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Computational-Based Design of Hydrogels with Predictable Mesh Properties

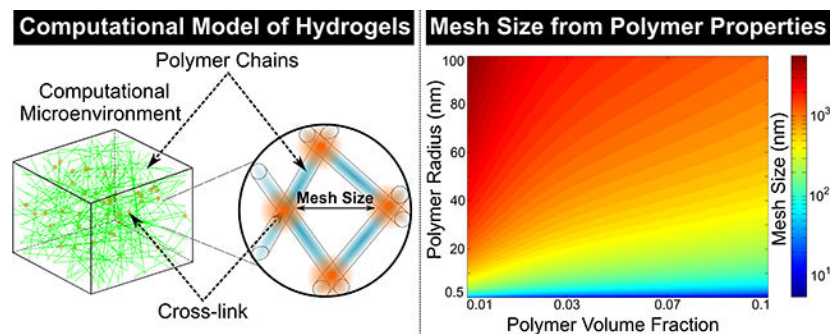
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Abstract

Hydrogel systems are an appealing class of therapeutic delivery vehicles, though it can be challenging to design hydrogels that maintain desired spatiotemporal presentation of therapeutic cargo. In this work, we propose a different approach in which computational tools are developed that creates a theoretical representation of the hydrogel polymer network to design hydrogels with predefined mesh properties critical for controlling therapeutic delivery. We postulated and confirmed that the computational model could incorporate properties of alginate polymers, including polymer content, monomer composition and polymer chain radius, to accurately predict cross-link density and mesh size for a wide range of alginate hydrogels. Additionally, the simulations provided a robust strategy to determine the mesh size distribution and identified properties to control the mesh size of alginate hydrogels. Furthermore, the model was validated for additional hydrogel systems and provided a high degree of correlation ($R^2 > 0.95$) to the mesh sizes determined for both fibrin and polyethylene glycol (PEG) hydrogels. Finally, a full factorial and Box-Behnken design of experiments (DOE) approach utilized in combination with the computational model predicted that the mesh size of hydrogels could be varied from approximately 5 nm to 5 μm through controlling properties of the polymer network. Overall, this computational model of the hydrogel polymer network provides a rapid and accessible strategy to predict hydrogel mesh properties and ultimately design hydrogel systems with desired mesh properties for potential therapeutic applications.

Graphical Abstract



Keywords

Hydrogels; Mesh Size; Computational Modeling; Cross-link Density; Mesh Size Distribution; Alginate; Fibrin; Polyethylene Glycol

Introduction

Hydrogel systems have many promising clinical applications due to their ability to provide control over the presentation of therapeutic agents to the surrounding tissue environment. Hydrogels consist of cross-linked polymer chains that do not dissolve but can swell in water or aqueous solutions and can provide a wide range of controllable mechanical properties to exert their desired therapeutic mechanism¹⁻³. One promising strategy to control therapeutic delivery involves designing hydrogels with predefined mesh sizes^{1, 3-5}. The mesh size of a hydrogel could be defined as the distance between two adjacent cross-links and has a critical role in determining the diffusion of agents within the hydrogel network^{1, 6-8}. Indeed, the mesh size is involved in determining the ability of therapeutic cargo to diffuse through the polymer system⁹⁻¹⁴ and the ability of encapsulated cells to interact with the surrounding environment for cell-based therapies¹⁵⁻¹⁷. The role mesh size has in providing controllable therapeutic delivery from hydrogel systems has prompted interest in designing hydrogels with desired mesh sizes.

Characterizing the mesh structure of a hydrogel system is one of the most critical parameters when designing hydrogels for drug delivery applications. Several approaches have been developed to characterize the mesh structure of hydrogel systems. Typical methods to determine the mesh size of hydrogels rely on swelling^{5, 18}, rheology^{15, 19-24}, small angle X-ray scattering²⁵, cryoporometry²⁶, dynamic light scattering²⁷, low field nuclear magnetic resonance (NMR)²⁶ and diffusion-based methods^{15, 28}. However, many of these techniques require access to complex and expensive experimental setups and provide limited information on the distribution of mesh sizes within the hydrogel. Furthermore, applying these methods to develop hydrogel systems with the desired mesh structure can be challenging. An alternative strategy involves developing a computational model of the hydrogel that creates a theoretical representation of the actual cross-linked polymer structure to determine mesh properties. A computational strategy that incorporates specific properties of the polymer system could be advantageous by providing a rapid and accessible method to predict the hydrogel mesh structure and ultimately to help the design of hydrogels with controllable mesh properties.

Here, we propose developing a computational strategy that can mimic the polymer network of the hydrogel system to determine mesh properties. Alginate, a naturally occurring anionic polymer comprised of α -L-guluronic (G-block) and β -D-mannuronic (M-block) residues, was utilized as our hydrogel model system when developing and validating the computational model given it has extensive use for various therapeutic delivery applications^{15, 29-35}. The computational model incorporated alginate polymer properties including polymer chain radius and monomer composition to determine hydrogel mesh properties. The model was then extended to predict the mesh properties of other popular

hydrogel systems. Thus, applications of this computational model could provide an appealing strategy for predicting the mesh size distribution of various polymer systems to help the design of hydrogels for a variety of potential therapeutic applications.

We hypothesize that incorporating properties of the polymer system into our computational model will allow for accurate prediction of hydrogel mesh size and allow for the identification of properties most relevant to the mesh structure. In this work, we begin by designing the computational model of the hydrogel mesh structure and determining how the ratio of M/G-blocks and polymer content effects cross-link density and mesh size distribution. The mesh size of alginate hydrogels composed of polymers with various molecular weight, G-block content and polymer volume fractions were determined to validate the simulations. Then the design of experiments (DOE) was used to determine how the mesh size predicted from the model changes for hydrogels composed of both high G-block content (MVG) and low G-block content (MVM) polymers. Finally, the model was further validated for fibrin and PEG hydrogels and then extended to predict the mesh size of other hydrogels from the properties of the polymer system.

Materials and Methods

Computational Model of the Hydrogel Mesh structure

The hydrogel mesh structure was modeled as randomly oriented overlapping cylinders within a unit cell as previously described with modifications³⁶⁻³⁷. Briefly, polymers were approximated as unidirectional randomly oriented cylinders where one point on a face of the unit cell was randomly selected as the start of the cylinder and another randomly selected point on a different face of the unit cell was assigned as the end of the cylinder. The radii of the cylinders were set at 0.8 nm based on the estimated average radius of alginate polymers³⁸⁻³⁹ and the volume contribution of each simulated polymer was approximated as the volume of the cylinder. Simulated polymers were added to the unit cell until reaching a desired volume fraction. For alginate hydrogels, the cylinders were divided into segments of either M-blocks or G-blocks with varying lengths based on the desired M/G ratio and the Kuhn length of an alginate monomer (0.515 nm)⁴⁰⁻⁴¹. Cross-links within the modeled alginate hydrogels were then determined from regions where two simulated polymers had four overlapping G-blocks (two from each polymer) based on the egg-box model for calcium alginate hydrogels⁴²⁻⁴⁵. Simulated hydrogels were assumed to have sufficient calcium to cross-link any four overlapping G-blocks and the number of cross-links within the unit cell was used to determine the cross-link density and the average distance between neighboring cross-links was calculated to determine mesh size. The cross-link density, mesh size and mesh size distribution were modeled for hydrogels over a wide range of polymer volume fractions and with alginate polymers containing varying G-block content. The normalized mesh distribution from the simulations was used to determine the probability of finding a mesh of size ξ (Equation 1):

$$P(\xi) = \frac{S(\xi)}{\int_0^{\infty} S(\xi) d\xi} * 100 \quad (1)$$

Where $S(\xi)$ is the generated best fit curve through the simulation data determined via nonlinear regression. For modeling the fibrin and PEG hydrogels, cylinders were set with a radius of 75 nm based on the estimated average radius of fibrin fibers⁴⁶ or 0.51 nm based on the estimated radius of PEG⁴⁷. For all simulations, the volume of the unit cell was large enough to ensure that mesh properties were calculated from at least 10,000 cross-links. All simulations were performed in Java using Eclipse and cross-link density, mesh size and mesh size distribution were calculated from multiple simulations ($n = 10$) and the average values were reported.

Hydrogel Formulation

The alginate polymers used in this study were obtained from Novamatrix (FMC). The MVG alginate containing a higher G-block content (~70% G-block alginate as specified by the manufacturer) used in this study included a lower molecular weight (LMW) LF 10/60 and a higher molecular weight (HMW) LF 20/40. The MVM alginate used in this study contained a lower G-block content (~35% G-block alginate as specified by the manufacturer). The alginate polymers used in this study were treated with activated charcoal to further purify the material as previously described^{15, 34}. Briefly, 1 g of alginate was dissolved in 100 ml of deionized (DI) water before being mixed with 0.5 g of activated charcoal (Sigma) for 30 minutes. The alginate solution was then sterile filtered (0.22 μm , Thermo Fischer Scientific) and lyophilized prior to starting hydrogel experiments. Hydrogels for the swelling, polymer volume fraction and storage modulus studies were prepared by dissolving alginate polymer powder in DI water overnight to minimize the contribution of salt when calculating polymer volume fraction. Hydrogels were prepared to create a final concentration of 1, 1.5, 2, 2.5 or 3% (w/v) polymer alginate solution and 0.42 mg of calcium sulfate (Sigma) per mg of alginate polymer. The alginate solution was then dispensed between two glass plates with 2-mm spacers height and allowed to gel for at least 25 minutes at room temperature before being cut with an 8-mm biopsy punch as previously described^{15, 23–24, 29}. Hydrogels were then transferred to 24-well plates and swelled in 0.5 ml of DI water at 37°C for an additional hour prior to determining hydrogel properties. For the diffusion of FITC dextran studies, alginate polymer was dissolved in DI water supplemented with 10 mM HEPES (Sigma) to control the effect of pH on the fluorescence of FITC. Hydrogels were created with 2 or 3% (w/v) polymer content and were dispersed in 12-well plates to gel. Hydrogels then had a square region (~1 cm^2) removed from the center of the hydrogel and were swelled in an additional 1 ml of 10 mM HEPES at 37°C for an hour. For the fibrin hydrogel studies, fibrinogen was dissolved in 0.9% NaCl (Sigma) and mixed with thrombin in 8-well plates (Lab-Tek) to create a final concentration of 2.5, 5, 7.5 or 10 mg/ml of fibrinogen and 1 unit of thrombin per mg of fibrin. Fibrin hydrogels were then incubated overnight at 37°C to ensure full gelation. For PEG hydrogels, PEG-diacrylate (PEGDA, 20K MW, Laysan Bio) was dissolved in HEPES-buffered saline (HBS, pH 7.4) to create a final concentration of 10, 15, 20 and 40% (w/v) containing 30 μg of 2,2-dimethoxy-2-phenyl-acetophenone (DMAP) in *N*-vinylpyrrolidone (NVP) per mg of PEGDA as previously described⁴⁸. PEGDA solutions were dispensed into rubber molds with approximately 10 mm diameter and 1.5 mm height and exposed to UV light (UVL-56 UV Lamp, 365 nm) for half an hour to ensure full gelation. PEG hydrogels were transferred to 12 well plates, swelled in 1 ml of DI water for an hour at 37°C prior to determining swelling properties.

Polymer Molecular Weight and Characteristic Ratio

Briefly, alginate polymers were prepared at 0.01, 0.05, 0.1 and 0.15% (w/v) in various NaCl solutions and were transferred to an immersion cup with a cone plate in a rheometer (HR3, TA Instruments) to determine viscosity. The polymer solutions were strained over a range of 5–50 hertz and values of viscosity were obtained from the linear region. A Huggins plot (η_{reduced}/c vs c) was constructed to determine the intrinsic viscosity of alginate polymers and the average molecular weight and characteristic ratio (C_n) were calculated from empirically derived equations in previous work⁴⁹ and reported.

Swelling, Polymer Volume Fraction and Rheological Characterization of Hydrogels

Alginate hydrogels were formulated from MVG or MVM alginate polymers as described above to obtain 1, 1.5, 2, 2.5 or 3% (w/v) final alginate polymer concentrations. Alginate hydrogels were then transferred to 24-well plates, each hydrogel was covered with 0.5 ml of DI water and incubated at 37°C for an hour. PEG hydrogels with 10, 15, 20 or 40% (w/v) final polymer concentrations were incubated at 37°C for an hour. Hydrogels were then removed ($n = 5-6$), and the hydrogels were weighed to obtain the wet weight (W_S). For swelling and polymer volume fraction studies, hydrogels were frozen overnight and lyophilized for at least 48 hours before weighing the hydrogels to obtain the dry weight of the polymer (W_D). The following equation (Equation 2) determined the volumetric swelling ratio:

$$Q = \frac{\frac{W_S - W_D}{\rho_W} + \frac{W_D}{\rho_P}}{\frac{W_D}{\rho_P}} \quad (2)$$

Where ρ_w is the density of water and ρ_p is the density of polymer as previously determined or specified by the manufacturer (Alginate: 1.6 g/ml^{16, 50}, PEG 1.13 g/ml). Polymer volume fraction $v_{2,3}$ is the volume the polymer occupies in the swelled hydrogel and was determined by taking the inverse of the volumetric swelling ratio. For fibrin hydrogels, the polymer volume fraction was determined from the ratio of fibrin concentration to the density of fibrin (0.4 g/ml)⁵¹. For rheological characterization, hydrogels were placed between parallel plates (axial force set at 0.01 N) in a rheometer. The hydrogels were strained from 0.001–5% at a frequency of 1 Hz and values of storage modulus (G') were obtained from the linear viscoelastic region. At least a dozen points were used to obtain the average G' value.

Mesh Size Calculations

Mesh size was estimated with two different approaches for alginate and PEG hydrogels. For PEG hydrogels, molecular weight between cross-links (M_C) was determined via the Peppas-Merrill equation^{5, 18} (Equation 3):

$$\frac{1}{M_C} = \frac{2}{M_n} - \frac{\bar{v}}{V_1} * (\ln(1 - v_{2,s}) + v_{2,s} + X_1 * v_{2,s}^2) / \left[v_{2,r} * \left[\left(\frac{v_{2,s}}{v_{2,r}} \right)^{\frac{1}{3}} - \frac{v_{2,s}}{2 * v_{2,r}} \right] \right] \quad (3)$$

Where M_n is the molecular weight of PEG (20,000), \bar{v} is the specific volume of the polymer (0.885 cm³/g for PEG), V_1 is the molar volume of the solvent (18 cm³/g), X_1 is the polymer-solvent interaction parameter (0.426 for PEG), $v_{2,s}$ is the polymer volume fraction in the swelled state and $v_{2,r}$ is the polymer volume fraction in the relaxed state.

Due to the anionic contributions of the alginate polymers, alginate hydrogel mesh size was determined from the swelling and rheometry data as previously described^{15, 20–21, 23–24}. M_C for alginate hydrogels was determined via the following equation¹⁹ (Equation 4):

$$M_C = C_p RT / G' \quad (4)$$

Where C_p is the polymer concentration, R is the gas constant and T is the measurement temperature. Mesh size (ξ) is then calculated as follows (Equation 5):

$$\xi = Q^{\frac{1}{3}} l \left(\frac{2M_C}{M_r} \right)^{\frac{1}{2}} C_n^{\frac{1}{2}} \quad (5)$$

Where l is the Kuhn length of the repeating unit (Alginate: 5.15 Å, PEG: 1.46 Å), M_r is the molecular weight of the repeating unit (Alginate: 194 g/mol, PEG: 44.05 g/mol) and C_n is the determined characteristic ratio. Mesh sizes were obtained using this equation for alginate and PEG hydrogels and were plotted with the mesh size predicted from the computational simulations described above.

Diffusion Experiments to Determine Mesh Size

Diffusion of a fluorescent tracer through hydrogels were performed to further validate mesh size as previously described with modifications⁵². Alginate hydrogels in 12-well plates with a square cutout (~ 1 cm²) in the center of the gel were monitored with a fluorescent microscope (Axio Vert A.4, Zeiss) at 4X magnification. Approximately 100 microliters of a 100 µg/ml solution of FITC-dextran (20 kDa, Sigma) were carefully transferred to the cutout region and time-lapse imaging of the alginate hydrogel adjacent to the cutout region was performed using a 488-nm excitation filter every minute for the course of 30 minutes. Images were taken immediately after the addition of FITC-dextran to the cutout region and the normalized fluorescent intensity of five equally spaced regions of the alginate hydrogel were determined for each image over the course of 30 minutes. The effective isotropic diffusion coefficient (D_{eff}) was determined from the data assuming that diffusion could be approximated as a one-dimensional, semi-infinite slab, and solving Fick's second law (Equation 6):

$$\frac{\partial C}{\partial t} = D_{eff} * \frac{\partial^2 C}{\partial x^2} \quad (6)$$

Assuming the following initial and boundary conditions (Equations 7–9):

$$t = 0, \quad C = 0 \quad \text{for all } x > 0 \quad (7)$$

$$x \rightarrow \infty, \quad C \rightarrow 0 \quad \text{for all } t > 0 \quad (8)$$

$$x = 0, \quad C = C_0 \quad \text{for all } t > 0 \quad (9)$$

Solving Fick's second law then yields the following solution as previously described⁵² (Equation 10):

$$\ln\left(\sqrt{t^3} * \frac{\partial C}{\partial t}\right) = -\frac{x_0}{4 * D_{eff} * t} + \ln\left(\frac{C_0 x_0}{4 * \sqrt{D_{eff}}}\right) \quad (10)$$

Where t is time (seconds), x_0 is the distance of the analyzed region to the edge of the reservoir and C_0 is the intensity of FITC-dextran. The derivative of concentration with respect to time was approximated using a finite difference formula (Equation 11):

$$\frac{\partial C}{\partial t} = \frac{C_{r+1} - C_r}{t_{r+1} - t_r} \quad (11)$$

C_r is the intensity measured for any region at time t_r . The diffusion coefficient was determined by plotting $1/t$ vs $\ln\left(\sqrt{t^3} * \frac{\partial C}{\partial t}\right)$ and applying linear regression to estimate the slope of the line ($-x_0/4D_{eff}$). D_{eff} was determined for each region and the mesh size of the hydrogel was then determined via the following equation²⁸ (Equation 12):

$$D_{eff} = \left(1 - 2 * \frac{r_s}{\xi}\right) * \exp\left(-y_p * \frac{\phi}{1 - \phi}\right) \quad (12)$$

Where r_s is the stokes radius of FITC-dextran in water (3.3 nm as specified by the manufacturer), y_p is approximated as one when the mesh size is much greater than r_s and ϕ is the polymer volume fraction. All images were analyzed with ImageJ (NIH) and the mesh size was calculated for each region and the average value per hydrogel ($n = 4$) was reported.

Design of Experiments (DOE)

A Box-Behnken DOE experimental design was created with JMP software (SAS institute) to analyze how combining both MVM and MVG alginate polymers at different concentrations effect hydrogel mesh properties. Input parameters included ratio of MVG vs MVM polymer

(0 to 1), G-block content of MVM alginate (30% to 50% G-block), G-block content of MVG alginate (60% to 100% G-block) and total polymer volume fraction (1% to 2%). Output parameters consisted of cross-link density and mesh size. The input variables were selected due to the effect G-block content of alginate polymers and polymer volume fraction has on modulating alginate hydrogel properties, including modulus, swelling and correspondingly mesh size⁵³. The input variables were examined at high, medium and low conditions based on the Box-Behnken model to generate prediction maps of the input variables effect on hydrogel mesh properties. The significance of the input parameters was determined via the response surface predictions generated by the DOE software as previously described⁵⁴. A full factorial approach was also utilized to determine how polymer radius and polymer volume fraction affect hydrogel mesh structure. Polymer chains with 0.5, 5, 10, 25, 50, 75 and 100 nm radii were modeled at 0.01, 0.025, 0.05, 0.075 and 0.1 polymer volume fraction to determine hydrogel mesh size. Heatmaps of the effect of inputs on hydrogel cross-link density and mesh size were plotted with MATLAB 2014a (Mathworks) and reported.

Confocal Reflection Microscopy

Confocal reflection microscopy was utilized to characterize the mesh size of fibrin hydrogels. Fibrin hydrogels were transferred to coverslips after gelation and were imaged with a Leica TCS SP8 STED 3x confocal microscope using the 615 nm laser line. Images were acquired for four regions of 2.5, 5, 7.5 and 10 mg/ml fibrinogen hydrogels and the average distance between adjacent cross-links was quantified with ImageJ (NIH) for at least 40 regions in each image to determine average mesh size. The average mesh size for each image was plotted with the mesh size predicted from the computational simulations described above.

Statistical Analysis

The coefficients of determination were calculated in order to determine the goodness of fit of the simulation model to the experimental data. DOE experiments were designed and analyzed using JMP 14.3 software (SAS institute) and significant interactions were asserted at $P < 0.05$. GraphPad Prism 8 software (GraphPad Software Inc.) was used to perform all regressions.

Results

Simulations Predict that Mesh Properties are Dependent on the G-Block Content of the Polymer and Polymer Volume Fraction.

A computational model of the alginate hydrogel was designed in order to predict various hydrogel mesh properties. The alginate hydrogel was modeled as overlapping randomly oriented cylinders and regions with overlapping G-blocks were designated as cross-links in order to determine cross-link density and mesh size (Fig. 1A). The computational model predicted that cross-link density increased exponentially for polymers with higher G-block content and higher polymer volume fractions, with cross-link density reaching approximately 188, 75, 38 and 12 μM for hydrogels composed of polymers with 100%, 70%, 50% and 30% G-block content respectively (Fig. 1B). Furthermore, the predicted mesh size from the computational model determined that hydrogel mesh size significantly

decreases with higher polymer volume fractions and with polymers containing higher G-block content (Fig. 1C). The data from the simulations and nonlinear regressions were used to generate the mesh size equation based on polymer volume fraction and G-block content (Equation 13):

$$\xi = S_f * e^{(-412.9 * v_{2,s})} + S_s * e^{(-44.1025 * v_{2,s})} \quad (13)$$

Where S_f and S_s determine the magnitude of the decay phases which are dependent on the fraction of G-block (G_P) content of the polymers given by the following equations (Equations 14–15):

$$S_f = (655.2 - 128.7) * e^{(-2.497 * G_P)} + 128.7 \quad (14)$$

$$S_s = (365.2 - 57.0) * e^{(-3.387 * G_P)} + 57.0 \quad (15)$$

The Computational Model Predicted the Mesh Size of Alginate Hydrogels with Varying Properties

The computational model closely predicted the mesh size of alginate hydrogels over a range of polymer and hydrogel properties. Alginate hydrogels with varying mechanical properties were prepared from polymers with different molecular weight and G-block content (Table 1). Hydrogels with higher polymer and G-block content were found to have an increased storage modulus, with 3% (w/v) hydrogels having a storage modulus of approximately 11.8, 32.5 and 51.1 kPa for MVM, MVG LMW and MVG HMW hydrogels respectively (Fig. 2A). The swelling ratio was found to significantly increase with lower polymer content, with all hydrogels experiencing an approximately 3-fold greater swelling for 1% (w/v) hydrogels compared to 3% (w/v) hydrogels (Fig. 2B). As expected, the polymer volume fraction was found to increase for hydrogels with higher alginate percentages (Fig. 2C). The Mesh size of hydrogels was determined from the characteristic ratio of the polymer (Table 1), swelling ratio and storage modulus and was compared to the predictions from the computational simulations (Fig. 2D). The simulations provided a high degree of correlation to the mesh sizes determined for MVG LMW ($R^2 = \sim 0.91$) and MVG HMW ($R^2 = \sim 0.96$) alginate hydrogels for polymer volume fractions higher than 0.01. The computational simulations also predicted the mesh size of hydrogels composed of MVM polymer, with polymer volume fractions above 0.01 showing a close correlation to MVM ($R^2 = \sim 0.87$) hydrogels (Fig. 2E).

Diffusion Based Methods for Determining the Mesh Size of Alginate Hydrogels Followed Model Predictions

The diffusion of FITC-Dextran within alginate hydrogels was observed to further validate the ability of the computational model to predict hydrogel mesh properties. FITC-dextran diffused from the reservoir into the surrounding hydrogel for 30 minutes (Fig. 3A). Alginate hydrogels were found to have slower diffusion of FITC-Dextran, and correspondingly

smaller mesh sizes, with higher polymer volume fractions and for hydrogels comprised of polymers with higher G-block content. Hydrogels comprised of MVG polymers were determined to have diffusion derived mesh sizes close to the mesh size predicted from the computational model, with MVG HMW hydrogels showing a closer correlation to simulations than MVG LMW hydrogels (Fig. 3B). The simulations also provided a reasonable approximation of the diffusion derived mesh size of hydrogels comprised of MVM polymers (Fig. 3C).

Simulations Provided Information on the Mesh Size Distribution of Alginate Hydrogels

The computational model provided a method to characterize the mesh size distribution of alginate hydrogels. The mesh size distribution was predicted for 1, 2 and 3% (w/v) alginate hydrogels composed of MVG HMW, MVG LMW or MVM alginate polymers (Fig. 4 A–C). In general, the simulations predicted that the mesh size distribution had a higher maximum frequency and lower standard deviation for hydrogels with higher alginate percent (w/v). Furthermore, hydrogels with less G-block content were observed to have a larger distribution of mesh sizes and a lagging region of mesh sizes extending to over 200 nm. The simulation data was used to determine $P(\xi)$, which is the predicted percent of the total mesh structure with a specific mesh size. $P(\xi)$ was also observed to have a greater amplitude, decreased standard deviation and smaller mean for hydrogels with more G-block content and alginate percent (w/v).

Prediction of Alginate Hydrogel Mesh Properties by Experimental Design

A DOE approach could be used in conjunction with the computational model to predict further mesh properties of alginate hydrogels consisting of polymers with both high and low G-block content. The Box-Behnken DOE model generated 27 unique compositions of alginate hydrogels to model, and the computational simulations predicted the corresponding cross-link density, mesh size and mesh size distribution for each condition. The polymer volume fraction, the ratio of MVG to MVM polymer, the G-block content of MVG polymer and the G-block content of MVM polymers were determined to affect both hydrogel cross-link density and mesh size. The DOE software predicted that cross-link density could be varied from approximately 5 to 100 μM through varying the polymer volume fraction and G-block content of the MVG and MVM alginate polymers (Fig 5A). The mesh size was predicted to be inversely correlated to cross-link density, with simulated mesh sizes varying from approximately 30 to 90 nm through altering the input parameters (Fig 5B).

The Computational Model Predicts the Mesh Size of Fibrin and PEG Hydrogels

The computational model also provided a close prediction of mesh properties for other hydrogel systems. Hydrogels with 2.5, 5, 7.5 or 10 mg/ml of fibrinogen were imaged with confocal reflection microscopy to determine hydrogel mesh size (Fig 6A). The mesh size determined for the fibrin hydrogels were found to range from approximately 2.2 μm —5.6 μm and were very closely predicted ($R^2 > 0.99$) by the computational model (Fig 6B). The mesh size of PEG hydrogels with varying polymer concentrations was also compared to the computational model, with the simulations providing a high degree of correlation ($R^2 = \sim 0.95$) to PEG hydrogels (Fig. 6C).

DOE Strategy to Design Hydrogels with Predefined Mesh Sizes Through Controlling Properties of the Polymer Network

A full factorial DOE approach was utilized to determine how the size of the polymer chains affects hydrogel mesh size. The full factorial model generated 35 unique compositions of hydrogels to model, and the computational simulations predicted the mesh size for each condition. The polymer volume fraction and the radius of polymer chains were determined to have a significant effect on the mesh size. The DOE results predicted that the mesh size was proportional to the polymer chain radius and could be varied from approximately 5 nm to 5.5 μm through varying the polymer chain radius and the polymer content (Fig 7). The data from the DOE were used to predict how mesh size depends on the polymer volume fraction and the polymer chain radius (R_P) given by the following equation (Equation 16):

$$\xi = (72.79 * R_P + 2.01) * e^{(-89.49 * v_{2,s})} + (27.11 * R_P + 0.08) * e^{(-8.36 * v_{2,s})} \quad (16)$$

Discussion

This study investigates the utility of a novel computational model that mimics the hydrogel mesh structure and allows one to determine the hydrogel mesh properties critical for the design of hydrogels systems as therapeutic delivery vehicles. This work demonstrates that the computational model could accurately predict the mesh size of alginate hydrogels over a broad range of polymer and hydrogel properties. Additionally, the model provided information about the mesh size distribution of hydrogels and provided adjustable parameters to design alginate hydrogels with controllable mesh properties. Finally, the computational model also provided a close prediction of the mesh size for fibrin and PEG hydrogels and was further extended to predict the mesh properties of other hydrogel systems. To our knowledge, this is the first study of a computational model of the hydrogel polymer network to predict hydrogel mesh properties as a strategy to provide control over therapeutic delivery applications.

This work involved designing a computational model of the alginate hydrogel. In particular, the mesh size of hydrogels has an essential role in determining the diffusion of encapsulated therapeutic agents and regulating mass transport for cell-based therapies^{5-6, 18}. In this study, we modeled the alginate polymers as a collection of rigid rods because the polymers in hydrogels experience tension following gelation due to various swelling forces acting on the polymer system⁵⁹, with alginate hydrogels forming very rigid chains following calcium crosslinking⁵⁵. Moreover, alginate polymers have previously been shown to be reasonably modeled as rigid rods, especially with polymers containing larger M block or G block regions⁵⁶⁻⁵⁸. Taken together, and while understanding some limitations, we still believe that modeling the crosslinked alginate polymers as rigid rods provides a computationally accessible strategy to approximate properties of the polymer system and derive hydrogel mesh properties. The model also determined cross-linking regions and correspondingly mesh size based on the egg-box model, where the α -l-gulonate (G) residues in the alginate polymer chain adopt a 2_1 helical conformation and the pair of G residues (“binding cavity”) surrounds the calcium ions⁶⁰. Previous work has shown that cross-link density significantly increases for alginate hydrogels with more calcium^{16, 61-62} or more polymer content²⁸.

Similarly, this model predicted that the regions of overlapping G-blocks (i.e. cross-links) dramatically increased for hydrogels with higher polymer percentage and G-block content. Furthermore, the cross-link density predicted from the simulations for hydrogels with higher G-block content followed a similar trend to theory derived from Flory-Rehner^{16, 63–64}. Finally, the model predicted that hydrogel mesh size increases for hydrogels comprised of lower G-block content and polymer volume fractions. Thus, this work provides a computational model that can incorporate properties of alginate hydrogels to provide a theoretical representation of the hydrogel mesh structure.

This study demonstrates that the model can accurately predict the mesh size of alginate hydrogels. Previous work has shown that alginate hydrogel properties can be modulated through controlling polymer properties, including molecular weight, polymer content and the ratio of M/G-blocks^{16, 21, 24, 28, 53, 65–66}. Thus, in this work, hydrogels with varying polymer concentrations were prepared from polymers of various molecular weight and G-block content to validate the model. The mesh size of alginate hydrogels was determined from mechanical properties^{15, 20, 23–24} and from the diffusion of FITC-dextran through alginate hydrogels^{28, 52}. Our results found that the mesh size determined from the simulations could accurately predict the mesh size of various alginate hydrogel compositions. Previous work has also shown that alginate hydrogels can display a wide range of mesh sizes from ~10 to 100 nm depending on the polymer and calcium cross-linking concentrations^{15–16, 20, 22, 26, 28, 65}. The computational model was found to closely predict hydrogels with higher polymer content and also provided a good prediction of the mesh properties from other work using alginate hydrogels with greater calcium concentrations and higher polymer volume fractions⁶⁷. However, the model underestimated the mesh size of hydrogels with lower polymer volume fractions. This is possibly the result of the simulations modeling the polymer structure as a collection of rigid cylinders, which is likely more valid for stiffer and more mechanically stable hydrogels composed of higher polymer volume fractions⁶. Overall, the computational model provided a robust prediction of the mesh size of alginate hydrogels for a wide range of hydrogel compositions and mechanical properties.

The computational model was then applied to predict the mesh size distribution of alginate hydrogels. Mesh size distribution has an important role in determining the probability of a solute encountering a sufficient opening within the hydrogel to allow for diffusion, the maximum size of therapeutic agents that can diffuse through the mesh structure and also has a role in modulating encapsulated cell and native tissue properties^{15–16, 68–70}. Previous studies using NMR to characterize the mesh size distribution of alginate hydrogels composed of 2% (w/v) MVG polymers have shown a broad peak of mesh sizes near 30 nm and a lagging mesh region extending over 100 nm^{7, 71}. The simulations determined a similar prevalence of mesh sizes between approximately 10 to 50 nm and a lagging region of mesh sizes exceeding 100 nm. The simulations also predicted mesh size distributions similar to other work with interpenetrating alginate hydrogels, with the model more closely following mesh size distributions predicted from cryoporometry²⁶. This study found that the model could determine hydrogel mesh size distributions.

The results of this study predicted how varying properties of binary hydrogels effects hydrogel cross-link density and mesh size. Hydrogels consisting of both MVG and MVM polymers have been shown to exhibit different properties than hydrogels comprised of only MVG or MVM polymer, allowing for additional control over hydrogel mechanical properties^{72–73}. In this study, a DOE design was utilized in conjunction with the computational model to predict how factors affected MVG and MVM binary hydrogel mesh properties. The ratio of MVG to MVM, polymer volume fraction and G-block content of the polymers were determined to have the most effect on both cross-link density and mesh size. In summary, the computational model determined important properties to control when designing alginate hydrogels with desired mesh properties.

This work also demonstrated that the computational model could be extended to predict the mesh properties of additional hydrogel systems. Fibrin and PEG hydrogels were used to further validate the model for hydrogels systems over a broad range of mesh properties. Fibrin is a naturally occurring polymer that aggregates into thick fibers and typically has hydrogels with mesh sizes in the micron range^{74–75}. PEG is a synthetic polymer with a small polymer chain radius and forms nanoporous hydrogels^{47–48, 76}. While the model does not account for structural defects and network non-ideality that can be present in PEGDA (or PEG-dimethacrylate) hydrogels^{77–78}, the computational model accurately predicted the mesh size of both fibrin and PEG hydrogels. In this study, PEGDA hydrogels were prepared from polymers with high molecular weight and have a greater fully extended polymer length following gelation and swelling⁷⁹ which could be reasonably represented by the unidirectional cylinders in the model. Furthermore, the higher PEG concentrations used in this work helped reduce the prevalence of network defects⁸⁰, with certain defects including intra-chain cycles and dangling ends not contributing to the network-dependent properties of the hydrogel⁸¹. Interestingly, previous work has also modeled hydrogels made from Peg-*b*-PLA with end-capped acrylate functionalities as ideal networks to predict hydrogel properties including degradation⁸². Nevertheless, and while understanding some limitations, the model still provided a reasonable estimate of the mesh properties of other fibrin^{75, 83–84} and PEGDA^{47, 77, 85} hydrogels, especially for PEGDA hydrogels with higher polymer molecular weight and polymer volume fractions. The model was then extended to predict the mesh structure of other hydrogel systems from the polymer radius and content. Interestingly, the simulations determined that hydrogels with a larger polymer chain radius had significantly greater mesh sizes. Other work has also observed that hydrogels comprised of polymer chains with a small polymer radius (< 1 nm) commonly have mesh sizes in the tens of nanometer range^{85–86}, while hydrogels comprised of polymers with a larger polymer fiber radius can have pore sizes into the micron range^{87–88}. Taken together, the computational model can closely predict the mesh properties of multiple hydrogel systems and provides a robust strategy to predict the mesh size of hydrogels.

Conclusions

This study investigates the use of a computational model of the hydrogel mesh structure to predict important hydrogel properties conducive to therapeutic delivery. This work found that the model could incorporate various parameters of alginate hydrogels, including polymer fraction and G-block content, to predict cross-linking density, mesh size and mesh

size distribution. The results from the simulations were found to closely correlate to the actual mesh size of alginate hydrogels, with higher polymer fractions providing a better correlation to mesh size. Furthermore, the simulations could predict the mesh size of hydrogels combining polymers with varying G-block content. Finally, the computational model was validated for fibrin and PEG hydrogel systems and was extended to predict the mesh properties of hydrogels from the polymer network. Future work will include applying additional modeling strategies and crosslinking paradigms into the simulations to further characterize specific hydrogel systems. Overall, this model provides a robust and accessible strategy to predict the mesh size of hydrogels and could have applications for designing hydrogels with controllable mesh properties for potential therapeutic delivery applications.

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Computation Model of Alginate Hydrogels

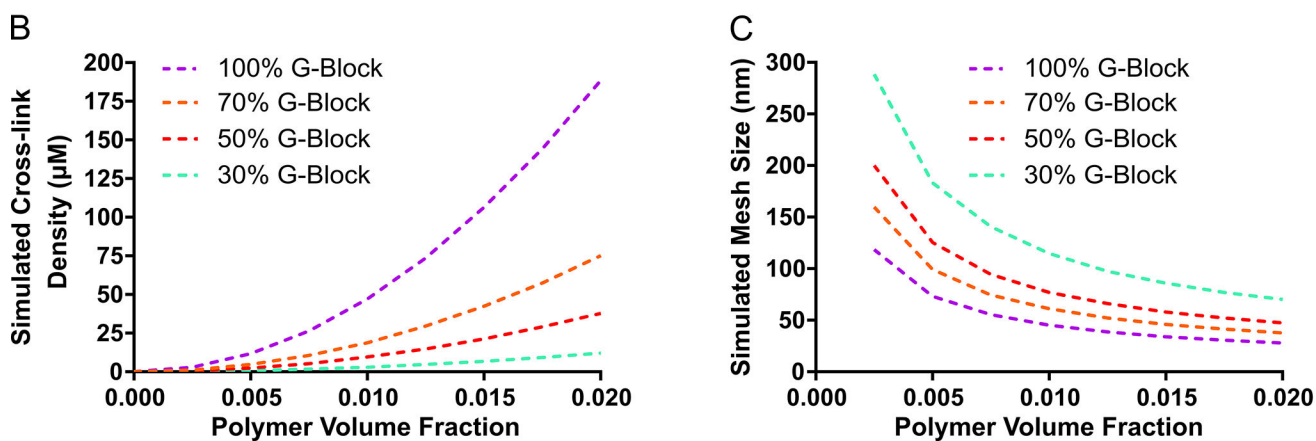
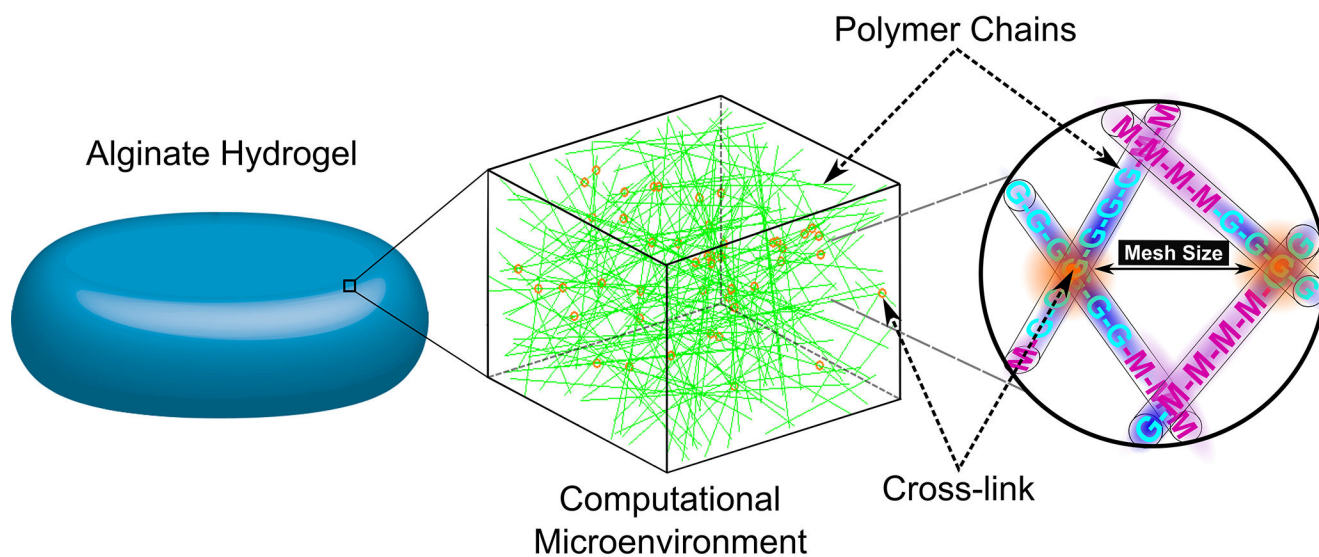


Fig 1. Design of a Computational Model that Mimics the Hydrogel Polymer Network and Enabled the Prediction of the Cross-link Density and Mesh Size of Various Alginate Hydrogels. The cross-link density and mesh size of alginate hydrogels composed of various polymer volume fractions and percent G-block content were determined via computational simulations. A schematic of the computational model of the hydrogel is provided (A). The cross-link density of alginate hydrogels was predicted to exponentially increase for hydrogels with polymers containing higher G-block content (B). The model anticipated a wide range of alginate hydrogel mesh sizes based on polymer volume fraction and G-block content (C). In figure panel B and C, dashed lines represent mean. (B–C, n = 10).

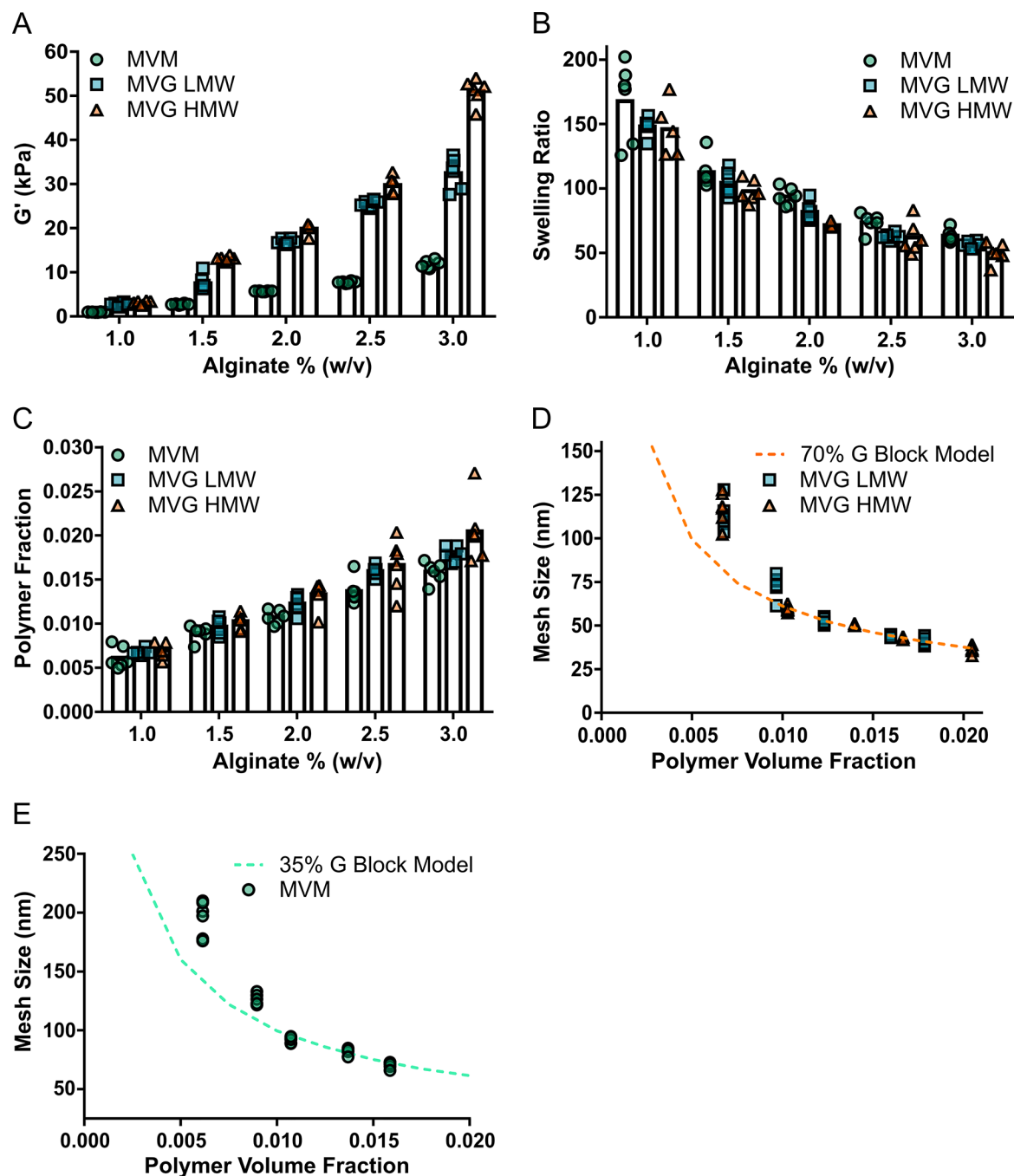


Fig 2. The Computational Model of the Hydrogel Polymer Network Provided a Strategy to Closely Estimate the Mesh Size of Alginate Hydrogels

The computational model of the hydrogel mesh size provided a strong correlation to the mesh size of alginate hydrogels comprised of different mechanical and polymer properties. The storage modulus increased for hydrogels with greater alginite percent, G-block content and polymer molecular weight (A). Hydrogels with greater polymer content were determined to experience less swelling (B) and greater polymer volume fractions (C). The mesh size of MVG (high G-block content) alginate hydrogels were closely predicted by the computational model (D). The computational model provided a good estimation of the mesh

size of the MVM (low G-block content) alginate hydrogels (E). In figure panel A, B and C, bar represents mean, scatter dot plots display individual measurements and error bars represent standard deviation. In figure panel D and E, scatter dot plots display individual measurements and dashed lines represent mean values determined from the simulations (A–E, $n = 5-6$).

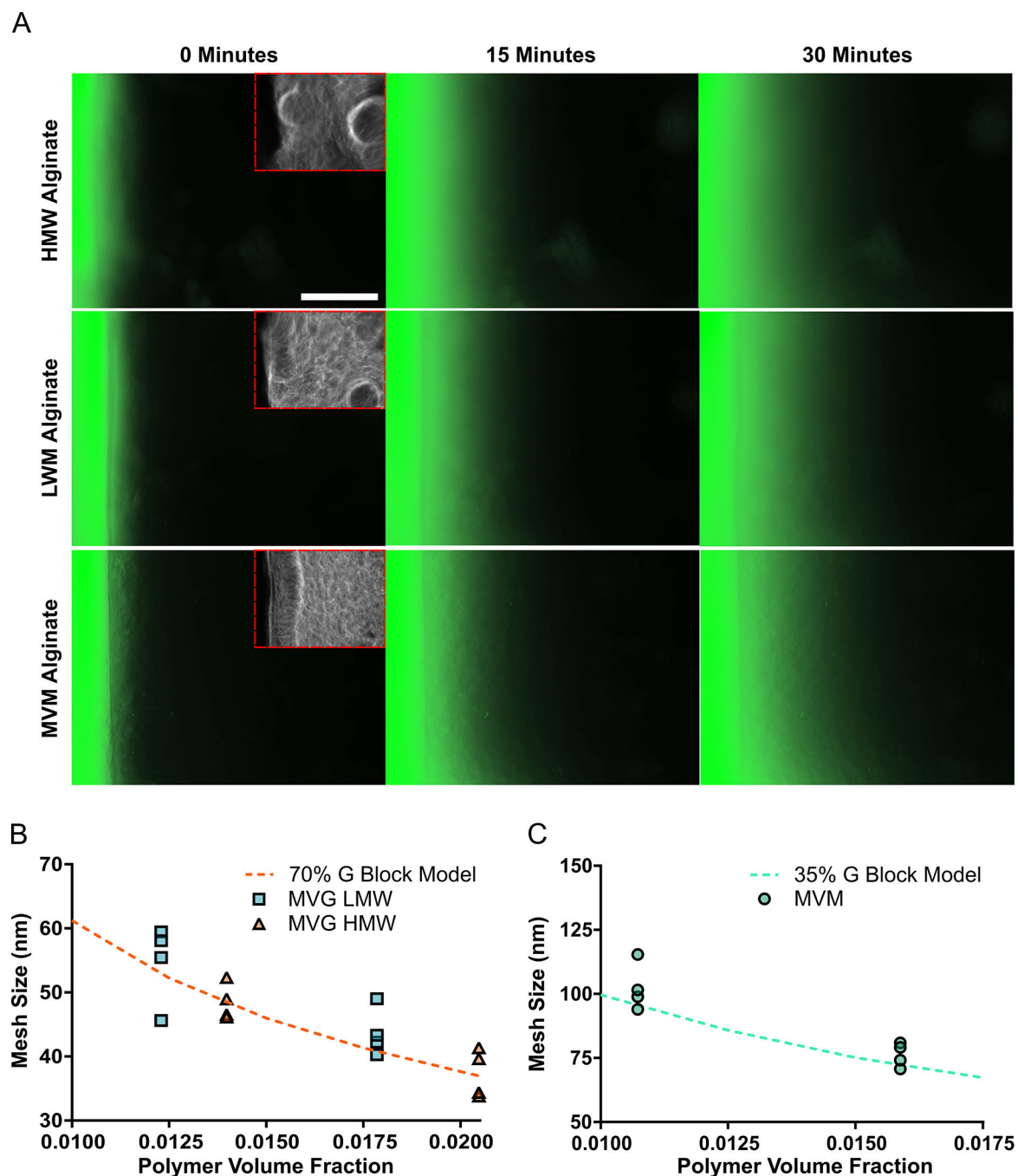


Fig 3. A Diffusion-Based Strategy for Determining Alginate Hydrogel Mesh Size Further Validated the Computational Model's Ability to Predict Hydrogel Mesh Properties

The mesh size of alginate hydrogels determined from the diffusion of FITC-dextran followed the mesh size predicted from the computational model. The diffusion of FITC-dextran from a reservoir into various alginate hydrogels was observed for 30 minutes. Scale bar represents 5 mm (A). The mesh size of MVG (high G-block content) hydrogels was determined from the effective diffusion coefficient of FITC-dextran within the polymer system and had similar mesh sizes predicted by the simulations (B). The computational model reasonably predicted the mesh size of MVM (low G-block content) hydrogels

determined from the diffusion of FITC-dextran through the polymer system (C). In figure panel B and C, scatter dot plots display individual measurements and dashed lines represent mean values determined from the simulations (B–C, $n = 4$).

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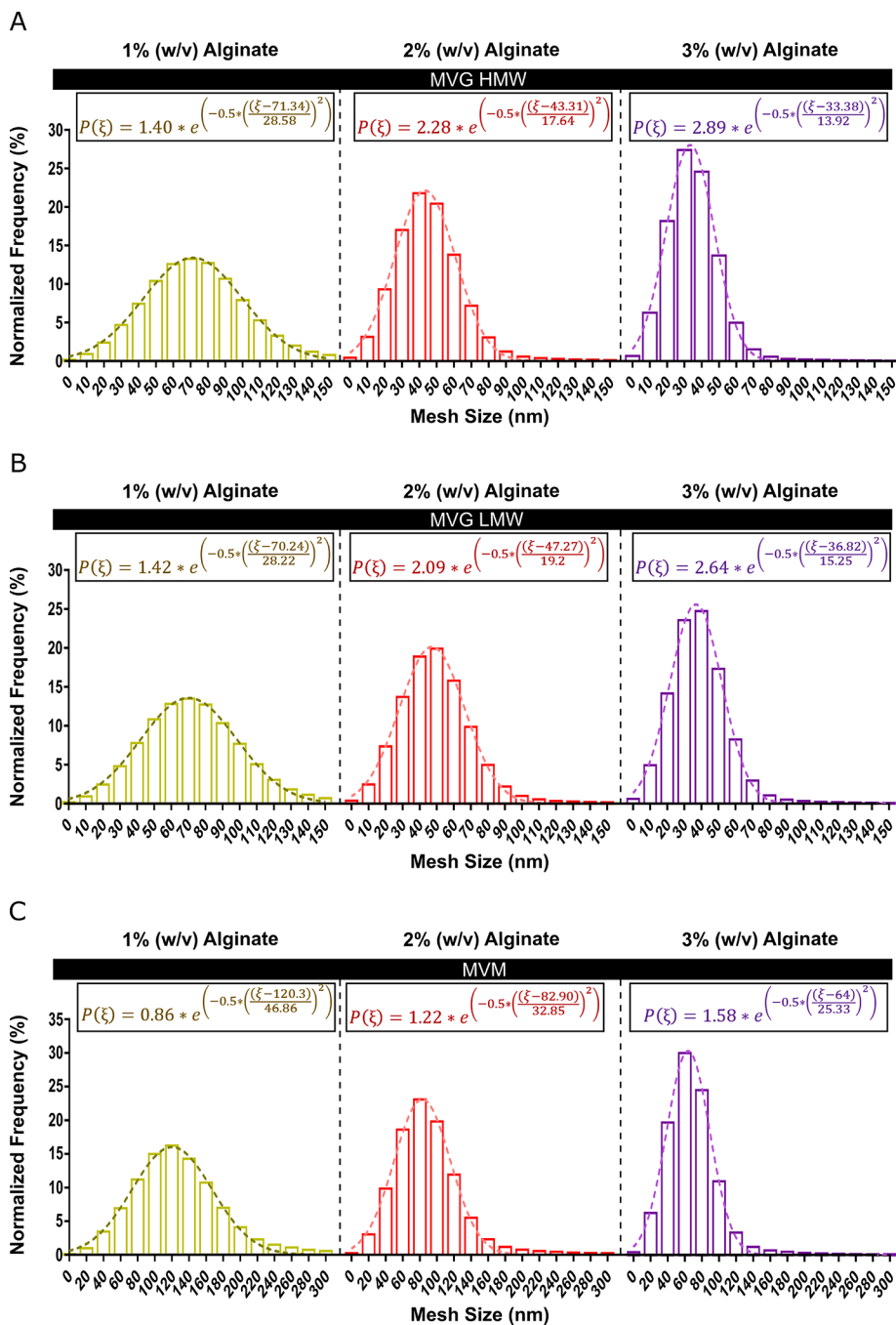


Fig 4. The Computational Model Provided a Method to Readily Characterize the Mesh Size Distribution of Various Alginate Hydrogels

The computational model determined the mesh size distribution and the probability of finding a mesh size ξ for 1, 2 or 3% (w/v) MVM (low G-block content) or MVG (high G-block content) hydrogels. The mesh size distribution was found to experience greater amplitudes, lower means and decreased standard deviations for MVG hydrogels with greater polymer volume fractions (A–B). MVM hydrogels were predicted to have a lagging region of large mesh sizes extending over 300 nm (C). (A–C) Dashed lines were determined from nonlinear regression (A–C, $n = 10$).

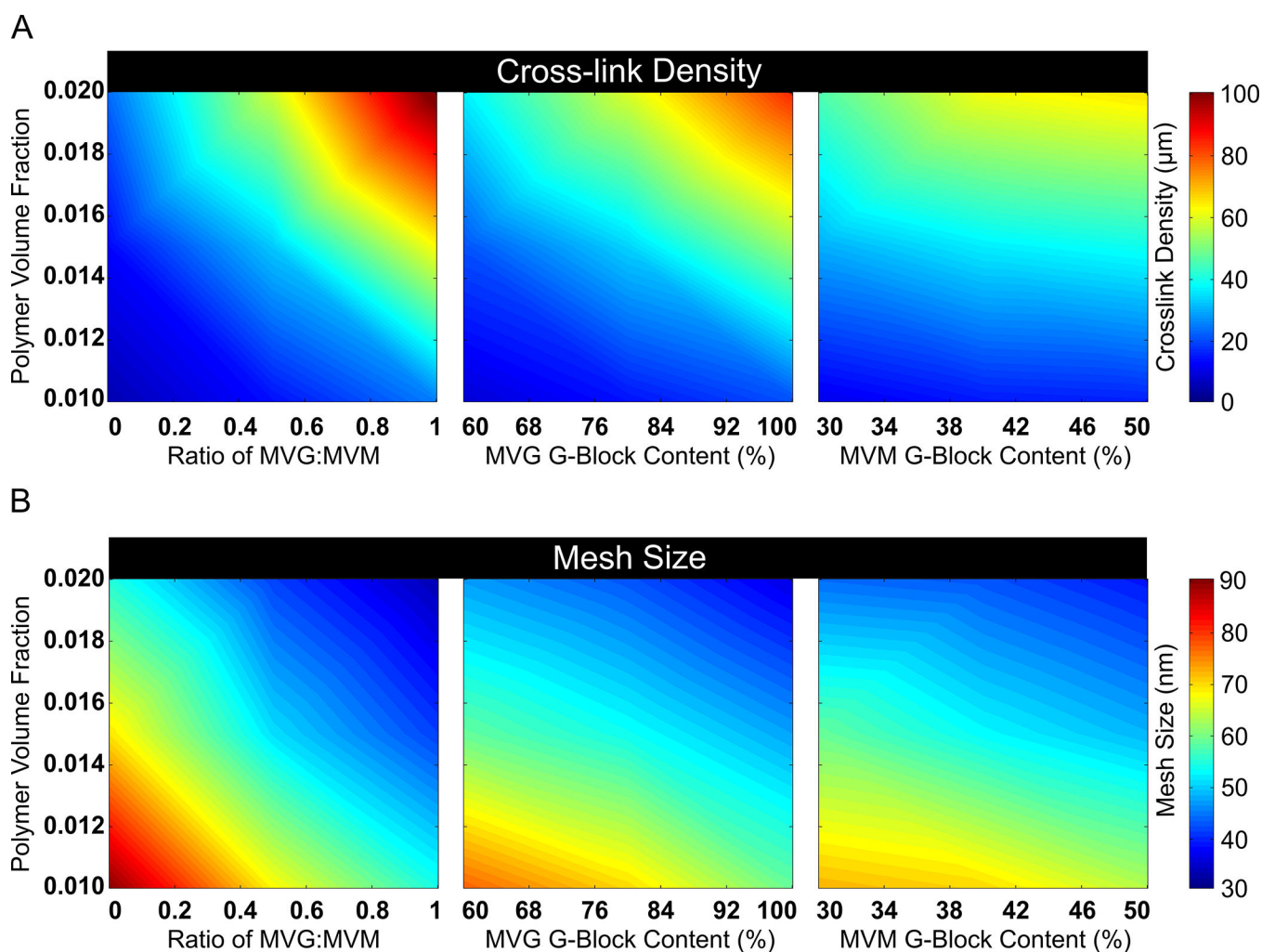


Fig 5. DOE Approach Predicted Hydrogel Design Properties to Control the Mesh Structure of Binary Alginate Hydrogels

DOE was used to identify properties that could alter the cross-link density and mesh size of alginate hydrogels with MVM (low G-block content) and MVG (high G-block content) polymers. Response surface maps of hydrogel properties effect on cross-linking density are provided (A). Altering polymer volume fraction, the ratio of MVG to MVM polymers and G-block content of hydrogels predicted a large range of adjustable hydrogel mesh sizes (B). In figure panel A, the heat color map represents the range of cross-link density. In figure panel B, the heat color map represents the mesh size range (A–B, n = 10).

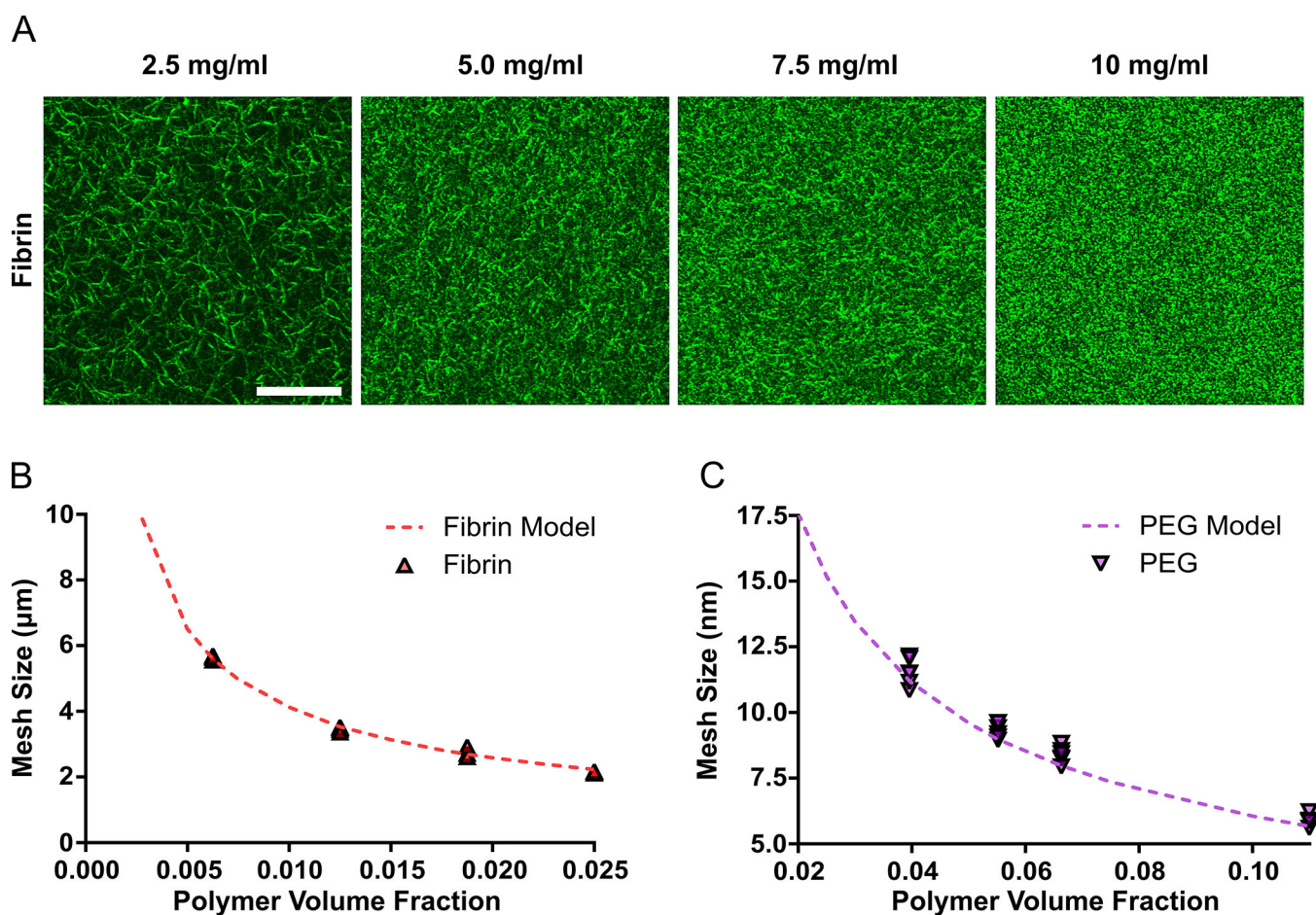


Fig 6. The Computational Model Simulated the Mesh Structure of Other Hydrogel Systems

The computational-based strategy for determining hydrogel mesh size was further validated for fibrin and PEG hydrogels. Confocal reflection of the fibrin meshwork for 2.5, 5, 7.5 and 10 mg/ml hydrogels is provided. Scale bar represents 20 μm (A). The computational model provided a very close estimation of fibrin hydrogel mesh size determined from the confocal images (B). The mesh size of PEG hydrogels was accurately predicted by the simulations for varying PEG concentrations (C). In figure panel B and C, scatter dot plots display individual measurements and dashed lines represent mean values determined from the simulations (B–C, $n = 4-6$).

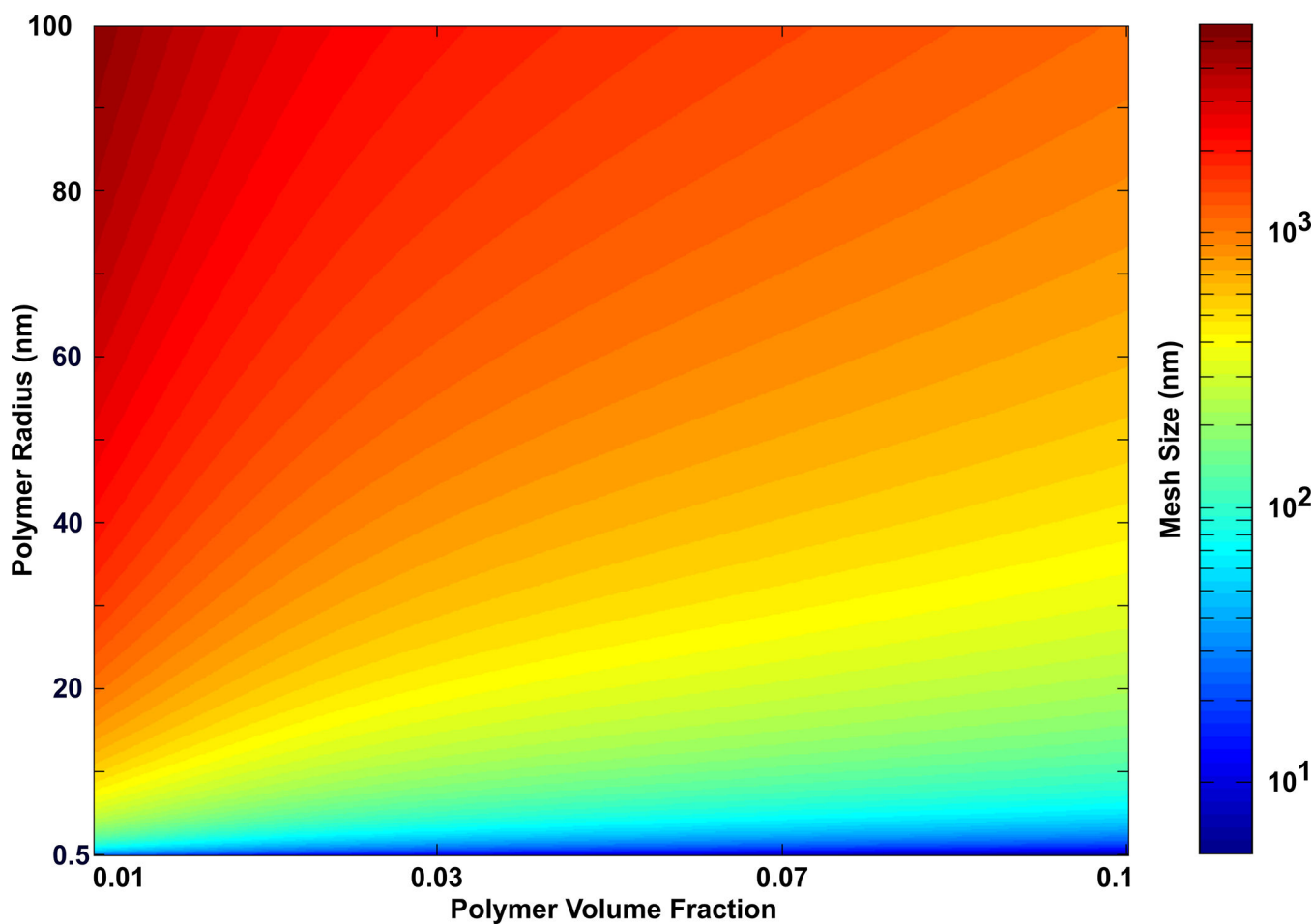


Fig 7. DOE Provided a Strategy to Predict the Mesh Size of Additional Hydrogel Systems

A full factorial DOE approach was utilized to predict the mesh size of hydrogels from the properties of the polymer system. The mesh size of hydrogels was predicted to increase with both polymer volume content and the polymer chain radius, with the polymer chain radius having the most effect on mesh size for the conditions tested. The heat color map represents the mesh size range ($n = 10$).

Table 1.

G-block content, Intrinsic Viscosity, Average Molecular Weight (MW) and Characteristic Ratio of Alginates

Alginate	G-block (%)	Intrinsic Viscosity (dL/g)	MW (kg/mol)	Characteristic Ratio (C_n)
MVG HMW	70%	3.94	238	22.96
MVG LMW	70%	1.17	41	18.81
MVM	35%	0.91	22	18.41

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