## BOOK OF EXTENDED ABSTRACTS

## THE 42<sup>nd</sup> ANNUAL SYMPOSIUM ON POLYMER SCIENCE/ENGINEERING

University of Waterloo Waterloo, Ontario

September 2<sup>nd</sup>, 2020

## INSTITUTE FOR POLYMER RESEARCH

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# 2020

# PROGRAM

#### INSTITUTE FOR POLYMER RESEARCH CELEBRATING 35 YEARS OF OFFICIAL INSTITUTE STATUS FORTY-SECOND ANNUAL SYMPOSIUM ON POLYMER SCIENCE/ENGINEERING 2020 University of Waterloo, Waterloo, Ontario Wednesday, September 2, 2020

12:50	Welcome and Opening Remarks				
1:00 – 1:20	Janine Thoma [Prof. Duhamel]				
	Persistence Length of Polymeric Bottle Brushes Determined by Pyrene Excimer				
	Fluorescence				
	(Winner of 2020 IPR Award for Academic Excellence in Polymer Science/Engineering)				
1:20 – 1:40	Elnaz Esmizadeh [Prof. Mekonnen]				
	Degradation benavior of polypropylene during reprocessing and its				
	biocomposites: thermal and oxidative degradation kinetics				
1:40 - 2:20	5-Min. Mini Presentations				
	1) Sainiwetha Saihrishnan [Prof. Mekonnen]				
	Thermo-mechanical Degradation of Polypropylene-Low Density				
	Polyethylene blend system after multiple extrusion				
	2) Javan Buratynski [Prof. Schipper]				
	Using Degradable Polymers to Separate Carbon Nanotubes				
	3) Abdullah Ba Salem [Prof. Duhamel]				
	A) Kristijan Lulic [Prof. Dubamol]				
	Self-Association of Oligoquinoline Foldamers Probed by Eluorescence				
	Anisotropy				
	5) Hunter Little [Prof. Duhamel]				
	Progress in Instrumentation for Time-resolved Fluorescence Anisotropy				
	6) Franklin Frasca [Prof. Duhamel]				
	Relating Pyrene Excimer Fluorescence to Conformation in Pyrene-Labeled				
	Polyamines				
	7) Tiana Trumpur [Prof. Forrest]				
	Solvent Induced Nanoscopic Roughness on Glassy Polymer Thin Films				
	8) Minghui Liu (Prof. Zhao)				
	Antimicrobial Activity of Quaternary Ammonium Compound/Polyurethane				
	[QAC/PU] Colloidal Complex Film Based on Synergetic Release Killing and				
	Contact Killing Mechanisms				
2:20 - 2:40	Remi Casier [Prof. Duhamel]				
	Copolymerization and its Effects on Polymer Dynamics in Solution				
2:40 – 3 :00	Break				

Mini-Symposium on Modification, Characterization, and Applications of Polysaccharides					
3:00 - 3:20	Chunxia Tang [Prof. Tam] Functionalized Cellulose Aerogel Beads for Heavy Metal Ions Removal (Winner of the 2020 IPR Award for Academic Excellence in Polymer Science/Engineering) Natun Dasgupta [Prof. Gauthier] Thermoresponsive Starch Nanoparticles for Oil Recovery from Tar Sands				
3:20 - 3:40					
3:40 - 4:00	<ul> <li>5-Min. Mini Presentations</li> <li>9) Maryam Bagheri [Prof. Simon] Polysaccharide-Ionic Liquid corrosion inhibitors</li> <li>10) Ewomazino Ojogbo [Prof. Mekonnen] Effects of fabrication method on the dispersion of CNCs in highly crosslinked rubber composites</li> <li>11) Sanjay Patel [Prof. Duhamel] Pyrene-labeled Starch Nanoparticles as Fluorescent Sensors for Explosive Detection</li> <li>12) Donghan Liu [Prof. Duhamel] Synthesis and Characterization of Furan Based Non-ionic Surfactant (FBNIOS)</li> </ul>				
4:00 – 4:20	Joanne Fernandez [Prof. Gauthier] Grafting of Starch Using a Complex Initiation System of CAN-KPS				
4:20 – 4:40	Muhammad Shahidul Islam [Prof. Tam] Cellulose Nanomaterials: Synthesis, Properties, and Applications				
4:40 - 5:00	Damin Kim [Prof. Duhamel] Better Understanding the Structure of Glycogen with Pyrene Excimer Fluorescence				
5:00	Closing Remarks				

# EXTENDED ABSTRACTS

Janine Thoma Chemistry Waterloo

## Persistence Length of Polymeric Bottle Brushes Determined by Pyrene Excimer Fluorescence

Winner of the 2020 IPR Award for Academic Excellence in Polymer Science/Engineering

#### Investigating the Conformation and Flexibility of Poly[oligo(ethylene glycol) Methacrylate] Polymeric Bottle Brushes in Solution

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#### Introduction

Polymer architecture is a key component in the design of a polymer because it dictates its physical properties and consequently, its desired applications. Branched polymers constitute a large family of macromolecules with complex architectures that can be separated into four main categories: star, dendritic and hyper-branched, brush, and network/gel polymers. This study focuses on the third category, namely brush polymers. Developments in the synthesis of brush polymers were spearheaded by the growing number of their applications. For example, they are used for drug delivery in the biomedical industry and to prepare new super-soft materials.<sup>1,2</sup> Brush polymers, also called polymeric bottle brushes (PBBs), offer functionality with respect to backbone and macromonomer chemical composition which allows for their customization depending on the specific property requirements of a given application. What gives a PBB its unique properties are its high grafting density and high degree of polymerization.

Contrary to studies of PBBs by microscopy where the macromolecule is adsorbed onto a 2-D surface, the characterization of PBBs in solution is of interest because it allows scientists to probe the flexibility of these densely packed systems in their native conformation. A PBB will adopt a different conformation, either a polymer coil or an extended rod, depending on whether it is prepared with a short or long macromonomer, respectively. The Kratky-Porod (KP) Worm-like chain model enables the derivation of Equation 1 which accounts for the relative extension of a polymer by relating its average end-to-end distance,  $r_{EE}$ , to its persistence length,  $l_p$ .

$$r_{EE}^{2} = 2l_{\rm p}Nb - 2l_{\rm p}^{2} \left[ 1 - \exp(-Nb/l_{\rm p}) \right]$$
(1)

In Equation 1, N and b are the number of bonds and the bond length, respectively.  $l_p$  quantifies the bending stiffness of a polymer and represents the characteristic length required to decorrelate the orientation of the tangent to the contour length of the main chain from its original orientation as shown in Equation 2. In Equation 2,  $\theta(x)$  represents the angle between the tangent at the original position (x = 0) and at a position x along the chain contour length.  $l_p$  is equal to twice the Kuhn length,  $l_K$ . A polymer in an extended conformation, such as a PBB with long sidechains, will have a larger  $l_p$  than a polymer in a coil conformation, such as a PBB with short sidechains. A variety of techniques can be used to calculate  $l_p$  for a polymer, including static light scattering (SLS), small angle X-ray scattering (SAXS), small angle neutron scattering (SANS), atomic force microscopy (AFM) for a 2D worm-like chain adsorbed on a flat substrate, and for the first time in the scientific literature, pyrene excimer fluorescence (PEF).

$$<\cos[\theta(x)]>=\exp(-x/l_p)$$
 (2)

In order to use PEF to determine  $l_p$  for a series of PBBs, small quantities of the macromonomers 1-pyrenemethoxy-penta(ethylene glycol) methacrylate (PyEG<sub>5</sub>MA) was copolymerized with a series of methyl ether oligo(ethylene glycol) methacrylates (EG<sub>x</sub>MA with x = 0, 3, 5, 7, 9, and 19) to yield a series of pyrene-labeled PBBs (Py-PEG<sub>x</sub>MA). In turn, this report will illustrate how a combination of PEF, the fluorescence blob model (FBM),<sup>3</sup> and the KP model can be applied to calculate  $l_p$  for the Py-PEG<sub>x</sub>MA samples as a function of side chain length (x). This study takes advantage of the fact that the scaling laws describing the properties of an entire polymer can be applied to sub-volumes of the polymer coil referred to as *blobs*. *Blobs* are often used in polymer physics to break down a polymer into even sub-volumes.<sup>4</sup> Within the framework of the FBM, a *blob* is defined as the volume probed by an excited pyrene and it is defined by its number of structural units  $N_{blob}$ . Application of the KP model to the *blobs* derived from the FBM, allows  $l_p$  to be calculated for each Py-PEG<sub>x</sub>MA sample as shown in Equation 3.

$$r_{EE,blob}^{2} = 2l_{p}N_{blob}b - 2l_{p}^{2} \left[1 - \exp(-N_{blob}b/l_{p})\right]$$
(3)

In Equation 2,  $r_{EE,blob}^2$ ,  $N_{blob}$ , and *b* represent the averaged squared end-to-end distance of the polymer segment inside a *blob*, the number of structural units encompassed inside a *blob* obtained with the FBM, and the length added to the main chain contour length by one macromonomer, respectively. Since the same PyEG<sub>5</sub>MA macromonomer was used for each Py-PEG<sub>x</sub>MA sample, an excited pyrene probed the same volume during its lifetime resulting in a constant  $r_{EE,blob}^2$  for each sample as depicted in Figure 1. Since  $r_{EE,blob}^2$  could be determined for a fully extended Py-PEG<sub>x</sub>MA with x = 19, Equation 3 was applied to calculate  $l_p$  for each Py-PEG<sub>x</sub>MA sample.



Figure 1. Random coil (left) and extended (right) polymer conformations in solution.

#### **Experimental Procedure**

The synthesis of a Py-PEG<sub>x</sub>MA sample begins by preparing the pyrene labeled macromonomer (PyEG<sub>5</sub>MA) (Scheme 1). The chemical composition of Py-EG<sub>5</sub>MA was confirmed by <sup>1</sup>H NMR. Py-EG<sub>5</sub>MA was then copolymerized by conventional radical chain polymerization in tetrahydrofuran (THF) with the methyl ether oligo(ethylene glycol) methacrylate (EG<sub>x</sub>MA) macromonomers using a grafting through technique and azobisisobutyronitrile (AIBN) as the initiator to obtain the Py-PEG<sub>x</sub>MA samples.



Scheme 1. Synthesis of the Py-EG5MA macromonomer.

The Py-PEG<sub>x</sub>MA samples were characterized using gel permeation chromatography (GPC) to obtain their number  $(M_n)$  and weight  $(M_w)$  average molecular weight, and polydispersity (D) as well as atomic force microscopy (AFM) to visualize the shape of the Py-PEG<sub>19</sub>MA samples. **Table 1.** Chemical structure of each Py-PBB and the number of atoms in each side chain. Ns.

Sample	Py-	Py-PEG₃MA	Py-PEG₅MA	Py-PEG7MA	Py-	Py-
	PEG <sub>0</sub> MA				PEG <sub>9</sub> MA	PEG <sub>19</sub> MA
Structure				$\left(\begin{array}{c} 1 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \\$	$\left( \begin{array}{c} 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 $	
Ns	3	12	18	21	30	60

#### **Results and Discussion**

Six Py-PEG<sub>x</sub>MA samples were prepared by conventional radical copolymerization of PyEG<sub>5</sub>MA and different EG<sub>x</sub>MA macromonomers to yield the chemical structures shown in Table 1. Each Py-PEG<sub>x</sub>MA sample was prepared with 4-5 different pyrene contents. Each polymer was characterized using gel permeation chromatography (GPC) to determine their absolute  $M_n$  and  $M_w$  in dimethylsulfoxide (DMSO). Their degrees of polymerization were found to be greater than 100.

The time-resolved fluorescence decays of all Py-PEG<sub>x</sub>MA samples were acquired in DMSO. They were analyzed according to the FBM to yield  $N_{blob}$  and the  $N_{blob}$  value averaged over all pyrene contents,  $\langle N_{blob} \rangle$ , was plotted as a function of the number of atoms in the side chain (*N*s) in Figure 2A.



**Figure 2.** Plot of A)  $\langle N_{blob} \rangle$  versus the number of atoms in the sidechain,  $N_S$ , and B)  $l_p$  versus  $N_s$  for the series of Py-PEG<sub>x</sub>MA calculated by the ( $\circ$ ) FBM and ( $\times$ ) GPC analysis in aerated DMSO. ( $\Box$ ) PyEG<sub>5</sub>-PEG<sub>0</sub>MA, ( $\diamond$ ) PyEG<sub>5</sub>-PEG<sub>3</sub>MA, ( $\triangle$ ) PyEG<sub>5</sub>-PEG<sub>5</sub>MA, ( $\times$ ) PyEG<sub>5</sub>-PEG<sub>7</sub>MA, ( $\circ$ ) PyEG<sub>5</sub>-PEG<sub>9</sub>MA, and (+) PyEG<sub>5</sub>-PEG<sub>19</sub>MA.

Py-PEG<sub>0</sub>MA with the shortest sidechain was the most flexible of all PBBs and adopted a random coil conformation in solution. As a result, it had the largest  $N_{blob}$  value of  $63 \pm 7$ . Larger  $N_s$  values resulted in a stiffening of the main chain as reflected by a decrease in  $N_{blob}$  until  $N_{blob}$  plateaued for  $N_s$  values greater than 30. The plateau region in Figure 2A was attributed to the extended conformation adopted by the Py-PEG<sub>x</sub>MA samples for large *x* or  $N_s$  values. Since the same EG<sub>5</sub> spacer was used to link the pyrenyl label to the polymethacrylate backbone, all Py-PEG<sub>x</sub>MA samples had the same *blob* size given by  $r_{EE,blob}$  in Equation 3. Although  $r_{EE,blob}$  took the same value for all Py-PEG<sub>x</sub>MA samples, the *blobs* were occupied by fewer structural units as  $N_s$  increased up to the point where the polymethacrylate backbone was fully extended over the  $r_{EE,blob}$  length scale and  $N_{blob}$  remained constant and equal to  $16 \pm 1$  for  $N_s$  values greater than 30.

The persistence length,  $l_p$ , was then calculated using Equation 3 and plotted as a function of  $N_s$  in Figure 1B.  $l_p$  was found to scale as  $N_s^2$  in Figure 2B as expected for Gaussian side chains.<sup>5</sup> A more quantitative analysis of the  $l_p$ -vs- $N_s$  trends shown in Figure 1B was achieved by applying Equation 4 which was first introduced by Nakamura and Norisuye.<sup>6</sup> Equation 4 indicates that the stiffness parameter ( $\lambda^{-1}$ ) taken as either  $l_p$  or the Kuhn length ( $l_K = 2 \times l_p$ ) is the result of two contributions. These contributions to  $\lambda^{-1}$  are the stiffness ( $\lambda_0^{-1}$ ) induced by the main polymethacrylate chain and the stiffness ( $\lambda_b^{-1}$ ) induced by the excess free energy against bending due to the sidechains.<sup>6</sup>

$$\lambda^{-1} = \lambda_0^{-1} + \lambda_b^{-1} \tag{4}$$

 $l_p$  values calculated using the FBM for Py-PBBs in DMSO were compared to those obtained with a GPC equipped with a viscometer in DMSO (Figure 3A). The excellent agreement found between the  $l_p$  values obtained in DMSO by fluorescence and the GPC analysis was taken as validation of the fluorescence study. The  $l_p$  calculated for Py-PEG<sub>19</sub>MA from the GPC was found to deviate from the scaling law shown for the shorter side chains. This discrepancy could be attributed to a saturation effect which appears when the side chains, which adopt a random coil conformation, have fully occupied the local volume surrounding the polymethacrylate backbone and prefer to move to a less dense environment perpendicular from the polymer backbone. The break point observed in Figure 3A indicated the point where further increase in  $N_S$  would no longer affect  $l_p$ .



**Figure 3.** (A) Plot of  $l_p$  versus  $N_s^2$  for a series of PEG<sub>x</sub>MA calculated from the ( $\circ$ ) FBM and ( $\times$ ) GPC analysis. B) AFM height image of PEG<sub>19</sub>MA on a mica substrate at room temperature.

Finally, atomic force microscopy (AFM) was also used to visualize the PEG<sub>19</sub>MA sample in Figure 3B. Visual inspection of Figure 3B indicates that the PEG<sub>19</sub>MA chains are elongated but still flexible thus complementing the results obtained from a combination of PEF and FBM as well as qualitatively agreeing with the  $l_p$  of 4.5 nm ± 0.4 nm obtained from GPC analysis.

This study represents the first example in the literature where fluorescence was used to calculate  $l_p$  for a series of PBBs with increasing side chain length.

#### References

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Degradation behavior of polypropylene during reprocessing and its biocomposites: thermal and oxidative degradation kinetics

## Degradation behavior of polypropylene during reprocessing and its biocomposites: thermal and oxidative degradation kinetics

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#### Abstract

Non-isothermal thermogravimetric analysis (TGA) was employed to investigate the degradation of polypropylene (PP) during simulated product manufacturing in a secondary process and woodplastic composites. Kissinger–Akahira–Sunose (KAS), Ozawa- Flynn-Wall (OFW), Friedman, Kissinger, and Augis models were employed to calculate the apparent activation energy (E<sub>a</sub>). Experimental investigation using TGA indicated that the thermograms of PP recyclates shifted to lower temperatures revealing the presence of accelerated degradation process induced by the formation of radicals during chain scission. E<sub>a</sub> increased with the extent of degradation ( $\alpha$ ) and the dependency intensified by reprocessing cycles. In biocomposites, despite the detectable degradation steps of wood and PP in thermal degradation, a partial coincidence of degradation was observed under air.

**Keywords:** Polypropylene; Recycling; Thermogravimetric analysis; Wood-plastic composites **1. Introduction** 

Green awareness influences today's industry significantly in several ways to fulfill raising human preference for more eco-friendly procedures and products. Recently, the practice of recycling commodity polymeric materials such as polypropylene (PP) has been encouraged by the ever-growing desire to save resources, reduce the cost, and reuse waste material [1]. Wood-plastic composites (WPC), which typically contain wood (fiber, flour, pulp), plastic, and additives (e.g. coupling agents, processing aids), are substituting plastics, solid wood, and steel in applications such as construction, industrial and agricultural products [2,3]. Various grades of polypropylene[4,5] are extensively utilized as WPC matrices. WPCs can thermally decompose easier with induced degradability due to the inherent low thermal stability of wood-based reinforcements, desired for reducing environmental disadvantage [6]. However, the degradation of wood components during the thermo-mechanical processing of WPCs may lead to several undesirable properties in the fabricated products, such as deterioration of color, unpleasant odor, and poor mechanical properties[5,7].

The knowledge of degradation origins and kinetics both in oxidative and non-oxidative is of fundamental importance to determine the lifespan of the product, the effective conditions of processing, and how to control unfavorable degradation[8]. Thus, the objective of this work was to investigate the thermal and oxidative degradation kinetics using the various models. An internal batch mixer was employed to simulate recycling to avoid oxidation further.

#### 2. Materials and Methods

Polypropylene homopolymer powder (Pro-fax 6301) and Maple wood flour (mesh 600) from LyondellBasell (USA) and Ontario Sawdust Supplies (Canada), respectively, were used in this study. The composition of the WPCs was shown by  $PP^{\times n \text{ or } v} W_x$ , where v is virgin,  $\times n$  is n-time processed and x is 40 or 60 (wt.% of wood). The materials were compounded in a HAAKE Rheomix 3000 batch mixer at the mixing temperature of 180°C and rotor speed of 80rpm. Thermogravimetric analyses (TGA) were performed on a TA instrument Q500 analyzer both under nitrogen and air atmosphere with a 40 ml/min flow, respectively.

#### 3. Results and discussion

#### 3.1. Effect of reprocessing cycles

Figure 1 shows the thermogravimetric (TG) and the first-order derivative (DTG) curves at a heating rate of  $\beta$ =5 °C/min and 20 °C/min of the virgin PP and the samples reprocessed by batch mixing under non-oxidative and oxidative environment. As expected, increasing the heating rate led to a shift of the thermograms towards higher temperatures both in oxidative and non-oxidative degradation. This is due to the shorter retention time at the higher heating rates which limit time-consuming molecular motions taking place before decomposition [9]. All curves exhibit one-step degradation that can be attributed to the radical random scission commonly occurring with the thermal degradation of the polyolefins [10]. In a nitrogen environment, PP degrades in a single step beginning at 300 °C and ending at 475 °C. On the other hand, it degrades from about 250 to 425 °C primarily in a single step in the presence of air. Under both environments, all PP samples have degraded completely without leaving any significant residue. It can also evidently be observed that the thermo-oxidative decomposition took place at lower temperatures compared with the inert environment. Besides, the degradative effect of reprocessing cycles is more pronounced in the experiments under the air.



Figure 1. TG curves for the effect of reprocessing cycles on the thermal/oxidative decomposition of PP

The activation energy ( $E_a$ ) values versus  $\alpha$  for virgin PP and its recyclates decomposed under N<sub>2</sub> and the air calculated from various models were demonstrated in Figure 2. A gradual increase in  $E_a$  values with an increase in the degree of conversion was observed for all the samples. Lower  $E_a$ values at early stages of degradation suggest that at low conversions decomposition, kinetics was limited by initiation at the weak links. The increasing trend with the evolution of  $E_a$  can be interpreted by a shift in the rate-limiting step from initiation at the weak links to the degradation initiated by random scission [11].

#### 3.2. Effect of wood

TGA/DTG curves of the wood fiber and PP-based WPCs in nitrogen and air atmospheres are presented in Figure 3. Three distinct stages were for the thermal degradation of the well-dried wood fiber under a nitrogen atmosphere. In the beginning, a small step degradation occurred below 100 °C; followed by a second step that occurred at 220-315 °C and the last one occurred at 315-400 °C which can be related to the volatilization of the wood extractives; the decomposition of the hemicelluloses; and the decomposition of cellulose, respectively [6,12,13].



Figure 2. Activation energies (E<sub>a</sub>) for virgin PP and its recyclates obtained by different models

The other main component of wood fiber, lignin is the most stable component to thermal decomposition without any specific peak. However, in the case of oxidative degradation, a small peak that occurred at 400-500 °C, which can be related to the degradation of lignin. A clear indication is that in the WPC systems, the degradation of the two components can be easily detected individually owing to the noticeable difference in their thermal stability. A slight rightward shift in the degradation peak temperature of cellulose and PP is observed in the case of WPCs. Under the exposure of PP-based WPC to the air atmosphere, the oxidative degradation of all components commences sooner as compared to the nitrogen atmosphere. Interestingly, T<sub>onset</sub> and T<sub>peak</sub> of wood and PP<sup>v</sup>W00 become close to each other in the presence of oxygen. Narrow temperature interval between the occurrence of oxidative degradation of wood fiber and PP is attributed to the partial co-occurance of their degradation in a wide temperature range. At the heating rate of 5 °C/min, the degradation profile of WPC did not fall between PP and wood fiber curves, but by increasing the heating rate, the curves gradually move toward the area between the two components. At the low heating rate, the volatile compounds evolved in oxidative degradation

of wood fiber reduce access of oxygen species to the polymer chains. With the lack of access to oxygen, PP requires a higher temperature to decompose than normal oxidative degradation.



Figure 3. Thermal/oxidative degradation of wood fiber and PP-based WPCs

To analysis the effect of wood on the degradation of PP precisely, the DTG curves were deconvoluted to separate the overlapped cellulose and PP peak. The values of the deconvoluted figures were employed to determine  $E_a$  proposed by iso-conversional models. For instance, the plots of the FWO method, which was employed for the calculation of the  $E_a$  values for PP deconvoluted from WPCs are illustrated in Figure 4. Under an inert environment, the shape of the  $E_a$  evolution curves for PP degradation deconvoluted from WPCs was similar to PP sample without wood (PP<sup>v</sup>W00). Under the air, however, the  $E_a$  evolution curve of PP deconvoluted from WPCs showed an irregular decreasing trend with the conversion.



Figure 4. Variation of Ea for individual PP degradation obtained from deconvolution WPCs (OFW model)

#### 4. Conclusions

The degradation behavior of a simulated recycled PP and wood-PP biocomposites have been analyzed by TGA. Three iso-conversional methods, KAS, OFW, and Friedman, and two conversion-independent methods, Kissinger and Augis were employed to calculate  $E_a$ . T<sub>onset</sub> and T<sub>peak</sub> reduction. The results revealed that the PP underwent substantial degradation after reprocessing, confirmed by a remarkable drop of  $\tau_{\infty}$  accompanied by an increase in MFR. A progressive reduction was observed for  $E_a$  with the successive reprocessing cycles at a whole range of conversion.

In the WPC systems, two probable mechanisms, heat sink effect of the residual ash and thermal insulating effect of the foam-like structure of PP were proposed to cause a slight increase in  $T_{peak}$  of PP during thermal degradation. An interesting observation in the oxidation of WPCs was that the TGA profile did not fall between PP and wood curves, at low heating rates. The reduction in oxygen access by the polymer chains due to the evolved volatile compounds from wood was the most probable reason for this. Overlapping tails of degradation peaks corresponding to cellulose and PP was observed for degradation of WPCs. The Lorentzian area deconvolution method employed to separate them indicated that under N<sub>2</sub>, the E<sub>a</sub> evolution curves for PP shifted upward at the whole range of conversion due to thermal absorbency of the residual chars. Under the air, a significant increase in E<sub>a</sub> values was observed at the initial steps of degradation, which was associated with the high volume of degradation volatiles generated from wood that reduced the oxygen diffusion. At high conversions, the effect of wood became negligible as there was no residual remained from the decomposition of wood.

#### Acknowledgments

We gratefully acknowledge financial support from Natural Resources Canada (NRCan) for this research work. We are grateful to Ontario Sawdust Supplies for the supply of wood fiber.

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Sainiwetha Saihrishnan Chemical Engineering Waterloo

Thermo-mechanical Degradation of Polypropylene-Low Density Polyethylene blend system after multiple extrusion



1525

and ment with work

Javan Buratynski Chemistry Waterloo

Using Degradable Polymers to Separate Carbon Nanotubes



5

**》** 

Abdullah Ba Salem Chemistry Waterloo

Determination of Aggregation Number for Pyrene-labeled Gemini Surfactants



## Kristijan Lulic Chemistry Waterloo

## Self-Association of Oligoquinoline Foldamers Probed by Fluorescence Anisotropy



#### Acknowledgements

- Dr.Jean Duhamel
- Lab Members of the Duhamel Group
- Dr. Huc's Group for foldamer samples

• Jingqi Wang





## Hunter Little Chemistry Waterloo

## Progress in Instrumentation for Time-resolved Fluorescence Anisotropy



Same øvalue

Same r, value

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Relating Pyrene Excimer Fluorescence to Conformation in Pyrene-Labeled Polyamines



#### Conclusions

#### Since the rate constant ds for pyrene excimer formation of the pyrene-labeled polyamines is directly proportional to the local concentration of pyrene, the linear relationship between ds and $\|Py\|_{loc}$ suggest that the expression for $\|Py\|_{loc}$ which was derived in an earlier publication [1], is valid.

#### Acknowledgements

Afton Chemical Duhamel Group Members

Thus, <k> can be used to assess the local amine concentration experienced in a polyamine by covalently labeling the polyamine with the

Afton

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Tiana Trumpur Physics and Astronomy Waterloo

## Solvent Induced Nanoscopic Roughness on Glassy Polymer Thin Films



## Minghui Liu Chemical Engineering Waterloo

## Antimicrobial Activity of Quaternary Ammonium Compound/Polyurethane [QAC/PU] Colloidal Complex Film Based on Synergetic Release Killing and Contact Killing Mechanisms







## Remi Casier Chemistry Waterloo

## Copolymerization and its Effects on Polymer Dynamics in Solution

#### Copolymerization and its Effects on Polymer Dynamics in Solution

Remi Casier and Jean Duhamel

#### INTRODUCTION

The bulk properties of polymers are typically described by their glass transition temperature  $(T_g)$ . The effect that copolymerization has on  $T_g$  has been extensively studied leading to several commonly used equations, such as the Fox<sup>1</sup> and Gordon-Taylor-Wood<sup>2,3</sup> equations, which are able to predict the  $T_g$  of copolymers based on their comonomer constituents.<sup>4</sup> Despite this, little has been done to study how copolymerization affects the dynamics of polymers in solution in a controlled manner. The solution dynamics of polymers are not only important to the ability of macromolecules to spontaneously form ordered structures in solution<sup>5</sup> but also plays an important role in the rheological properties of polymer solutions.<sup>6</sup> The goal of this study is to help answer two questions: How does the structure (size) and content of each comonomer affect the dynamics of the resulting copolymer in solution. To this end, a series of copolypeptides were prepared with amino acids of varying size. By studying how the internal dynamics of the copolypeptides could be viewed as model systems to probe the solution behaviour of copolymers in a controlled setting.



**Figure 1.** The chemical structures of pyrene-labeled A) poly(glycine-*co-D*,*L*-glutamic acid) (Py-PGlyGA), B) poly(*D*,*L*-alanine-*co-D*,*L*-glutamic acid) (Py-PAlaGA), C) poly(*D*,*L*-glutamic acid) (Py-PGA), and D) poly(carboxybenzyl-protected *D*,*L*-lysine-*co-D*,*L*-glutamic acid) (Py-PLys(Z)GA). All stereocenters in the chemical structures are racemic.
Copolypeptides were prepared via a ring-opening polymerization of the *D*,*L*-*N*-carboxyanhydride (NCA) derivatives of *t*-butyl-protected glutamic acid (GA) with glycine (Gly), alanine (Ala), or carboxybenzyl-protected lysine (Lys(Z)). The *D*,*L*-isomers of all the amino acids were used to prevent the polypeptides from forming secondary structures (ex.  $\alpha$ -helices) in solution, thus ensuring they all adopted a random-coil conformation. After deprotecting the GA's with trifluoroacetic acid, a molar fraction *x* of the structural units was labeled with 1-pyrenemethylamine. Each polypeptide was prepared with several pyrene contents. The structures of the pyrene-labeled polypeptides used in this study are given in Figure 1.

The copolypeptides were characterized by <sup>1</sup>H NMR and GPC to determine their structural unit composition and absolute molecular weights, respectively. Their pyrene contents were determined by UV-Vis absorption measurements. A summary of the parameters describing the pyrene-labeled polypeptides is given in Table 1. The pyrene-labeled polypeptides were characterized by steady-state and time-resolved fluorescence. The monomer and excimer time-resolved fluorescence decays were globally analyzed with the fluorescence *blob* model (FBM). Briefly, the FBM divides a macromolecule into a series of homogeneous subvolumes (*blobs*). The FBM analysis of the fluorescence decays yielded  $N_{blob}$ , the number of structural units inside a *blob*, and *k*<sub>blob</sub>, the rate constant at which those structural units encountered one another inside a *blob*. Although the *blob* volume was expected to be the same for all the polypeptides used in this study, the  $N_{blob}$  values could be different. Therefore the product  $k_{blob} \times N_{blob}$  provided a better description of the polypeptide dynamics, as this product represents the number of diffusive encounters between the structural units per unit time in a *blob*.

1 2			2 I I		0		
Polypeptide	Polypeptide Composition		# <b>Py</b>	x	Mn	מח	А
	mol% GA	mol% comonomer	labelings	(mol%)	(kg·mol <sup>−1</sup> )	Dr	D
PGlyGA	48	4	4	3.5-5.3	13	150	-
PAlaGA	42	4	4	6.6–15.2	23	240	1.05
	59	5	5	6.6–10.4	16	150	1.21
	76	5	5	3.3-5.0	10	90	1.10
PGA	100	5	5	6.0–12.3	100	784	1.07
PLys(Z)GA	43	5	5	4.5-13.2	37	180	1.04

**Table 1:** The composition, pyrene contents (x), number-average molecular weight  $(M_n)$ , degree of polymerization (DP), and dispersity (D) of the polypeptides shown in Figure 1.

#### **RESULTS AND DISCUSSION**

In order to probe the effects of copolymerization on the internal dynamics of polypeptides, a series of copolypeptides (Figure 1) were prepared such that each contained glutamic acid (GA) while the comonomer was varied. In this manner the effect that each of the comonomers had on the dynamics of a poly(glutamic acid) (PGA) base polymer could be studied in a systematic manner. To quantify the differences between each of the chosen comonomers, each comonomer was assigned a value based on the number of non-hydrogen atoms in their side chains. This so-called *side chain size* (*SCS*) equaled 0 for Gly, 1 for Ala, 5 for Gly, and 15 for Lys(Z).

*Comonomer Side Chain Size:* To determine what effect comonomer size has on the internal structure and dynamics of polypeptides, a set of the polypeptides in Figure 1 were prepared such that each of them contained a similar amount of comonomer (ca. 44 ( $\pm$ 3) mol%). This ensured that any observed differences between them must have originated from changes in the comonomer size

and was not due to changes in composition. For each polypeptide, the Nblob value (# structural units in a *blob*) remained relatively constant as a function of pyrene content, indicating that the presence of pyrene did not alter the polypeptides behaviour. The Nblob values of each polypeptide obtained for different pyrene contents were then averaged and plotted as a function of their comonomer SCS in Figure 2A. Figure 2A shows that  $N_{blob}$ decreased quickly with increasing SCS, reaching a plateau for SCS above 4 atoms. A similar trend of decreasing Nblob with increasing SCS had



**Figure 2.** Comparison of the average A)  $N_{blob}$  and B)  $k_{blob} \times N_{blob}$  values of ( $\diamondsuit$ ) Py-PGlyGA, ( $\bigtriangleup$ ) Py-PAlaGA (42 mol% GA), ( $\circlearrowright$ ) Py-PGA, and ( $\square$ ) Py-Lys(Z)GA in as a function of the number of non-hydrogen side chain atoms in DMSO. The solid line indicates the  $N_{blob}$ <sup>theo</sup> value of  $10 \pm 1$  for an extended polypeptide in a coiled conformation. The dashed lines were added to guide the eyes.

also been observed with  $poly(n-alkyl methacrylate)s^7$  and is thought to be a result of the increasing side chain sterics which led to a straightening of the backbone on the few nm length-scale of a *blob*. The plateau  $N_{blob} = 11 (\pm 1)$  value obtained for PGA and PLys(Z)GA indicated that the side chains of GA and Lys(Z) were sufficiently large to fully elongate the polypeptide backbone. Consequently, any side chain containing 5 or more atoms would be expected to affect the backbone in a similar manner. To support this conclusion, molecular mechanics optimizations were conducted on an elongated PGA backbone. By monitoring *in silico* how far apart two pyrenyl labels could be located from one another along an elongated PGA backbone and still form excimer, a theoretical  $N_{blob}$ <sup>theo</sup> value was determined to equal 10 (±1), matching the experimentally determined  $N_{blob}$  value.

The dynamics of the polypeptides were measured through the term  $k_{blob} \times N_{blob}$ , which is plotted as a function of SCS in Figure 2B. Similar to the trends observed with  $N_{blob}$ ,  $k_{blob} \times N_{blob}$  was largest for PGlyGA and quickly decreased to a plateau with increasing SCS. The rapid decrease in  $k_{blob} \times N_{blob}$  observed for small SCSs indicated that the addition of a fairly small side chain was sufficient to fully stiffen the polypeptide backbone resulting in the near halving of the rate of encounter between the structural units within blobs from 0.26 ns<sup>-1</sup> for PGlyGA to 0.15 ns<sup>-1</sup> for PAlaGA. In agreement with the trends of  $N_{blob}$ , the plateau of  $k_{blob} \times N_{blob}$  at larger SCSs meant that incorporation of any comonomer with SCSs of 5 atoms or more would result in a copolypeptide exhibiting the same slow dynamics.

*Comonomer Composition:* The above section demonstrated that the incorporation of a monomer with a small side chain (ex. Gly and Ala) into a polypeptide was able to increase both the conformational freedom and dynamics experienced by the structural units of a polypeptide. However, since all the polypeptide contained ca. 44 mol% of these small comonomers, the relationship between the amount of comonomer and the polypeptides behaviour in solution could not be determined. To address this issue, a series of PAlaGAs (Table 1) were prepared with an

alanine content ranging from 24 to 58 mol%. Nblob was found to remain constant with pyrene content, thus demonstrating that the results obtained with the FBM were independent of the pyrene content (x). The effect that the Ala-content had on the chain structure is shown in Figure 3A. Although it was expected that the incorporation of Ala would increase Nblob, the response of N<sub>blob</sub> with respect to the alanine content was rather surprising. Figure 3A shows that the incorporation of 24 mol% Ala into a PGA backbone increased Nblob from



**Figure 3**: Comparison of A)  $\langle N_{blob} \rangle$  and B)  $\langle k_{blob} \rangle N_{blob} \rangle$  as a function of ( $\circ$ , 0 *mol% Ala*) Py-PGA and Py-PAlaGAs (*mol% Ala* = ( $\Box$ ) 0.24, ( $\diamond$ ) 0.41, and ( $\Delta$ ) 0.58) in DMSO. The dashed lines represent the average values of the Py-PAlaGAs. The solid line indicates the  $N_{blob}$ <sup>theo</sup> value of 10 ± 1.

11 (±1) to 15 (±1) due to the reduction of side chain sterics. However, upon the further addition of Ala,  $N_{blob}$  remained constant taking an average value of 16 ± 1. The constant  $N_{blob}$  value of the PAlaGAs indicated that the presence of relatively low levels of Ala were sufficient to entirely disrupt the elongated structure of PGA and therefore the incorporation of more Ala had no effect on the polypeptides conformation. Figure 3B shows that like  $N_{blob}$ ,  $k_{blob} \times N_{blob}$  of the PAlaGAs was more or less independent of the alanine content taking a similar, but slightly larger value compared to that of PGA.

#### **CONCLUSIONS**

The FBM was used to characterize how comonomer size and composition affected the internal dynamics and conformational freedom of copolypeptides. It was found that comonomers with large side chains restricted the backbone freedom of the copolypeptides resulting in the elongation of the backbone and a slowing in the dynamics. A point was reached however where the comonomer was sufficiently large to stiffen the polypeptide on the length-scale of a *blob*, at which point the copolypeptide no longer responded to further increases of the monomer size. Next, the response of a copolypeptide to monomer composition was probed by varying the amount of comonomer incorporated into a polypeptide. Aside from a rapid increase in  $N_{blob}$  due to the inclusion of a relatively small (< 25 mol%) amount of flexible comonomer, the solution behaviour of the copolypeptides did not change as a function of comonomer content. This suggested that the flexibility of a copolypeptide is dictated by the smallest (most flexible) comonomer, even if that comonomer had a relatively low incorporation.

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### Functionalized Cellulose Aerogel Beads for Heavy Metal Ions Removal

Winner of the 2020 IPR Award for Academic Excellence in Polymer Science/Engineering

### Functionalized Cellulose Aerogel Beads for Heavy Metal Ions Removal

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#### Abstract

Water pollution is a serious issue in the world, which is highly associated with the cumulatively discharge of industry wastewater. Heavy metal ions, such as  $Cu^{2+}$ ,  $Pb^{2+}$ , and  $Cr^{6+}$  are among the major components in the industry wastewater, which can threaten human health due to their high toxicity, non-degradability and bio-accumulation. Absorbents, such as activated carbon and petroleum-based polymers are commonly used absorbents or flocculants. However, their low adsorption capacity, high cost, poor recyclability and non-degradability make them a less attractive choice. Therefore, novel absorbents that satisfy all the practical requirements should be developed for future applications.

Developing bio-based absorbents possessing high removal capacity towards heavy metals is one of the key strategies to manage water pollution worldwide. Cellulose is one of the potential alternatives as it is abundant on earth, renewable and bio-degradable. Functionalized cellulose nanofibrils (CNF) as absorbents have been widely reported for wastewater treatment applications due to their larger specific surface area, abundant surface hydroxyl groups and structural flexibility. However, most of the reported cellulose nanomaterial- based absorbents directly used their single fibril form, which are difficult to recover due to their submicron size. Besides, pristine cellulose nanomaterials have low removal capacity due to the low affinity of hydroxyl groups towards heavy metals.

In my PhD research, functional groups are decorated on CNF to improve their metal ions adsorption capacity followed by assembled into aerogels which can be readily separated from the bulk solution. Two cellulose aerogel beads systems are designed depending on the crosslinking

methods and the functional groups. In one study, we grafted a hyperbranched cationic polymer, PEI on CNF using (3-glycidyloxypropyl) trimethoxysilane (GPTMS) as a crosslinker. The obtained mixture was injected into liquid nitrogen for rapid freezing, followed by sublimation to obtain cellulose aerogel beads (diameter between 3-4 mm). The preparation process is shown in Schematic 1. The chemical structure and composition, morphology, mechanical property, Cu (II) ion adsorption capacity and mechanism as well as regeneration performance of the aerogel beads was used instead of cellulose aerogel monoliths for Cu (II) removal to improve the removal efficiency by reducing copper ion diffusion path length; (2) small beads with larger surface to volume ratio than monoliths that enhanced the adsorption rate; (3) a scalable and simple one pot chemical crosslinking reaction conducted at room temperature and in aqueous solution was proposed; (4) the mechanical performance of the aerogel beads were precisely quantified, and to our knowledge, this is the first reported study; (5) the beads possessed excellent adsorption, regeneration and mechanical property, which is very promising for packed column adsorption.



Schematic 1 Illustration of the preparation and chemical structure of the CGP beads

In another study, we developed an efficient and simple and one step procedure to prepare carboxyl decorated and chemically crosslinked CNF aerogel beads. Pristine CNF aerogel beads (average diameter of 3 mm) were prepared by extruding the CNF slurry solution into liquid nitrogen

followed by freeze drying, and maleic anhydride (MA) was used as the crosslinker. The CNF aerogel beads were immersed in maleic anhydride to prepare robust carboxylated CNF aerogel beads. The chemical structure, chemical composition, morphology, mechanical property, Cu (II) ion adsorption capacity and mechanism as well as regeneration performance of the aerogel beads were investigated. The general preparation process and the key concept of this work is illustrated in Schematic 2.



**Schematic 2** Illustration of the preparation process, chemical structure and Cu<sup>2+</sup> removal of the CNF-MA beads

Natun Dasgupta Chemistry Waterloo

### Thermoresponsive Starch Nanoparticles for Oil Recovery from Tar Sands

#### Thermoresponsive Starch for Oil Recovery from Tar Sands

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The bitumen deposits in Canada are considered to be the third largest in the world, at 175-300 billion barrels.<sup>1,2</sup> The Athabasca oil sands lying along the Athabasca River, deposited about 110 million years ago,<sup>3</sup> form the largest bitumen deposit in the world. The sand particles forming the core of the oil sands consist of over 90% quartz with some amounts of potash feldspar, chert and muscovite. Clay minerals also present mostly comprise kaolinite, illite and small amounts of montmorillonite, as shown in Figure 1. Montmorillonite are fine clays with a particle size of less than 44  $\mu$ m. The porosity of the sand grains is 35% by volume, which allows the bitumen and water contents to be around 18-20% by weight altogether. Oil sands that are rich in bitumen have about 18% bitumen by weight, but on average the bitumen content of Athabasca oil sands is 12%.<sup>3</sup> An oil content of 5-10% is considered intermediate, while sands with 2-5% oil are considered lean.



Figure 5. (Left) Cross-sectional view of oil sand particles; (right) bitumen contents in high- and low-grade Athabasca oil sands (the dark spots represent bitumen). Adapted from Ref. 3.

Since bitumen is a heavy oil consisting of large hydrocarbon molecules, the Clark hot water extraction (CHWE) process currently used requires heating water to high temperatures, which consumes a lot of energy with environmental consequences. The procedure also involves harmful chemicals such as sodium hydroxide. Because a limited amount of water can be reused and recycled, this leads to the formation of large tailing ponds of fine sand, oil residues, and several harmful chemicals used in the bitumen extraction process posing great hazard to the environment.

These issues can be tackled by using polymeric surfactants exhibiting a lower critical solution temperature (LCST). These polymers, when coupled with hydrophilic starch nanoparticles, become amphiphilic at temperatures above the LCST. This characteristic can be exploited to extract oil from tar sands without heating the water to high temperatures. Since these smart materials are tunable, characteristics such as the value of the LCST can be controlled. This makes it possible to carry out the extraction process at lower temperatures (45 °C), but also to reuse the water for multiple cycles of oil extraction. The method developed thus minimizes the use of tailing ponds, protecting life below water and on the land.

Starch is a natural, renewable and biodegradable material consisting of two key polymers: amylose and amylopectin. Amylose is a linear polysaccharide typically making up 15-35% of starch in plants.<sup>4</sup> It has a large number of D-anhydroglucose units attached by  $\alpha(1\rightarrow 4)$  glycosidic linkages. Unlike amylose, amylopectin has a highly branched structure including linear segments with  $\alpha(1\rightarrow 4)$  glycosidic linkages, but also about 5% of  $\alpha(1\rightarrow 6)$ -linkages as branching points. Starch can be isolated from rice, oats, peas, wheat, tapioca, potatoes, corn and other plants.<sup>4</sup> In year 2000, starch production was about 48 million tons per year. Starch is of interest not only as food, but also to the industry for high-value applications.<sup>5</sup> The use of starch also ensures that these materials are environmentally friendly and non-toxic.

Polysaccharides grafted with thermoresponsive polymers display unique properties, in addition to good biocompatibility and biodegradability. Starch molecules useful for the extraction of oil from tar sands were obtained by modification with thermoresponsive poly(di(ethylene glycol) methyl ether methacrylate) (PMEO<sub>2</sub>MA) segments through RAFT (reversible addition-fragmentation chain transfer) grafting. PMEOM<sub>2</sub>A exhibits an LCST, such that the thermoresponsive polymer-grafted starch is amphiphilic above the LCST and hydrophilic below the LCST. The synthetic methods used provide easy control over the characteristics of the grafted starch (number and length of grafted PMEO<sub>2</sub>MA segments), and therefore over their hydrophilic-lipophilic balance (HLB). The starch-*g*-PMEO<sub>2</sub>MA samples were characterized by <sup>1</sup>H NMR, UV-visible spectroscopy, TEM and DLS analysis, and the grafted PMEO<sub>2</sub>MA chains were cleaved from the starch substrates for analysis by gel permeation chromatography.

Two different strategies were developed to control the characteristics of the target starch-*g*-PMEO<sub>2</sub>MA molecules, by using starch modified with suitable functional groups as RAFT agent for the methacrylate monomer (Figure 2). The first approach was to vary the degree of substitution (DS) of the starch-based RAFT agent, while maintaining a constant starch-*g*-PMEO<sub>2</sub>MA composition. A RAFT agent with a high DS should indeed generate a larger number of shorter polymer chains due to its multiple reactive sites, while a RAFT agent with a low DS would generate longer polymer chains due to the presence of fewer initiating sites. The other approach was to vary the amount of monomer added to the starch-based RAFT agent for a set DS, since the length of the PMEO<sub>2</sub>MA chains should increase with the amount of monomer added to the

reaction. A third approach explored to control the solution properties of the materials was the addition of a block of poly(2-hydroxyethyl acrylate) (PHEA) to form a hydrophilic shell on the particles. It was expected that PHEA, being hydrophilic, may prevent macrophase separation of the thermoresponsive graft polymer above its LCST.



Figure 6. Control of starch-g-PMEO<sub>2</sub>MA characteristics through variations in the DS of the RAFT agent and the amount of monomer added.

Oil extraction experiments were done using a clamp attached in an oven maintained at 45°C with 1 g oil sand samples in 20 mL vials as shown in Figure 3. Overall, over 80% extraction efficiency was obtained using starch-g-PMEO<sub>2</sub>MA (15 wt%, DS 0.022) in the presence of 60 mg of toluene, 15 mL of water and 0.5 M NaCl. The extraction efficiency should be higher for lower clay samples. Most importantly, the aqueous phase can be reused in multiple cycles, thus decreasing water and thermoresponsive starch consumption in the extraction process.



Figure 7. (Left) Vial with 1 g of tar sand (10  $\pm$  1% bitumen content), 15 mg starch-g-PMEO<sub>2</sub>MA

(15 wt%, DS 0.022), 0.5 M NaCl, 60 mg toluene and 15 mL water attached to a tumbler in an

oven at 45 °C; (right) bitumen floating on the surface 24 hours after the extraction.

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# Polysaccharide-lonic Liquid corrosion inhibitors



WATERLOO

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Ewomazino Ojogbo Chemical Engineering Waterloo

Effects of fabrication method on the dispersion of CNCs in highly crosslinked rubber composites



### Sanjay Patel Chemistry Waterloo

Pyrene-labeled Starch Nanoparticles as Fluorescent Sensors for Explosive Detection



### Donghan Liu Chemistry Waterloo

# Synthesis and Characterization of Furan Based Non-ionic Surfactant (FBNIOS)



A series of nonionic surfactants were successfully prepared with a hydropholic octyl chain, a furan core, and hydrophilic oligo(ethylene oxide)s of different lengths. Their CMC was determined by a combination of pyrene fluorescence and surface tension. These nonionic surfactants show properties similar to those of typical surfactants (micelle formation, lower surface tension) but use a furan core obtained from renewable, biodegradable cellulose.

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Joanne Fernandez Chemistry Waterloo

## Grafting of Starch Using a Complex Initiation System of CAN-KPS

#### Grafting of Starch Using a Cerium–Persulfate Initiation System

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#### Introduction

Starch is a biocompatible, renewable and biodegradable polysaccharide. Its abundance and relatively low cost have led to substantial interest in its use as a base material for a wide range of applications.<sup>1</sup> It is composed of two main components, amylose and amylopectin. Amylose is a linear polymer with D-glucopyranose units joined by  $\alpha$ -1,4-glucosidic linkages, whereas amylopectin is highly branched with both  $\alpha$ -1,4- and  $\alpha$ -1,6-glucosidic linkages.<sup>2</sup> The content of amylose and amylopectin in natural starch varies depending on the plant species, but "waxy" starch has essentially no amylose.<sup>1</sup> The characteristics, rheological behaviour and mechanical properties of starch can differ considerably depending on its composition.



Figure 1. Structure of the starch components (a) amylose and (b) amylopectin.

In its native state, starch has some characteristics that are undesirable for many industrial applications such as poor long-term stability caused by water absorption, and poor mechanical properties.<sup>3</sup> Due to these issues, it cannot be used directly in most cases. Modification by various means, including polymer grafting, is required to achieve more desirable properties. Through modification, starch can acquire characteristics comparable to synthetic petroleum-based

polymers.<sup>4,5</sup> Poly(acrylic acid)-grafted starch, in particular, possesses remarkable water absorbency, and has been used as a platform for controlled drug delivery as well as in paper and textile finishing.<sup>6</sup>

Graft copolymers derived at least in part from renewable resources such as starch have found numerous applications in the industry. The synthesis and properties of various graft copolymers derived from cooked starch, in combination with acrylic acid or acrylonitrile, were explored in this investigation. Initiation was best achieved through a combination of cerium (IV) and potassium persulfate. Grafting of vinyl monomers on starch with cerium (IV) ions is generally assumed to involve a redox initiation process. This is an efficient initiation method, as (ideally) radical centers are formed only on the starch substrate, albeit it is only known to work with a few specific monomers.<sup>7</sup> It was thus determined in the current investigation that acrylic acid could not be grafted directly onto starch using cerium (IV) ions, but the addition of an aldehyde to the reaction enabled grafting with high efficiency. Furthermore, potassium persulfate was introduced to reoxidize the cerium (III) to cerium (IV) in the initiation cycle, thus reducing the amount of initiator required in the reaction.<sup>8</sup> This presentation will focus on the synthesis of poly(acrylic acid)-modified starch and polyacrylonitrile-modified starch via cerium-persulfate initiation and aldehyde-promoted grafting. Various aldehydes were investigated to determine their influence on the grafting efficiency. The modified starch was characterized by nuclear magnetic resonance (NMR) spectroscopy and dynamic light scattering (DLS). In addition, gel permeation chromatography (GPC) was used to determine the molecular weight of the poly(acrylic acid) side chains generated in the grafting reaction.

#### **Experimental**

Using an initiation system with cerium (IV) and potassium persulfate, graft copolymers of starch with vinyl monomers were synthesized as shown in Scheme 1.



Scheme 1. Synthesis of poly(acrylic acid-modified starch. PTSA: p-toluenesulfonic acid; CAN:

ceric ammonium nitrate; KPS: potassium persulfate; PAA: poly(acrylic acid).

The Ce<sup>4+</sup>/aldehyde redox initiator system (Scheme 2) was also explored to increase the overall grafting efficiency of the reaction.



Scheme 2. The Ce<sup>4+</sup>/aldehyde reaction proceeds through coordination of the carbonyl with loss of an  $\alpha$ -hydrogen.

#### **Results and Discussion**

The synthesized starch graft copolymers were characterized by various techniques. Through proton nuclear magnetic resonance spectroscopy (Figure 2), the molar substitution of the graft copolymers was determined after purification by dialysis. The peaks at 1.7 and 2.3 ppm are for the  $-CH_2$ - and -CH- protons of the poly(acrylic acid) side chains, respectively. The molar substitution achieved was calculated by comparing the integrated values of the side chain peaks with that of the anomeric proton (H<sub>a</sub>) on the glucopyranose unit at 5.4 ppm.



Figure 2. <sup>1</sup>H NMR spectra for (a) starch and (b) poly(acrylic acid)-g-starch after dialysis.

Through the addition of various aldehydes (Table 1), it was found that glyoxal and

butyraldehyde both yielded high monomer grafting efficiencies on starch. This correlates with the known relative reactivity of aldehydes RCHO as R = H < phenyl < alkyl.<sup>9</sup>

Aldehyde	% Monomer conversion (modified starch and homopolymer)	% Monomer grafted to starch	% Monomer converted to homopolymer	Wt% PAA in starch- <i>g</i> -PAA
Glyoxal	99 ± 1	82 ± 7	17 ± 7	52 ± 2
Butyraldehyde	97 ± 2	81 ± 9	17 ± 10	52 ± 4
Glutaraldehyde	100 ± 0	59 ± 4	41 ± 4	44 ± 2
Acetaldehyde	80 ± 6	39 ± 2	41 ± 8	34 ± 2

**Table 1.** Comparison of the grafting efficiency in the presence of various aldehydes.

To better understand the effect of glyoxal on the grafting reaction, its concentration was varied from 0 to 8 weight percent with respect to starch. Without glyoxal, poly(acrylic acid) homopolymer was formed but it was not bound to the starch. When glyoxal was introduced in the system, the poly(acrylic acid) chains became bound to the starch substrate.



**Figure 2.** Effect of varying the concentration of glyoxal on (a) the % monomer conversion (modified starch and homopolymer), and (b) the % monomer grafted to the starch.

#### Conclusions

Graft copolymers of starch with acrylonitrile and acrylic acid were successfully synthesized using a cerium/persulfate initiation system. Through the addition of aldehydes (e.g. glyoxal) to the reaction, the yield of side chains bound to the starch substrate increased markedly. A wider range

of molar substitution levels will be investigated, and further characterization of the graft copolymers by transmission electron microscopy will be carried out.

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Cellulose Nanomaterials: Synthesis, Properties, and Applications

### Cellulose Nanomaterials: Synthesis, Properties, and Applications

#### ABSTRACT

Nanotechnology has been recognized as one of the most promising and versatile areas for potential high-tech development in the 21<sup>st</sup> century. As research advances, the demand for non-petroleum based, carbon neutral biodegradable products from sustainable resources having low environmental, animal/human health and safety risks is ever rising in the field of nanotechnology<sup>1</sup>. The environmental safety issues and the emphasis on "green synthesis (solvent free)" from industries and government have recently generated a significant interest in the development of nanomaterials based on renewable and sustainable natural resources such as cellulose nanocrystals (CNCs) as the alternatives of other polymers or resins. Cellulose is one of the most ubiquitous and abundant renewable biopolymers produced on earth, which is considered as an almost inexhaustible source of raw material to meet the ever-rising demand for a wide spectrum of biocompatible products and materials. Soft and hardwood from higher plants is the most commercially exploited source of cellulose<sup>2</sup>.

Considering the increasing interest in sustainability and nanotechnology, the use of CNCs in water treatment applications has gained increasing attention. CNCs have been actively researched for their use in various water treatment systems, such as adsorption, absorption, flocculation, membrane filtration, catalytic degradation and disinfection<sup>3</sup>. CNCs are attractive candidates because of its high specific surface area, high specific strength, hydrophilicity, biodegradability and surface functionalization capabilities<sup>2</sup>. The major types of pollutants present in water bodies include organic dyes, heavy metal ions, pharmaceutics, pesticides, polycyclic aromatic hydrocarbons, and biomolecules. Adsorption is an attractive and economical technique, and when implemented using a well-designed system, it can offer excellent results<sup>3</sup>. The

production of conventional adsorbents, such as activated carbons, can be energy intensive, expensive, and emit greenhouse gases. Thus, producing alternative low-cost adsorbents from industrial and agricultural by-products offers many new opportunities. The use of sustainable nanomaterials, such as CNCs will reduce our dependence on activated carbons and also reduce the carbon foot print as several CNCs based adsorbents have demonstrated excellent adsorption capacity<sup>3</sup>.

Cellulose is a long-chain, high molecular weight, and hydrophobic polysaccharide (an isotactic homopolymer) composed of repeat units of dimers or disaccharides of anhydro-Dglucopyranose unit (AGU), known as cellobiose where each anhydro-D-glucopyranose unit is connected by  $\beta$ -1,4 glycosidic linkage and every other monomer is rotated 180<sup>0</sup> with respect to its neighbouring unit (Figure-1A). Cellulose microfibrils are the structural units in the higher plants which are formed by a biogenetic bottom-up process in the nature where the cellulose chains connect or assemble with each other and stabilize through hydrogen bonds and van der Waals forces, further assemblies of fibrils in turn lead to formation of cellulose fibers to form a hierarchical system. These microfibrils, which are several micrometers in length and 5-10 nm in diameter, contain highly ordered (crystalline) regions alternating with disordered (amorphous) regions (Figure-1B and 1C). These crystalline regions can be separated and extracted from the desired cellulose source through a "top down" deconstruction of cellulose fibers by mechanical, chemical or a combination of mechanical, chemical and/or enzymatic (endoglucanases for example) treatments (Figure-1B and 1C), yielding the cellulose nanocrystals (CNCs). The simplest and widely employed process to extract CNCs involves the acid hydrolysis, where concentrated sulphuric acid (H<sub>2</sub>SO<sub>4</sub>) and hydrochloric acid (HCl) are most commonly used. Sulphuric acid hydrolysis introduces abundant anionic sulfate half ester groups (-OSO<sub>3</sub><sup>-</sup>) onto the CNC surface

(Figure-1C), which provides electrostatic repulsion among CNC particles in aqueous dispersion and enables the stabilization of CNCs over a wide range of pH values<sup>2</sup>. Currently, the majority of the commercially supplied CNCs (commonly called as the 'Pristine CNC') possess surface – OSO<sub>3</sub><sup>-</sup> groups and thus are negatively charged.



*Figure-1: (A)* Chemical structure of cellulose biopolymer representing reducing and nonreducing ends. *B)* Schematic illustration of CNC production from fibre cell walls by mechanical and chemical treatments, respectively. *C)* Amorphous and crystalline portions in cellulose microfibrils, intrachain & interchain hydrogen bonding (dotted line), and the numbering of the carbons in each AGU showing the  $1^0$  alcoholic groups<sup>2</sup>. The CNC rods in twisted form at the bottom right corner was adapted from<sup>4</sup>.

In terms of renewability of CNCs, the presence of abundant surface hydroxyl groups enables them for different chemical modification such as oxidation, esterification, etherification, silylation, polymer grafting, etc. (reaction scheme shown in Figure-3). From Figure 1C, it is evident that each AGU of a cellulose chain contain three hydroxyl groups which can be categorized into two chemical class namely, primary (1<sup>0</sup>) and secondary (2<sup>0</sup>) alcoholic groups. In most instances the hydroxyl groups at the C2 and C3 positions are secondary alcohols while the hydroxyl group in the C6 position is a primary alcohol. Although, the relative reactivity of the hydroxyl groups has the following order; (C6)–OH >> (C2)–OH > (C3)–OH, but most of the modification occurs at the (C6)–OH<sup>5</sup>. Such potential chemical modifications also allow their incorporation and dispersion into different hydrophilic/hydrophobic polymer matrices effectively rendering CNCs the status of an ideal candidate as a novel material<sup>5</sup>.



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Figure-2: Surface functionalization of CNCs through various covalent modifications.

Currently, the three major suppliers of pristine CNCs are -

- ➤ CelluForce Inc: the world's largest producer of CNCs (NCC<sup>TM</sup>),
- ➢ Alberta Innovates Technology Futures (AITF), and
- > USDA (U.S. Department of Agriculture) Forest Products Laboratory (FPL).

As the majority of the surface  $1^0$  alcoholic groups are occupied by  $-OSO_3^-$  groups in the pristine CNCs (reported S = 0.86 %, w/w)<sup>6</sup>, and the chemical modification(s) preferably occurs at the 'free' 'most reactive'  $1^0$  alcoholic group(s) on C6 protruding from the CNC rods, desulfation of such pristine CNC is necessary prior to any chemical modification.

CNCs are the unique nanomaterials among the other nanomaterials of the same range due to its exceptional properties discussed above. Due to its nanoscale size, high aspect ratio, and unique mechanical and chemical properties, CNCs have a plethora of applications notably in the paper, polymer, plastics, chiral templating, flocculants, aerogels, hydrogels, materials science, electronics, drug delivery, cosmetics, pharmaceuticals and tissue engineering.

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Better Understanding the Structure of Glycogen with Pyrene Excimer Fluorescence

#### Characterization of the Conformation of Glycogen in Solution by Pyrene Excimer Fluorescence Damin Kim and Jean Duhamel

IPR Symposium, University of Waterloo, Waterloo, ON, N2L 3G1, Canada

#### Introduction

Glycogen is a major carbohydrate that animals and humans alike use as energy storage. The structure of glycogen has been investigated for many years. Glycogen is a polysaccharide made of linear segments of anhydroglucose units (AGUs) joined by  $\alpha$ -(1,4) glycosidic bonds. These oligosaccharides are connected to each other by  $\alpha$ -(1,6) glycosidic bonds. The general structure of glycogen is similar to that of amylopectin, the main carbohydrate used for energy storage by plants. However, the degree of branching of glycogen is twice larger than for amylopectin. Following the suggestion by Whelan in 1970 that glycogen could be viewed as a sphere made of concentric tiers,<sup>1</sup> the Tier Model was proposed. The fundamental purpose of the Tier Model was to explain how glycogen could store about 55,000 glucose units inside a sphere with a diameter of 40 nm.<sup>2</sup> The Tier Model uses the degree of polymerization, or chain length (*CL*), of the side chains constituting glycogen. *CL* can vary from 9 to 16 AGUs depending on the source of glycogen.<sup>3</sup> The molecular weight of glycogen can be calculated with Equation 1 where 162 g/mol, *r*, and *t* represent the molar mass of an AGU, the degree of branching, and the number of tiers. Degrees of branching higher than 2 result in densities that are much too high for glycogen. After optimization, the degree of branching higher than 2 result in densities that are much too high for glycogen. After optimization, the degree of branching was found to equal 2 (i.e. r = 2 in Equation 1).<sup>4</sup>

$$M_n = 162 \times \left(\frac{1 - r^t}{1 - r} \times CL\right) \tag{1}$$

Considering that each AGU increases the tier thickness (*tt*) by 0.126 nm and that the branching point generated by an  $\alpha$ -(1,6) glycosidic bond adds 0.35 nm to *tt*,<sup>5</sup> *tt* could be calculated according to Equation 2.

$$tt = 0.126 \times CL + 0.35 \tag{2}$$

The density ( $\rho$ ) of a glycogen molecule made of *t* tiers can then be calculated with Equation 3.

 $\rho = \frac{3 \times 162 \times CL}{4\pi (t \times tt)^3 \times N_A} \quad \text{it is wrong wrong}$ 

(3)

*X*-ray crystallography has demonstrated that the crystalline lamellae of amylopectin are constituted of mainly double helices that are hexagonally packed.<sup>6</sup> Assuming that the oligosaccharide branches of glycogen form helices that also pack hexagonally, the density of each tier can be calculated with Equation 4 where  $d_{h-h}$  is the interhelical distance.

$$\rho_{tier} = \frac{162 \times CL}{N_A \left(\frac{\tan(\pi/3)}{2} d_{h-h}^2 \times tt\right)} = \frac{162 \times CL \times r^{t-1}}{\frac{4\pi}{3} \times tt^3 \left(t^3 - (t-1)^3\right)}$$
(4)

A hypothetical glycogen molecule made of 13 tiers would be too dense and would leave no room in the peripheral tier for an enzyme to enter the glycogen macromolecular volume and graft a new oligosaccharide branch or depolymerize the polysaccharide into AGUs. Therefore, the maximum number of tiers that glycogen can have is 12.<sup>3</sup>

Two glycogen samples, one extracted from oyster and the other from corn (phytoglycogen), were probed by a combination of pyrene excimer fluorescence (PEF), fluorescence blob model (FBM), and molecular mechanics optimizations (MMOs) to assess the validity of the Tier Model proposed for glycogen. Gel Permeation Chromatography and intrinsic viscosity measurements were also conducted to determine the molecular weight and density of glycogen, respectively.

#### **Experimental**

*Materials*: Glycogen from oyster (Type II) was purchased from Sigma Aldrich and phytoglycogen was generously donated by Mirexus. Glycogen from oyster and corn were used after precipitation in ethanol. All other chemicals involved in the pyrene-labeling reaction were purchased from Sigma and used without any purification.

*Labeling of Glycogen with 1-Pyrenebutyric Acid*: Esterification, purification, and characterization of pyrene-labeled glycogen was conducted in the same manner as described earlier for pyrene labeled nanosized amylopectin fragment.<sup>7</sup> The only modification added to the procedure was the temperature at which glycogen was dispersed in DMSO. Glycogen was dispersed in DMSO at 80 °C compared to 60 °C for nanosized amylopectin fragments.

#### **Results and Discussion**

The molecular weight distribution of unlabeled glycogen from oyster and corn was determined by gel permeation chromatography in DMSO. The number average molecular weight (M<sub>n</sub>) of glycogen from oyster and corn equaled  $4.1 \times 10^6$  and  $8.5 \times 10^6$  g/mol, respectively. Both glycogens exhibited a narrow molecular weight distribution with a PDI of 1.02 for glycogen from oyster and 1.15 for phytoglycogen. The intrinsic viscosity ([ $\eta$ ]) of glycogen from oyster was determined with an Ubbelohde viscometer to equal 9.1 ± 0.5 mL/g. The pressure detector on the GPC was used to determine [ $\eta$ ] for phytoglycogen found to equal 7.8 ± 0.5 mL/g. Using the known *CL* of 10 and 12 for, respectively, glycogen from oyster<sup>2</sup> and corn,<sup>8</sup> the number of tiers, that each glycogen molecule was made of, could be calculated according to Equation 3. The results of these calculations are listed in Table 1.

	Experimentally determined # Tier				
	Intrinsic Viscosity	GPC	Fluorescence	Average	
Glycogen from oyster	11.3	11.2	12.2	11.6 (± 0.8) ~12	
Phytoglycogen	12.2	12.6	12.6	12.2 (± 0.3) ~12	

Table 1. Number of tier in glycogen determined by intrinsic viscosity, GPC and fluorescence.

The glycogen samples were further investigated by GPC. The average hydrodynamic radius ( $R_h$ ) of glycogen equaled 18.1 (±0.3) nm for oyster and 23.5 (± 1.9) nm for phytoglycogen. The difference in size could be attributed to the different CL values which have been reported to equal 10 and 12 AGUs for glycogen from oyster<sup>3</sup> and corn,<sup>8</sup> respectively. According to Equation 2, these CL values imply a tt of 1.61 and 1.86 nm for glycogen from oyster and corn, respectively. In turn, they would suggest a number (t) of tiers given by the ratio  $R_h/tt$  equal to 11.2 (±0.2) and 12.6 (±1.0) for glycogen from, respectively, oyster and corn, in excellent agreement with the t values obtained by [ $\eta$ ] measurements (see 1<sup>st</sup> column in Table 1). The shape factor given by the ratio of the radius
of gyration over  $R_h$  ( $R_g/R_h$ ) equaled 0.86 and 0.94 for glycogen from oyster and corn, respectively. Hyperbranched polymers generally have a shape factor between 0.98 and 1.2.<sup>9</sup> Glycogen from oyster had a somewhat lower value than expected for hyperbranched polymers, but it agreed with other  $R_g/R_h$  reported in the literature.<sup>10</sup> The Mark Houwink Sakurada (MHK) exponent was found to equal 0.34 for phytoglycogen. This exponent was similar to that of amylopectin reflecting the highly branched nature of glycogen.<sup>11,12</sup> The MHS exponent was also smaller than 0.5 which is what would be expected for a polymer in a  $\theta$ -solvent.<sup>13</sup> The small exponent of 0.34 confirmed the compact nature of phytoglycogen due to its highly branched architecture.

The validity of the Tier Model was further investigated for glycogen by PEF. The two glycogen samples were labeled with different amounts of pyrene. The fluorescence spectra were acquired for the pyrene-labeled glycogen samples and they are shown in Figure 1A for glycogen from oyster. As the pyrene content increased, more pyrene-pyrene encounters took place in solution and more excimer was formed resulting in an increase in the fluorescence intensity of the excimer at 480 nm with respect to that of the monomer at 378 nm. This effect could be quantified by plotting the ratio of the fluorescence intensity of the pyrene excimer ( $I_E$ ) over that of the monomer ( $I_M$ ) as shown in Figure 1B. The  $I_E/I_M$  ratio increased continuously with increasing pyrene content in Figure 1B reflecting the increase in local pyrene concentration experienced by an excited pyrene. Interestingly, the  $I_E/I_M$  ratios for glycogen obtained from oyster or corn overlapped at a same pyrene content. This observation indicated that at the molecular level, an excited pyrenyl label experienced a same environment for both glycogen samples suggesting that both samples had a similar interior.



**Figure 1.** (A) Emission spectra of glycogen from oyster labeled with different pyrene contents (*x* increasing from 0.02 to 7.8) (B)  $I_E/I_M$  ratio as a function of pyrene content for pyrene-labeled glycogen from oyster ( $\bullet$ ) and from corn ( $\circ$ ).

While interesting, the  $I_{\rm E}/I_{\rm M}$  ratios only allow a somewhat limited analysis of the fluorescence response for any pyrene-labeled macromolecule because it combines two effects. Indeed, PEF depends on first, the conformation of the macromolecule in solution, a denser conformation bringing the pyrenyl labels closer from each other and resulting in stronger PEF, and second, the flexibility of the polymeric backbone, a stiffer backbone hindering pyrene-pyrene encounters and resulting in less efficient PEF. This is when the fluorescence blob model (FBM) analysis of fluorescence decays acquired with a macromolecule randomly labeled with pyrene comes into play. The FBM assumes that the macromolecular volume can be compartmentalized into *blobs* among which the pyrene labels distribute themselves randomly according to a Poisson distribution. FBM analysis of the fluorescence decays yield the parameters  $N_{\rm blob}$  and  $k_{\rm blob}$ .  $k_{\rm blob}$  is the rate

constant of PEF and reflects the flexibility of the macromolecular backbone while  $N_{blob}$  is the number of structural units of the macromolecule found inside a *blob*.  $N_{blob}$  is thus a measure of the internal density of the macromolecule.

An earlier study applied PEF to determine that amylose adopts a helical conformation in DMSO.<sup>14</sup> To reach this conclusion, the  $N_{blob}$  value of  $10 \pm 1$  found for pyrene-labeled amylose in DMSO was compared to that of a molecular model built with HyperChem for pyrenyl labels covalently attached onto an amylose helix. The theoretical  $N_{blob}$  ( $N_{blob}$ <sup>theo</sup>) found by conducting molecular mechanics optimization (MMOs) to probe the maximal number of AGUs separating two pyrenyl labels and still allowing PEF was found to equal 11, in perfect agreement with the experimental  $N_{blob}$  of  $10 \pm 1$ . The good agreement between the experimental and theoretical  $N_{blob}$  values indicated that amylose adopted a helical conformation in DMSO. These PEF experiments were repeated with the pyrene-labeled glycogens and the  $N_{blob}$  values retrieved by applying the FBM analysis to the fluorescence decays were plotted as a function of pyrene content in Figure 2 A. Within experimental error,  $N_{blob}$  remained constant with pyrene content and regardless of the origin of the glycogen sample. It equaled 41 (±3), a much larger value than for amylose with an  $N_{blob}$  value of  $10 \pm 1$ . The four-fold larger  $N_{blob}$  value obtained for glycogen compared to amylose indicated that the environment probed by an excited pyrene was much denser for glycogen than for amylose.



**Figure 2.** (A) Plot of  $N_{blob}$  as a function of pyrene content for pyrene-labeled glycogen from oyster (•) and corn ( $\circ$ ) (B) Illustration of the hexagonal arrangement for the glycogen helices. (C) Plot of  $N_{blob}^{theo}$  (•) as a function of  $d_{h-h}$  and the experimentally determined  $N_{blob}(\bullet)$ .

As for amylose, MMOs were conducted with Hyperchem to determine  $N_{blob}^{theo}$  for glycogen. An oligosaccharide helix was built with 10 AGUs. Since the primary alcohol is more reactive than the secondary alcohol, 1-pyrenebutyric acid was attached to the C6 hydroxyl of the AGU. The reference pyrene was attached to the AGU located at the top of the helix and the secondary pyrene was attached onto the next AGU and the two pyrenes were induced to overlap. A pair of pyrenes, where one of the pyrenes had more than 6 carbons overlapping the frame of the other pyrene, was considered to result in PEF. The largest number of AGUs separating two pyrene labels while still allowing PEF was taken as  $N_{blob}^{theo}$ .  $N_{blob}^{theo}$  for PEF taking place inside a same helix equaled 6.5  $\pm$  0.5 AGU. Assuming that the glycogen helices were arranged in a hexagonal array,  $N_{blob}^{theo}$  was determined as a function of the interhelix distance ( $d_{h-h}$ ) when PEF occurred between two pyrenyl labels attached on two different helices. The reference pyrene was attached on the helix located at the center of the hexagonal array shown in Figure 2B and the second pyrene was attached on one of the adjacent helices. When  $d_{h-h}$  was greater than 34 Å, PEF could not take place between helices and  $N_{blob}^{theo}$  equaled 7. Upon decreasing  $d_{h-h}$ , more anhydroglucose units were included in the calculation of  $N_{blob}^{theo}$  and  $N_{blob}^{theo}$  was shown to increase quickly with decreasing  $d_{h-h}$  in Figure 2C. When  $d_{h-h}$  equaled 16.2 Å, the experimental  $N_{blob}$  matched  $N_{blob}^{theo}$ .

Equating each side of Equation 4 for the density of each tier ( $\rho_{tier}$ ) using a  $d_{h-h}$  value of 16.2 Å yielded *t* which was found to equal 12.2 and 12.6 for glycogen from oyster and corn, respectively. This derivation assumed that pyrene targets preferentially the parts of glycogen that are denser in AGUs located mainly at the periphery of the molecule. The *t* values obtained by a combination of PEF, FBM, and MMOs were in perfect agreement with those found by [ $\eta$ ] and GPC measurements.

## Conclusions

Glycogen was obtained from two different sources and the number of tiers constituting both glycogen samples were investigated with three different techniques including GPC,  $[\eta]$ , and PEF. The three techniques confirmed that these glycogen samples contained 12 tiers, regardless of the source of glycogen. This study represents the first example where PEF was applied to demonstrate the applicability of the Tier Model for glycogen. The PEF study yielded the interhelical distance in the 12<sup>th</sup> outer tier of glycogen equal to 16.2 Å.

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