential oils are utilized as green aldehyde feedstocks 321 as in the work of Paula et al. (Paula et al., 2020), as FTIR analysis, qualitative Baeyer-tests and ¹H NMR analysis indicated the presence of unsaturation in the obtained chitosan derivatives. However, during
 their ¹H-¹³C HSQC NMR analysis not even a trace of unsaturation was observed. The fact that their ¹H ¹³C HSQC NMR analysis was performed at 70 °C under acidic conditions, in contrast with the shorter ¹H
 NMR analysis which was performed at room temperature, might be a possible explanation for this
 rather strange observation, as most prenyl groups could have been hydrated during the HSQC analysis.



Figure 11. The proposed chemical structures for the chitosan-citronellal adduct (a) and the chitosan-citral adduct (b) before depolymerization are depicted. ¹H NMR (400 MHz) spectrum of the chitosan-citronellal adduct (c) ($DS_{ch} = 5$ % and $DS_{COS} = 3$ %, based upon ¹H NMR integration, see Section S11.15.) and chitosan-citral adduct (d) at 20 mg/mL in 1 vol% d-TFA in D_2O , before (blue) and after (red) depolymerization.

327 2.4. Overview of the synthesized N-alkyl chitosans and their COS derivatives.

328 Table 1. An overview of the synthesized N-alkyl chitosans, their COS derivatives, and their calculated DS values based upon ¹H

329 NMR integration. The displayed standard deviations are based on the interpeak ¹H NMR variance. The samples of which the

added side chain structurally changed during depolymerization are indicated with an * (see section 2.3.3.2.). N.D. = not-

determined; DS_{ch} = degree of substitution of the chitosan sample; DS_{cos} = degree of substitution of the COS sample.

	R =	DS_{ch}	DScos		R =	DS_{ch}	DScos		R =	DS_{ch}	DS cos
		(%)	(%)			(%)	(%)			(%)	(%)
А	H ₂			F	H ₂			К		8	6
	,_N*	2	2		, N ⁺	5	5		\mathbb{N}^{+}	±	±
	I									0.13	0.10
В	H ₂	9	9	G	H ₂	7	4*	L	H ₂	6	5
	N ⁺	±	±		N ⁺	±	±		,·N ⁺	±	±
		1.38	0.59			1.20	0.09			0.48	0.24
С	H ₂		10	н	H ₂	9	7	М	H ₂	8	7
		11	±			±	±		· N ⁺	±	±
			0.04			1.33	0.44			0.60	0.12
D	H ₂	8	8	1		10	9	Ν	H ₂		
	, N⁺	±	±		N ⁺	±	±			5	3*
		1.13	0.47		/ V V	0.86	0.14				
Е	H ₂	11	11	J	H ₂	5	4	0	H ₂		
	, N ⁺	±	±		,·N ⁺	±	±		, N ⁺	N.D.	N.D.*
		0.95	0.55			0.94	0.14				



332

An overview of all the synthesized and characterized derivatives is given in Table 1, together with their DS calculated via ¹H NMR (see Section S11.). Very similar values are obtained before and after 335 depolymerization, making the presence of insoluble NMR invisible fractions before depolymerization 336 highly unlikely at these low degrees of substitution. This is in contrast with the NMR invisible fractions 337 observed by Novoa-Carballal et al. for the chitosan-PEG copolymer (Novoa-Carballal et al., 2013). 338 Moreover, a slight decrease in DS can be observed for certain samples after depolymerization. This 339 might be attributed to the increased organic solubility of highly modified COS fractions, which no longer 340 precipitate upon acetone addition (see section 2.3.3.2). As expected, due to the better-defined 341 baseline/peaks, as well as the increased mobility of the less mobile atoms, the interpeak variability in 342 DS decreased after depolymerization. A nice example of this phenomenon can be found in the 343 supplementary information (e.g. see Section S11.2.). Finally, for nine of the fifteen chitosan derivatives

a DS value within the range of 7-11 %, close to the envisioned 10 %, was obtained.

345 3. Conclusion

346 In this study, fifteen N-alkyl chitosan derivatives with varying linear, branched or unsaturated side 347 chains, incorporating between three and ten carbon atoms, were successfully synthesized. A similar DS 348 of approximately 10 % for nine compounds, determined via ¹H NMR, opens avenues for structure-349 activity studies on the effects of the different added alkyl chains. Next, these fifteen derivatives were 350 subjected to selective acid-mediated depolymerization and subsequent fractionation by precipitation, obtaining high-purity N-alkyl COS hydrochloride salts. While small amounts of these modified COS were 351 352 used for NMR structural elucidation, other unexplored properties, like foam stabilization or 353 antimicrobial activity, will be investigated in the future. The reduced chain rigidity in these 354 depolymerized compounds facilitated the unambiguous structural determination of thirteen 355 compounds while indicating the structure of the remaining two, through a combination of various NMR 356 techniques. NMR spectroscopy is a crucial tool for the analysis of modified polysaccharides, yet correct 357 structural characterization of the obtained derivatives remains a big challenge. Notably, relying solely 358 on ¹H NMR analysis, particularly on high molecular weight samples, may lead to inconclusive results, 359 as demonstrated by the discrepancies in common assignments for this well-known class of chitosan 360 derivatives found in the literature. Yet, these detailed structural characteristics are essential to realize 361 the full potential of these biobased derivatives in the transition toward a climate-neutral future 362 (Gericke et al., 2024).

363 4. Experimental

364 4.1. Materials

All chemicals and reagents were commercially available and obtained in analytical purity or higher from 365 Tokyo Chemical Industry Co., Sigma-Aldrich and Carl Roth. Low molecular weight chitosan (KitoGreen®) 366 367 (Mw = see Section S3., DA = 0.08, origin = fungal, 5.27 mmol NH₂/gram, insoluble fraction: 12-15 m% 368 chitin-glucan complex) was supplied by KitoZyme (Herstal, Belgium) and utilized without further 369 purification. The DA was determined based on ¹H NMR analyses (Lavertu et al., 2003). The amount of 370 reactive NH₂ units in a gram of material was determined via an average of three potentiometric 371 titrations (H. Wang et al., 2011). For a certain amount of equivalents, the obtained 5.27 mmol 372 NH₂/gram was utilized to calculate the amount of reagent that should be added. All solvents were utilized as received without further purification. ¹¹B NMR was utilized to check for the presence of any 373 374 leftover boron salts in the products obtained via reductive amination.

375 4.2. NMR analysis

1D and 2D NMR analyses were recorded at room temperature, unless stated otherwise, on a 400 MHz
 Bruker Avance III HD NanoBay, equipped with 1H/BB z-gradient probe (BBO, 5 mm), nuclear magnetic
 resonance (NMR) spectrometer. All spectra were processed using TOPSPIN 3.6.2. The standard pulse
 sequences supplied by Bruker were utilized for all 1D and 2D NMR analyses. The phasing of two-

dimensional ¹H-¹³C HSQC NMR spectra was manually corrected if required. Samples were analyzed at
 20 mg/mL in D₂O containing 1 vol% d-TFA.

382 4.3. Relative molecular weight determination

383 The relative molecular weight to pullulan standards of the starting chitosan together with the obtained 384 products was determined via LC/GPC ELSD. For the calibration curve, 1 mg of a pullulan standard (342 385 Da, 1 kDa, 6 kDa, 10 kDa, 50 kDa, 110 kDa, 200 kDa, 400 kDa and 800 kDa, PSS standards kit) was 386 dissolved in 1 mL of an 0.1 vol% TFA aqueous solution to obtain a final concentration of 1 mg/mL. For 387 the chitosan samples, 5 mg of the sample was dissolved in 5 mL of an 0.1 vol% TFA aqueous solution 388 to obtain a final concentration of 1 mg/mL. All solutions were left overnight to equilibrate and filtered 389 using a 25 mm polyethersulfone (PES) syringe filter (0.22 µm membrane) before injection. 20 µL was 390 automatically injected into an Agilent 1260 Infinity II HPLC system equipped with PSS NOVEMA Max 391 analytical Linear M columns (10 µM particle size). A guard column and two GPC columns (8 x 300 mm) 392 with a pore size of 100 Å, 3000 Å and 3000 Å were used respectively. The analyses were conducted at 393 25 °C with a flow rate of 0.5 mL/min. Detection was done using an Agilent 1260 Infinity ELSD detector, 394 the evaporation temperature was set at 80 °C and the nebulizer temperature was at 42 °C with an 395 evaporator gas flow of 2.5 standard liters per minute (SLM). All raw data were directly processed via 396 Matlab R2023b (Sadao Mori, 1999).

397 4.4. Synthesis of *N*-alkyl chitosan

398 Two grams of chitosan was dissolved in 200 mL distilled water containing 1 vol% acetic acid and 0.05 399 w/v% cetylpyridinium chloride in a 500 mL round bottom flask by an overhead stirrer at 800 rpm. 400 Afterward, 0.125 eq. of aldehyde (0.5 eq. for acetone) was added and the mixture was stirred for 1 401 hour. Next, 113 mg (0.1 eq.) of PICB was added and the mixture was stirred overnight. The following 402 day, 20 mL of MeOH was added and the pH was adjusted to 3 with 1 N HCl. Subsequently, the mixture 403 was stirred at 800 rpm for 2 hours at 60 °C to quench any residual PICB. Afterward, the chitosan 404 derivative was precipitated by pouring the reaction mixture into 600 mL of acetone. Finally, the 405 precipitate was collected via vacuum filtration, dried overnight at 50 °C and ground into a fine powder.

406 4.5. Oxidative depolymerization of (*N*-alkyl) chitosan

407 The oxidative depolymerization of (*N*-alkyl) chitosan was performed as previously described (Chapelle 408 et al., 2019). Briefly, 62.5 mg of (*N*-alkyl) chitosan was dissolved overnight in 5 mL demineralized water 409 containing 19.4 μ L acetic acid. Next, 625 μ L of a freshly prepared solution containing 0.2 eq. NaNO₂ 410 was added and the mixture was stirred at 50 °C for three hours. Finally, as the obtained COS no longer 411 precipitated upon the addition of acetone, COS were obtained after evaporation of the solvent.

412 4.6. Acid hydrolysis of (*N*-alkyl) chitosan

60 mg of (*N*-alkyl) chitosan was dissolved in 2 mL of 2 N HCl in a 20 mL capped vial. Afterward, the mixture was heated under reflux conditions for 24 hours. Next, the mixture was poured into 10 mL of acetone and the obtained precipitate was collected via centrifugation, dried overnight at 50 °C and ground into a fine powder.

417 CRediT authorship contribution statement

Casper Van Poucke: Conceptualization, Methodology, Formal analysis, Investigation, Data Curation,
Writing - Original Draft, Writing – Review & Editing, Visualization, Project administration, Funding
acquisition. Evert Verdegem: Investigation, Formal analysis. Sven Mangelinckx: Conceptualization,
Methodology, Validation, Resources, Data Curation, Writing - Review & Editing, Supervision. Christian
V. Stevens: Conceptualization, Methodology, Validation, Resources, Data Curation, Writing – Review &
Editing, Supervision, Project administration, Funding acquisition.

424 Declaration of competing interest

- 425 The authors declare that they have no known competing financial interests or personal relationships
- 426 that could have appeared to influence the work reported in this paper.

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- 431 Data availability
- 432 Data will be made available on request.
- 433 References

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