A new method for rapid degree of substitution and purity determination of chloroform-soluble cellulose esters, using $^{31}$P NMR†

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Chloroform-soluble palmitic and decanoic acid esters of cellulose were synthesized from the reaction of MCC with acid chlorides in LiCl/DMA and the ionic liquid [amim]Cl, as novel cellulose solvents. A process of derivatization of the remaining hydroxyl groups, as phosphate esters and subsequent $^{31}$P NMR analysis, allowed for simultaneous degree of substitution (DS) determination and quantification of the aqueous-quench acid by-product impurity, after the appropriate calculation. The full mathematical treatment for DS determination is presented, including scripts for the Python and Java programming languages for rapid interpretation of results. This method for DS determination was validated against traditional analyses of the palmitoyl cellulose and the fully substituted $p$-nitrobenzoyl palmitoyl diester product, from additional reaction with $p$-nitrobenzoyl chloride. DOSY NMR and SCORE analyses were also employed to demonstrate the utility of this rapid $^{31}$P derivatization and analytical process over traditional 1D NMR analyses.

**Introduction**

Cellulose, derived from natural sources (commonly bacteria and plants), has been long promoted as an abundant renewable resource for advanced materials. Cellulose is chemically modified at its C-2, C-3 and C-6 hydroxyl groups to form esters, ethers, carbonates or carbamates. However, determination of accurate values for degree of substitution (DS) and regioselectivity in the functionalization of cellulose can often be a complex task. This becomes more troublesome the more complex the reaction pathway or the substituent is, but is absolutely crucial for the characterisation of novel and technically advanced materials. Optimisation of these reactions from novel polar media such as lithium chloride/N,N-dimethylacetamide (LiCl/DMA) or imidazolium-based ionic liquids (ILs) such as 1-allyl-3-methylimidazolium chloride ([amim]Cl) is significantly more challenging than analysis of a purified product. A typical case in point is the esterification of cellulose with long-chain ‘fatty’ acid chlorides (Fig. 1). Several problems exist for the efficient optimization of these reactions using traditional DS determination methods such as gravimetric analysis, $^1$H NMR, elemental analysis (EA), IR or GPC. The main problem occurs during reaction workup. At this stage, addition of a quenching reagent, typically water, destroys any remaining reactive electrophile (e.g. acid chloride or anhydride) converting it to carboxylic acid. The main role of the water is to precipitate the product and thus separate it from the liquid reaction media as many solvents for cellulose chemistry are comprised of compounds with low vapour pressures (e.g. ionic liquids, LiCl, and DMA). Co-precipitation of product and by-products undoubtedly occurs, often requiring tedious purification procedures. Furthermore, this is compounded by the fact that $^1$H NMR, as the best method for DS determination at present, can suffer from poor resolution of by-products from the anhydroglucose (AGU) unit and main substituent resonances, frequently causing inaccuracies in DS determination. This is particularly a problem for long-chain acyl substituents such as fatty acid esters. Building upon traditional methods, recent articles have appeared based upon saponification of weighed samples of cellulose esters under standardised conditions, followed by GC analysis of the liberated acids. These techniques, however, also suffer from the fact that the samples must be acid free to give accurate results. One method, recently published, that attempts to alleviate these problems, is an improved IR analysis procedure. This involves resolution of the free acid and ester carbonyl stretches using monoethanolamine to form a conjugate base–acid pair, with the

![Fig. 1](image-url) - Typical fatty acid esterification of cellulose to give the ester and acid by-product, from [amim]Cl I, as reaction media.

† Electronic supplementary information (ESI) available: Python and Java scripts for determination of DS values, including spectroscopic data for the cellulose ester products, for a visual measure of sample quality. See DOI: 10.1039/c0ay00336k

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free acid in the analytical sample and leaving the esters untouched. The C=O stretch of the conjugate base–acid pair is shifted out of the region of the ester allowing increased accuracy in acid impurity determination. The IL 1-ethyl-3-methylimidazolium acetate ([emim][OAc]) was used for dissolution of their samples, as one of a range of accessible ILs capable of dissolving wood and cellulose, allowing for analysis of esters of cellulose in the low to medium range (0–1.5) in addition to those which are fully organically soluble.

Despite the high novelty and utility of this method, IR analysis is still limited in its accuracy and in the information that it provides. However, the use of cellulose-dissolving media, such as ILs, in the analysis procedure is an important concept as it may allow for the analysis of normally insoluble products such as low DS cellulose esters.

In the present publication, it is our objective to validate and demonstrate the utility of 31P derivatization and NMR analyses for the optimisation of cellulose functionalization reactions, in comparison to traditional methods. For this, we have initially chosen a short set of long-chain cellulose esters in the organically soluble DS range of 1.5 to 3 and of variable purity. Based upon our existing and growing knowledge of wood and wood biopolymer dissolution and functionalization in ILs, aided by 31P labelling and NMR analysis, it is our ultimate goal to develop this technique for the study of a wider range of wood biopolymer chemical modification reactions.

**Experimental**

**Materials**

Allyl chloride, 2-chloro-4,4,5,5-tetramethyl-1,3-dioxaphospholane (2-Cl-TMDP), anhydrous N,N-dimethylacetamide (DMA), endo-N-hydroxy-5-norbornene-2,3-dicarboximide (e-HNDI), anhydrous lithium chloride (LiCl), microcrystalline cellulose (MCC), N-methylimidazole and p-nitrobenzyol chloride were purchased from Aldrich (Finland), and were used without further purification. Palmitoyl and decanoyl chloride were purchased from Aldrich (Finland), and were used without further purification. [amim]Cl I was prepared according to a previous report.

**Measurements**

Quantitative 31P and 13C NMR spectra were recorded at 27 °C using inverse gated proton decoupling sequences on a Varian Unity Inova 600 spectrometer (600 MHz proton frequency) equipped with a 5 mm direct detection broadband probe-head. Quantitative 31P spectra were collected with 512 transients using 90° pulse flip angle, 80 000 Hz spectral width, 1 s acquisition time, and 10 s relaxation delay. The quantitative 13C spectrum was collected with 20 000 transients utilizing a 90° pulse flip angle, 36 000 Hz spectral width, 1 s acquisition time and 60 s relaxation delay. Quantitative 1H spectra and DOSY (diffusion-ordered spectroscopy) data were recorded at 27 °C on a Varian Unity Inova 500 spectrometer (500 MHz proton frequency) equipped with 5 mm triple-resonance (1H, 13C, and 15N) gradient probe-head. For 1H measurements, 60 transients were collected with 45° pulse flip angle, 1.9 s acquisition time, and 13 s relaxation delay. Spectral width for the 1H spectra was 8 000 Hz.

DOSY datasets were recorded using Bipolar Pulse Pair Stimulated Echo pulse sequence (BPPSTE) using 1 ms gradient duration, 0.2 ms gradient recovery delay, 150 ms diffusion time, 224 transients, 1 s acquisition time, 2 s relaxation delay, and spectral width of 8 000 Hz. In DOSY datasets, 60 diffusion gradient amplitudes were used (ranging from 0.5 to 20 G cm−1).

DOSY data were analyzed with the Speedy COnponent REsolution (SCORE)-algorithm incorporated into the DosyToolBox 0.53-software running on Matlab 7.5.0 (MathWorks, Natick, MA, USA). SCORE analysis was carried out with standard parameters over the region of 0–6 ppm for a 2-component system.

**Methods**

**Synthesis of cellulose esters**

*High purity palmitoyl cellulose 2.* Microcrystalline cellulose (MCC, 0.617 g, 3.81 mmol) was heated in DMA (anhydrous, 30 ml) at 130 °C for 3 h under inert atmosphere. The sample (2%/w/w in solution) was allowed to cool to 90 °C upon which LiCl (anhydrous, 2.080 g, 6.73% w/w in solution) was added in one portion and the sample allowed to cool to room temperature with stirring. The sample was allowed to stir for a further 18 h under inert atmosphere before addition of palmitoyl chloride (2.5 ml, 8.24 mmol) and pyridine (anhydrous, 1.2 ml, 14.90 mmol). The resulting mixture was allowed to stir for 72 h at room temperature and under inert atmosphere. The reaction mixture was poured into methanol (200 ml) and allowed to stir for 30 min. The mixture was filtered. The filtrand was dissolved in chloroform (100 ml) and precipitated by pouring into methanol (300 ml). The solution was filtered and the filtrand washed with methanol (100 ml) and deionised water (100 ml). The resulting solid was dried under vacuum at 55 °C for 18 h to give high purity palmitoyl cellulose 2 (1.97 g, 219% WPG (weight % gain), 92% yield at a DS of 1.67), as a white powder (found: C, 69.87, H, 11.0). Calc. for C32.72H60.10O6.67 (DS 1.67): C, 70.1, H, 10.8%; νmax (ATR)/cm−1 3481, 2922, 2851 and 1742; δ6H (500 MHz, CDCl3, Me4Si) 0.68–1.09 (3H, br t, J 6.8, CH3), 1.09–1.98 (26H, br m, COCH3(CH2)7CH2), 1.98–2.70 (2H, br m, COCH2(CH2)3CH2), 2.70–5.62 (7H, br m, AGU); δ31P: 1.67 ± 0.012 (calculated from 31P NMR analysis of the phosphitylated product, performed in triplicate); δH: 1.65 (calculated from 1H NMR analysis); δC: 1.62 (calculated from C-content determined from the EA); decanoic acid impurity: 0.94 ± 0.078%/w/w (calculated from 31P NMR analysis of the phosphitylated product, performed in triplicate).

*High purity p-nitrobenzoyl-palmitoyl cellulose 3.* Palmitoyl cellulose (2, 100 mg, 0.179 mmol) was dissolved in a mixture of chloroform : pyridine (1 : 1, 2 ml). p-Nitrobenzoylchloride (150 mg, 0.809 mmol) was added in one portion and the mixture was allowed to stir at room temperature for 18 h under inert atmosphere. Methanol was added until the product precipitated and the solution became clear. The solution was removed with a pipette and this precipitation process repeated 2 times. The solid was then finally precipitated from chloroform : hexane (1 : 1) with methanol addition. The solution was decanted and the product dried under vacuum at 55 °C for 18 h to give high purity p-nitrobenzoyl-palmitoyl cellulose 3 (128 mg, 28% WPG.
and the mixture agitated for 1 min. The sample was heated at 2500 rpm, using an Janke & Kunkel Vibrofix VF1 Electronic Orbital shaker. Decanoyl chloride (640 μl, 3.09 mmol) was added in one portion and the solution was vortexed for 10 s). 31P NMR spectra (243 MHz for calculation of the free hydroxyls per weight unit of substrate (OHg, mol g⁻¹)) is as follows:

\[ \text{OH}_g = \frac{\text{IS}_{\text{mol}} \times \text{IS}_{\text{vol}} \times \text{IR}}{1000000 \times \text{W}_s} \]  

Typical 31P NMR analysis procedure. CDCl₃ (1 ml) was added to palmityl cellulose (2, 25.0 mg) in a 10 ml screw-top vial. The sample was agitated for ∼5 min until the sample was fully dissolved. Pyridine (150 μl, 1.87 mmol) was added in one portion and the sample agitated until visibly homogeneous (∼5 s). The sample was allowed to cool to room temperature, whereby 2-Cl-TMMDP (200 μl, 1.26 mmol) was added in one portion and the solution was vortexed (∼5 s). 31P NMR spectra (243 MHz for 31P) were recorded with 700 μl samples, in a 5 mm o.d. NMR tube.

31P NMR DS and impurity determination. 31P labelling and NMR analysis is already an established method for analysis of lignin functional groups. These reports have demonstrated the ability of 31P NMR analysis with the 2-Cl-TMMDP as phosphitylating reagent for the resolution of aliphatic, individual phenolic and individual carboxylic acid resonances. Although aliphatic resonances are clearly resolved from the individual phenolic resonances, the resolution of individual aliphatic resonances is poor, under the present reagents and conditions, providing only limited information about regioselectivity of reaction on cellulose, or other biopolymers. The ability of this technique to distinguish the 3 main C–OH hydroxyl types over a substantial ppm range (20 ppm) offers great flexibility for the analysis of a wide range of functionalities. An equation for percentage DS determination from 31P NMR analysis has previously been published by us, in relation to wood analysis from pre-dissolution into ILs, however, no report detailing or validating this method has yet been published. A minor error in the published mathematical treatment also exists, which we would like to take the opportunity to correct. Consequently this error has little effect on the calculated values.

A description for the calculation of DS values and impurities from cellulose functionalization reactions follows; the experimental variables required for DS determination from phosphate ester derivatization and 31P NMR analysis are: (1) MWₙ (substituent molecular weight). This is the molecular weight (g mol⁻¹) of the substituent, not including the linking oxygen atom (between the cellulose backbone and the substituent); (2) ISvol (internal standard volume, µl); (3) ISₘₐₜ (internal standard molarity, mmol); (4) IR (integration ratio of remaining functionalized cellulose hydroxyls against internal standard); and (5) Wₖ (sample weight, mg). In addition, the constants DSₘₐₜ (maximum achievable DS value of 3 for unsubstituted cellulose) and OHₐ (the free hydroxyls per weight unit of cellulose, 3 × 162 = 0.01852 mol g⁻¹) are also required. The eqn (1) for calculation of the free hydroxyls per weight unit of substrate (OH₈, mol g⁻¹) is as follows:

\[ \text{OH}_8 = \frac{\text{IS}_{\text{mol}} \times \text{IS}_{\text{vol}} \times \text{IR}}{1000000 \times \text{W}_s} \]  

The final eqn (2) required to calculate the DS from 31P labelling and NMR analysis (DSₚₚ), is as follows:

\[ \text{DS}_{31P} = \frac{\text{DS}_{\text{max}}}{1 - \frac{\text{OH}_8}{\text{OH}_C}} \]  

In addition to calculating the DS of pure products, the integration ratio (against internal standard) for any impurity (I₁), e.g. carboxylic acids, in combination with the molecular weight of the impurity (MW₁, g mol⁻¹) can be used to determine the weight of the impurity (W₁, mg) in the original sample. This is determined using eqn (3):

\[ \text{W}_1 = \frac{\text{IS}_{\text{mol}} \times \text{IS}_{\text{vol}} \times \text{MW}_1 \times \text{I}_1}{1000 000 000} \]  

If impurity exists in a sample a corrected DS value can be eventually attained by simply subtracting W₁ from the original sample weight (Wₕ) and recalculating OHₘ, as shown in eqn (4):

\[ \text{OH}_8 = \frac{\text{IS}_{\text{mol}} \times \text{IS}_{\text{vol}} \times \text{I}_R}{1000 000 \times (\text{W}_s - \text{W}_1)} \]  

The above equations may be adapted to starting polymers other than unsubstituted cellulose, for any given reaction. A good example of this is if an additional substituent is added to an existing substituted cellulose product. In this case, if you have predetermined the DS value of the initial substituent, by 31P labelling and NMR analysis, one can analyse the product of the second substitution reaction in a similar manner. To determine...
the DS of the second substituent from these results you can replace OH\(_C\) in the second calculation of eqn (2) with OH\(_X\) from the first calculation of eqn (1), and for the second calculation of eqn (2), DS\(_{\text{max}}\) is replaced with the DS of the first substituent. The result will be DS\(_{\text{p}}\) values for both substituents from analysis of both products. Other non-cellulosic polymers or polymer mixtures, such as xylan, lignin, starch or wood, may also be analysed by replacing OH\(_C\) and DS\(_{\text{max}}\) values from their experimental or theoretically calculated values. As some polymers may have a non-integer number of hydroxyls per monomer unit, or non-regular monomer units, DS\(_{\text{p}}\) may be represented as a percentage substitution of the total sample by replacing DS\(_{\text{max}}\) with 100.

Eqn (1) and (2) have been incorporated in to scripts for the Python and Java programming languages\(^\text{13}\) in order to allow for rapid processing of data (see ESI\(^\dagger\)).

**Results and discussion**

As the objective of this article is to assess the \(^{31}\)P labelling and NMR analysis procedure, against traditional NMR analyses, suitable materials or reactions were required to demonstrate the advantages of using the \(^{31}\)P procedure over the traditional analyses, while still allowing for validation of the DS values against each other. To achieve this, a common acylation reaction in cellulose chemistry was chosen. That is esterification of cellulose with long-chain fatty acids.

**DS\(_{\text{p}}\) determination and method validation using high purity palmitoyl cellulose 2**

To initially determine the accuracy of the \(^{31}\)P method on a CDCl\(_3\) soluble sample, MCC was reacted to give palmitoyl cellulose 2, employing the standard method of dissolution of MCC into LiCl/DMA followed by reaction with palmitoyl chloride, in the presence of excess pyridine (Fig. 2). Care was taken to prepare the best quality \(^1\)H NMR spectra possible for DS determination (see ESI\(^\dagger\)).

As a secondary method, outside \(^{31}\)P labelling and analysis, the remaining hydroxyls in the palmitoyl cellulose product 2 were further esterified as \(p\)-nitrobenzoyl esters. This was achieved by quantitatively reacting the chloroform soluble starting material with the \(p\)-nitrobenzoyl chloride (Fig. 2) to give, after purification, fully substituted \(p\)-nitrobenzoyl-palmitoyl cellulose 3. The reaction was determined to be complete (almost no remaining free hydroxyl groups) by the absence of any OH-stretch in the IR spectra (see ESI\(^\dagger\)) or absence of any aliphatic phosphate ester resonances, after \(^{31}\)P derivatization and NMR analysis. In addition to providing an alternative DS\(_{\text{p}}\) value, this functionality also provided the opportunity to analyse the product by EA, observing both carbon and nitrogen percentages, and quantitative \(^{13}\)C NMR, with the carbonyl resonances providing a suitable region for integration and DS\(_{\text{p}}\) determination (see ESI\(^\dagger\)). The inclusion of the chromophore also would allow for GPC with UV detection, although this is not a suitable method for determining DS values. Although it was possible to determine DS\(_{\text{p}}\) by straightforward integration of the carbonyl resonances, with sufficient exponential line broadening, the signal to noise ratios and resolution, after 20 000 transients (~66 h collection time), were not adequate enough to allow for accurate determination of regioselectivity, as has been previously observed for cellulose acetates.\(^\text{14}\)

\(^{31}\)P derivatization and NMR analysis of palmitoyl cellulose 2, shown in Fig. 3, were performed by dissolution of the substrate 2 into CDCl\(_3\) and reaction with 2-Cl-TMDP under basic (pyridine) conditions. The internal standard \(e\)-HNDI was also included in the phosphorylating mixture, which also reacted with the phosphate acid chloride (2-Cl-TMDP) or anhydride (TMDP-anhydride), in solution, to form its corresponding phosphate ester (\(e\)-HNDI-TMDP). The spectrum was calibrated with TMDP-anhydride at 132.2 ppm. \(e\)-HNDI-TMDP resonates at 152.0 ppm under these solvent conditions. Alkoxy phosphate esters (alkoxy-TMDP) are located in the region between 151.5 and 143.5 ppm and both decanoic acid and palmitic acid-phosphate mixed anhydrides (decanoate or palmitate–TMDP) resonate at 134.9 ppm. The reaction was performed in triplicate and the original sample weights were corrected, based upon the presence of 0.94\% w/w palmitic acid impurity (as determined by the \(^{31}\)P NMR integrations of palmitate–TMDP). DS\(_{\text{p}}\) for this compound 2 was determined to be 1.67 with a standard deviation for the full procedure of 0.012. It is expected that as you approach DS values close to 0 or have very large substituents, the standard deviation will increase although for the chloroform soluble regime (DS ca. 1–3) the error will remain approximately the same.

From the initial palmitoyl cellulose substrate 2 and its subsequent characterization, the complete set of determined DS values are listed in Table 1, including short self-explanatory descriptions of the methods used to obtain them. From a comparison of these methods, we can see that EA predictably shows the most

![Fig. 2 Synthesis of palmitoyl cellulose 2 and \(p\)-nitrobenzoyl-palmitoyl cellulose 3.](image-url)
deviation from the mean value. This is very much dependent on both purity of materials and skill of the analyst. Due to the simplicity of integration, and the ability to resolve low molecular weight hydroxylated impurities from the polymeric material in solution, \(^{31}\)P derivatization and NMR analysis offer a more accurate and thorough characterization method, in combination with the traditional methods.

DS\(_{31P}\) determination and DOSY analysis of crude decanoyl cellulose 5

In order to demonstrate the true value of \(^{31}\)P derivatization and NMR analysis, e.g., in the optimization of cellulose acylation reactions, crude decanoyl cellulose 5 was synthesized, by carrying out a similar procedure to that used for the synthesis of palmitoyl cellulose 2, but avoiding any purification steps, beyond quenching the reaction by heating with water.

As ILs, such as [amim]Cl, are predicted to be potential environmentally benign media for cellulose chemistry, based upon their recyclability and low vapour pressure, [amim]Cl was chosen as media for this reaction. A typical workup procedure for cellulose reactions, from ILs, involves stirring with a solvent such as water to quench the reactants and precipitate the product. Additionally, heating with the ‘quenching’ solvent is often required to remove traces of IL from the product. This procedure will typically preserve the yield and polydispersity of the product by precipitating most of the functionalised polymer provided the functionality is sufficiently hydrophobic, as is the case with most acyl species. This will also co-precipitate a large portion of the quenched reactants, with the desired product. These co-precipitated species are often impossible to resolve using traditional \(^1\)H NMR analysis, preventing rapid, consistent and reliable optimisation of these reactions.

The crude decanoylated cellulose product 5 was subjected to DOSY NMR to attempt to resolve any small organic species, including the decanoic acid quenching by-product, from the functionalized biopolymer. Inspection of the SCORE analysis results (Fig. 4), for a 2-component system, clearly shows resolution of the decanoic acid quenching by-product from the desired product 5. The SCORE analysis estimates this impurity at 20.2% mol/mol (of hydrogen) which, when converted into % w/w, gives a value of 18.6% w/w. From the extracted product 5 spectra (Fig. 4), no accurate DS determination is possible.

When the same mixture was phosphitylated with 2-Cl-TMDP and analysed by \(^{31}\)P NMR, according to the above method, accurate values of 25.2% w/w for decanoic acid impurity and 2.29 for DS\(_{31P}\) of the impure product were rapidly determined (see ESI†). A proper comparison between the % w/w impurity

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Table 1 DS of palmitoyl cellulose 2 based upon \(^1\)H NMR (DS\(_{31H}\)), \(^{13}\)C NMR (DS\(_{13C}\)) or elemental (DS\(_{EA}\)) analyses of the parent compound 2, the \(p\)-nitrobenzoylated product 3 or TMDP-phosphitylated product 4, by \(^{31}\)P NMR (DS\(_{31P}\)) analysis

<table>
<thead>
<tr>
<th>Analyte</th>
<th>Method</th>
<th>Integral regions/element</th>
<th>Integral regions (ppm)</th>
<th>Calculation</th>
<th>DS</th>
</tr>
</thead>
<tbody>
<tr>
<td>2</td>
<td>DS(_{31H})</td>
<td>COCH(_3)(CH(_2))(_2)CH(_3) vs. AGU region</td>
<td>0.5–1.9 vs. 2.5–5.5</td>
<td>((CH(_2))(_2))/(AGU/7)</td>
<td>1.65</td>
</tr>
<tr>
<td>2</td>
<td>DS(_{EA})</td>
<td>Calculated from C-content</td>
<td>—</td>
<td>—</td>
<td>1.62</td>
</tr>
<tr>
<td>3</td>
<td>DS(_{31H})</td>
<td>Ar(_H) vs. AGU regions</td>
<td>8.5–7.5 vs. 2.7–6.0</td>
<td>3-(Ar(_H))/(AGU/7)</td>
<td>1.67</td>
</tr>
<tr>
<td>3</td>
<td>DS(_{13C})</td>
<td>COAr region vs. COAlk</td>
<td>162.7–164.4 vs. 172.0–173.5</td>
<td>COAlk × 3/(COAr + COAlk)</td>
<td>1.67</td>
</tr>
<tr>
<td>3</td>
<td>DS(_{EA})</td>
<td>Calculated from C-content</td>
<td>—</td>
<td>—</td>
<td>1.53</td>
</tr>
<tr>
<td>4</td>
<td>DS(_{31P})</td>
<td>e-HNDI-TMDP vs. alkoxy-TMDP</td>
<td>151.5–152.5 vs. 151.0–142.0</td>
<td>See eqn (1) and (2)</td>
<td>1.67</td>
</tr>
<tr>
<td>2</td>
<td>DS(_{mean})</td>
<td>Mean analysis value</td>
<td>—</td>
<td>—</td>
<td>1.65</td>
</tr>
</tbody>
</table>
values from the $^{31}$P and DOSY experiments is not meaningful as the DOSY procedure is unable to produce quantitative data. When the DOSY spectrum for the palmitoylated cellulose sample 2 was collected, it was not possible to process the data, using the SCORE algorithm, with the purpose of obtaining any meaningful results that would allow for quantification of the palmitic acid impurity. This was due to the very low quantity of acid by-product impurity present in the sample. On the other hand, however, the $^{31}$P NMR analysis method allowed for determination of this impurity to be 0.94 $\pm$ 0.078% w/w. The significant advantage of DOSY over $^1$H NMR or $^{31}$P NMR seems to be only in a qualitative discrimination between two abundant components in a mixture, where one component is a high molecular weight polymer. This lack of quantitivity with DOSY is mostly dictated by differences in $T_1$ and $T_2$ relaxation that occur during the DOSY pulse sequence. $T_1$ in particular can be greatly affected by the molecular weight of the resonating species. This $^{31}$P-based analysis method limits the optimisation of these reactions to a aqueous quench, drying of the crude product and one in situ derivatization and NMR analysis procedure, to simultaneously determine DS and purity of the product. This is unrivalled by any other NMR procedure, such as DOSY or $^1$H NMR, which must serve as complementary qualitative techniques, due to long collection times and poor resolution, in the analysis of these crude reaction mixtures.

Conclusions

A new method for DS determination of functionalized cellulose products has been developed, based upon derivatization of the remaining hydroxyl groups, as phosphate esters. Under standardized solvent and $^{31}$P NMR acquisition conditions, these phosphate esters can be quantitatively integrated against an internal standard, allowing for calculation of the DS of the functionalized cellulose starting material. This method is at the moment applicable to the analysis of chloroform soluble products and was validated against a palmitoylated cellulose sample, which was analysed in detail by standard DS determination methods. A crude decanoylated cellulose reaction product was also analysed by this $^{31}$P derivatization and NMR procedure, allowing for rapid determination of both decanoic acid impurity and DS of the unpurified product. Within a potential comprehensive list of polymeric substrates and chemical modifications, this study initially highlights the potential utility of this method at least for optimising cellulose acylation reactions, with a view to more widespread application and process optimization.

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