



BRIEF REPORT

REVISED Image analysis method for heterogeneity and porosity characterization of biomimetic hydrogels [version 2; peer review: 2 approved]

Maryam Jamshidi , Cavus Falamaki 

Chemical Engineering Department, Amirkabir University of Technology, P.O. Box 15875-4413, Tehran, Iran

V2 First published: 15 Dec 2020, 9:1461
<https://doi.org/10.12688/f1000research.27372.1>

Latest published: 12 Apr 2021, 9:1461
<https://doi.org/10.12688/f1000research.27372.2>

Abstract

This work presents an image processing procedure for characterization of porosity and heterogeneity of hydrogels network mainly based on the analysis of cryogenic scanning electron microscopy (cryo-SEM) images and can be extended to any other type of microscopy images of hydrogel porous network. An algorithm consisting of different filtering, morphological transformation, and thresholding steps to denoise the image whilst emphasizing the edges of the hydrogel walls for extracting either the pores or hydrogel walls features is explained. Finally, the information of hydrogel porosity and heterogeneity is presented in form of pore size distribution, spatial contours maps and kernel density dot plots. The obtained results reveal that a non-parametric kernel density plot effectively determines the spatial heterogeneity and porosity of the hydrogel.

Keywords

image processing, transformation, noise, filter, pixel, frequency, kernel, distribution, hydrogel, pore



This article is included in the **NEUBIAS - the Bioimage Analysts Network** gateway.

Open Peer Review

Reviewer Status  

Invited Reviewers

1

2

version 2

(revision)
12 Apr 2021



report



report



version 1

15 Dec 2020



report

1. **Miroslava Dušková-Smrčková** , Institute of Macromolecular Chemistry, Academy of Sciences of the Czech Republic, Prague, Czech Republic
2. **Susana Rocha** , KU Leuven, Leuven, Belgium

Any reports and responses or comments on the article can be found at the end of the article.

Corresponding author: Cavus Falamaki (c.falamaki@aut.ac.ir)

Author roles: **Jamshidi M:** Conceptualization, Investigation, Methodology, Software, Visualization, Writing – Original Draft Preparation; **Falamaki C:** Writing – Review & Editing

Competing interests: No competing interests were disclosed.

Grant information: The author(s) declared that no grants were involved in supporting this work.

Copyright: © 2021 Jamshidi M and Falamaki C. This is an open access article distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

How to cite this article: Jamshidi M and Falamaki C. **Image analysis method for heterogeneity and porosity characterization of biomimetic hydrogels [version 2; peer review: 2 approved]** F1000Research 2021, 9:1461
<https://doi.org/10.12688/f1000research.27372.2>

First published: 15 Dec 2020, 9:1461 <https://doi.org/10.12688/f1000research.27372.1>

REVISED Amendments from Version 1

There are some clarifications made to better explain the hydrogel material network and the artefacts associated with the imaging. Furthermore, it is mentioned how to use the code to extract hydrogel pore or wall thickness features. Lastly, the result of measurement with the method is compared quantitatively with the reported values in the source paper to validate the method. Figure 1 and Figure 2 have been updated.

Any further responses from the reviewers can be found at the end of the article

Introduction

Biomimetic engineered hydrogels often serve as 3D microporous extracellular microenvironment mimics in regenerative medicine, tissue engineering¹ and *in-vitro* cancer studies². The physical properties of these gels may provide a mechanical cue to regulate the cell phenotypic activities and functions via cellular mechanotransduction^{3,4}. Moreover, it has been shown that a change in the stiffness/elasticity of hydrogel is associated with the morphological change in the structure of the hydrogel mesh⁵. This article presents an image analysis method applied to characterise hydrogel structure heterogeneity and porosity resulting from the treatment of fully hydrated hydrogels

during plunge-freezing to acquire their cryogenic scanning electron microscopy (cryo-SEM) images. It should be mentioned that during the process of sample preparation for cryo-SEM microscopy, the sample undergoes a probable morphological alteration, furthermore the process itself might introduce artifacts according to the nature of the material and swelling rate. Readers are encouraged to refer to the studies specialized on the imaging of different hydrogel network by Kalasova *et al.*⁶ and Pradny *et al.*⁷.

Methods

Image processing

The image processing algorithm of the present work has been written as a code in the Python 3.8 language and was applied to analyse the cryo-SEM images of hydrogel, however it is applicable to analyse any other type of images of hydrogel porous network. The present study uses cryo-SEM images of fully hydrated hydrogels adopted with permission from Kaberova *et al.*⁸ as input data. To detect the pores precisely, the edges of the hydrogel walls are highlighted, and a band pass frequency filter is applied to optimize the removal of noise with preservation of the edges of hydrogel walls. To detect and measure the thickness of the hydrogel walls, the look up table of the binary image should be inverted to bring hydrogel walls in foreground (white) refer to Figure 1.

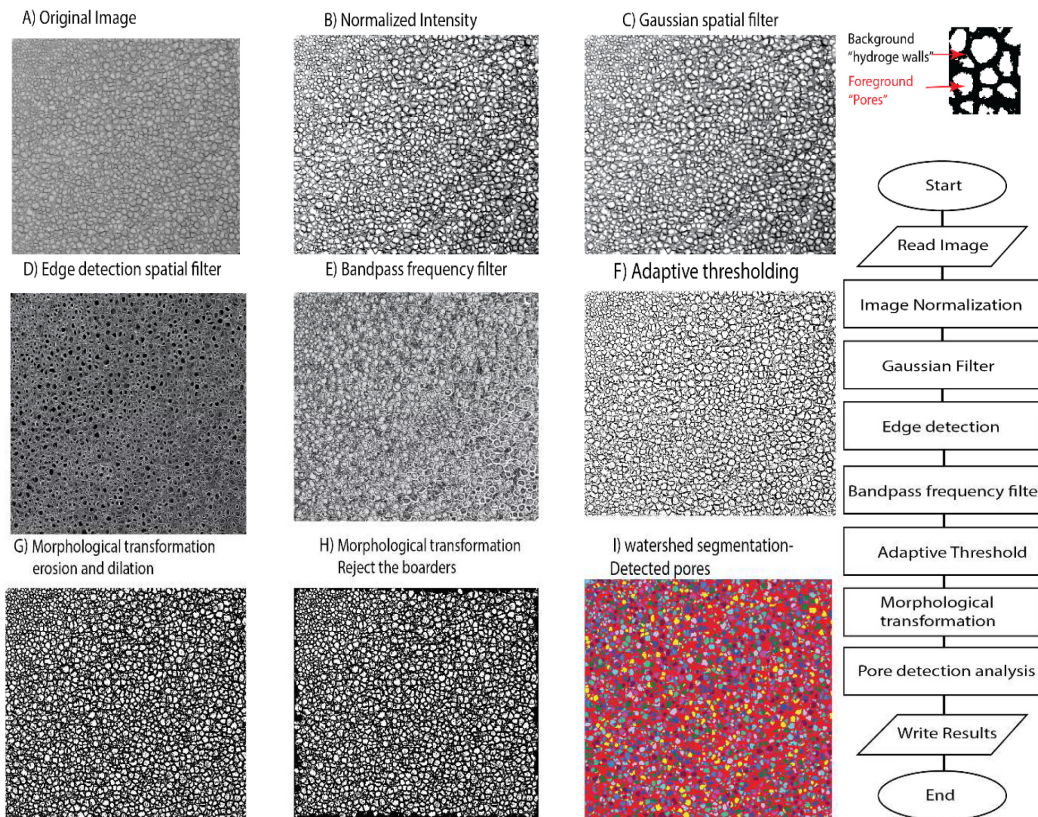


Figure 1. Image processing algorithm of sample cryo-SEM micrograph of glycidyl methacrylate hydrogel crosslinked with 0.3 mol% (ethylene glycol) dimethacrylate. Images on the left show the output images of each step in the processing algorithm and the relevant flowchart is illustrated on the right to distinguish foreground (pores) from background (hydrogel walls). The source image adapted from Kaberova *et al.*⁸ with permission.

Pre-processing. After loading the image from the specified path, it was first normalized to stretch the Gray level histogram between 0 and 255 and enhance the image contrast. Next, to filter out the noise, a Gaussian ($\sigma = 0.7$, 3×3) spatial filter convolution was applied to the image. The filtered image is a weighted average of the neighbourhood pixels that better preserve the edges in lower contrast areas while removing the noise⁹. Afterwards, to extract the edges of the hydrogel walls, a range of the non-linear 3×3 edge detection filters including Sobel was applied to the image to highlight the locations of sharp intensity transitions. Both the noise and edges belong to the high pass frequencies of the image in essence¹⁰ and the edge emphasising filter might introduce some artifacts in these range. Therefore, in the next step, a band pass frequency filter was used to highlight the desired quasi- high range of the frequencies for a better edge detection (refer to Figure 1).

Thresholding. The binarization of cryo-SEM images of gels is challenging as there is no standard method for thresholding. Since in most of the cryo-SEM images there is uneven illumination, the adaptive threshold technique was undertaken to segment the background (hydrogel walls, value=0 black) and foreground pores (pores, value=1 white). The algorithm calculates the optimal value of threshold based on the weighted mean and standard deviation of the pixel values within the neighbouring window of fixed size for each pixel and outperforms the conventional methods¹¹. In the present work, the adaptive Gaussian threshold has been applied to the image on a window size of 25×25 pixels. The window size should be optimized depending on the density of the details (information) and a lower or greater size may be chosen.

Morphological transformation. The pores touching the borders of the image were excluded and then basic morphological transformation (erosion $\times 1$ and opening $\times 5$) was applied to ensure that the isolated pixels both in the background and foreground (pores) were eliminated. For all transformations, a 3×3 elliptical structuring element was applied. Erosion transformation was applied to separate the touching pores and remove the remaining very small pores. Following, the holes within the detected pores were filled up. And lastly, the image was reconstructed based on the erosion and opening results and then a watershed algorithm was used to segment the pores and measure pore properties on the final image.

To validate the method, the screenshot images of a hydrogel network from a previously published work where the corresponding pore diameters have been reported, were analysed with the proposed method using Otsu's thresholding method and setting 0.05 max watershed threshold. The obtained results of the proposed method ($17 \mu\text{m}$) were compared

with the available reported measurement ($15 \mu\text{m}$ from Figure 4 of the reference)¹².

Hydrogel porosity and heterogeneity analysis

After performing the pore detection analysis, each single detected pore was associated with the centre of mass and area and the results were exported in a text file. The cryo-SEM image of the sample hydrogel shown in Figure 1A reveals that the hydrogel structure is heterogenous in spatial domain. Therefore, pore size distribution and statistical analysis alone might not represent the spatial heterogeneity and clustering of the detected pores. To quantify and visualize the spatial heterogeneity of the hydrogel, the kernel density estimation function was fitted on the centre of mass of the detected pores on the spatial domain of the cryo-SEM image.

Results

Comparing the obtained results of the proposed method with the available reported measurements of the pore sizes of fluorescent images of hydrogel elsewhere¹², the method has been validated with acceptable accuracy ($17 \mu\text{m}$ compared to the reported value was $15 \mu\text{m}$ for average pore size). Moreover, the average equivalent diameter of the pores of the analysed source image (adopted from Kaberova *et al.*⁸) was equal to $12.36 \mu\text{m}$ with the corresponding pore size distribution range of $2\text{--}36 \mu\text{m}$. The comparison of the results with the range reported by Kaberova *et al.*⁸ ($2\text{--}40 \mu\text{m}$) shows a good agreement and reliability of the presented method as it is shown in Figure 2B. The results of spatial heterogeneity quantification are presented in the form of contour plots (Figure 2). A higher density value indicates the presence of more pores in the unit of area; therefore, it might represent the location of smaller pore clusters and compactness of the pore clusters. On the other hand, a lower value indicates a less dense area, or an area covered with larger pores. It can be observed from the kernel density contours of a sample image (Figure 2D) that the distribution of most of the larger pores are almost uniformly dispersed; however, the smaller pores formed clusters at the top-left corner of the image.

Conclusion

The algorithm provides an image analysis method for biomaterial science research to investigate the structural heterogeneity of hydrogels. This simple and flexible analysis method allows optimization of different parameters to ideally analyse a broad range of images. We have also demonstrated that based on the data extracted from the image, the kernel density estimation function is a powerful graphical tool to visualize and compare spatial heterogeneity and porosity of the hydrogel. The application of this method can be extended to structural analysis of any other porous network. Furthermore, it is worth mentioning that applying more sophisticated

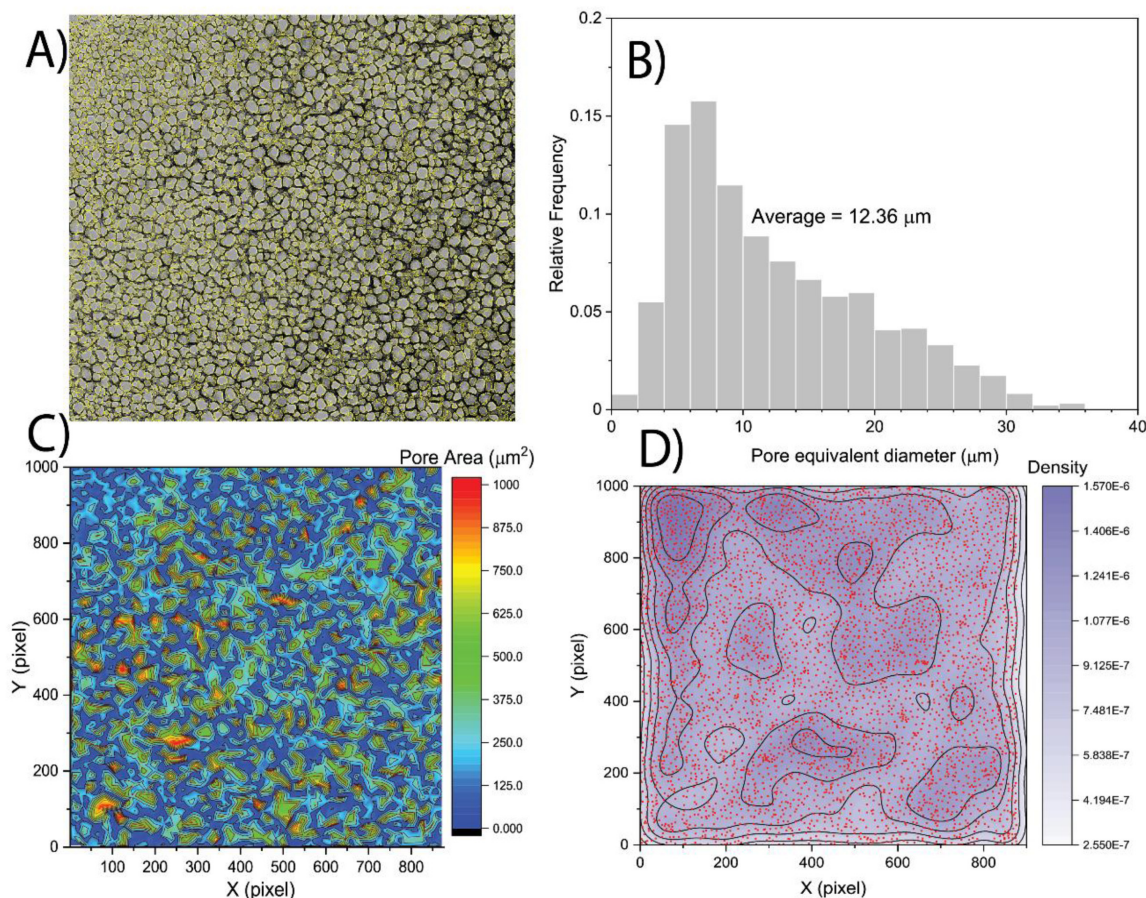


Figure 2. Porosity and heterogeneity analysis of glycidyl methacrylate hydrogel crosslinked with 0.3 mol% di(ethylene glycol) dimethacrylate. A) Detected pores of cryo-SEM micrographs. **B)** Pore size distribution. **C)** Spatial contour maps of pore area. **D)** Kernel density estimation dot plots demonstrating the spatial density of detected pores. Data adapted from Kaberova *et al.*⁸ with permission.

segmentation methods¹³ to combine classical transformation with deep learning models in the future works might improve the accuracy and performance of the method to distinguish touching pores and separate hydrogel walls.

Data availability

Zenodo: niliou/Hydrogel-pore-size: Hydrogel pore size distribution. <https://doi.org/10.5281/zenodo.4308907>¹⁴.

This project contains the following underlying data:

- Hydrogel pore size source images (original source image files in TIF format)

Data are available under the terms of the [Creative Commons Zero "No rights reserved" data waiver](https://creativecommons.org/licenses/by/4.0/) (CC0 1.0 Public domain dedication).

Code availability

Source code available from: <https://github.com/niliou/Hydrogel-pore-size.git>

Archived source code at time of publication: <https://doi.org/10.5281/zenodo.4308907>¹⁴.

License: MIT Licence

Acknowledgments

The authors would like to thank Niloufar Jamshidi (PhD candidate, UNSW) for her technical assistance particularly in developing the code. This publication was supported by COST Action NEUBIAS (CA15124), funded by COST (European Cooperation in Science and Technology).

References

1. Mantha S, Pillai S, Khayambashi P, *et al.*: **Smart Hydrogels in Tissue Engineering and Regenerative Medicine**. *Materials (Basel)*. 2019; **12**(20): 3323. [PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
2. Park KM, Lewis D, Gerecht S: **Bioinspired Hydrogels to Engineer Cancer Microenvironments**. *Annu Rev Biomed Eng*. 2017; **19**: 109–33. [PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
3. Cavo M, Fato M, Peñuela L, *et al.*: **Microenvironment complexity and matrix stiffness regulate breast cancer cell activity in a 3D *in vitro* model**. *Sci Rep*. 2016; **6**: 35367. [PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
4. d'Angelo M, Benedetti E, Tupone MG, *et al.*: **The Role of Stiffness in Cell Reprogramming: A Potential Role for Biomaterials in Inducing Tissue Regeneration**. *Cells*. 2019; **8**(9): 1036. [PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
5. Marmorat C, Arinstein A, Koifman N, *et al.*: **Cryo-Imaging of Hydrogels Supermolecular Structure**. *Sci Rep*. 2016; **6**(1): 25495. [PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
6. Kalasová D, Břínek A, Šlouf M, *et al.*: **Wide-Cone Angle Phase-Contrast X-Ray Computed Tomography of Synthetic Polymer Materials**. *IEEE Transactions on Instrumentation and Measurement*. 2020; **69**(11): 8910–8. [Publisher Full Text](#)
7. Přádný M, Dušková-Smrčková M, Dušek K, *et al.*: **Macroporous 2-hydroxyethyl methacrylate hydrogels of dual porosity for cell cultivation: morphology, swelling, permeability, and mechanical behavior**. *Journal of Polymer Research*. 2014; **21**(11): 1–12. [Publisher Full Text](#)
8. Kaberova Z, Karpushkin E, Nevoralová M, *et al.*: **Microscopic Structure of Swollen Hydrogels by Scanning Electron and Light Microscopies: Artifacts and Reality**. *Polymers (Basel)*. 2020; **12**(3): 578. [PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
9. Basu M: **Gaussian-based edge-detection methods-a survey**. *IEEE Transactions on Systems, Man, and Cybernetics, Part C (Applications and Reviews)*. 2002; **32**(3): 252–60. [Publisher Full Text](#)
10. Eng HL, Ma KK: **Noise adaptive soft-switching median filter**. *IEEE Trans Image Process*. 2001; **10**(2): 242–51. [PubMed Abstract](#) | [Publisher Full Text](#)
11. Cai Z, Ou Y, Ling Y, *et al.*: **Feature Detection and Matching With Linear Adjustment and Adaptive Thresholding**. 2020; **8**: 189735–46. [Publisher Full Text](#)
12. Bodenberger N, Kubiczek D, Abrosimova I, *et al.*: **Evaluation of methods for pore generation and their influence on physio-chemical properties of a protein based hydrogel**. *Biotechnol Rep (Amst)*. 2016; **12**: 6–12. [PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
13. Bai M, Urtasun R, editors: **Deep watershed transform for instance segmentation**. In *Proceedings of the IEEE Conference on Computer Vision and Pattern Recognition*. 2017; 5221–5229. [Reference Source](#)
14. Maryam, *et al.*: **niliou/Hydrogel-pore-size: Hydrogel pore size distribution (Version R2)**. *Zenodo*. 2020. <http://www.doi.org/10.5281/zenodo.4308907>

Open Peer Review

Current Peer Review Status:  

Version 2

Reviewer Report 02 July 2021

<https://doi.org/10.5256/f1000research.55764.r86092>

© 2021 Rocha S. This is an open access peer review report distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.



Susana Rocha 

Molecular Imaging and Photonics, KU Leuven, Leuven, Belgium

The influence of mechanical and physical cues on cell behaviour has led to an increasing interest in the quantitative analysis of structural features of hydrogels. The authors here present an algorithm to analyze 2D Cryo-SEM images of a glycidyl methacrylate hydrogel (previous published elsewhere).

The detailed explanation of the different steps of the algorithm, and the fact that this is open source, are strong points to recommend the indexing of this article. However, there are some points that should be addressed.

1. The pre-processing and thresholding steps are well explained, but there is some information missing in the 'morphological characterization' section. The authors mention: 'And lastly, the image was reconstructed based on the erosion and reopening results and then a watershed algorithm was used to segment the pores and measure pore properties.' For persons unfamiliar with image processing, a 'watershed algorithm' does not say much. Please add some more information, especially on what parameters can be calculated, how are these parameters obtained, and what is the advantage of using a watershed algorithm instead of other analysis (e.g. bubble analysis, as described in the references added in this report).
2. It is not clear how the 'result' is calculated. At the end of the section 'Morphological characterization,' the authors write: 'The obtained results of the proposed method (17 μm) were compared....'. It is not clear what the '17 μm ' refer to. In Figure 2B the average pore equivalent diameter is 12.36 μm , so how do the authors obtain the 17 μm value?
3. What do the authors mean with 'pore equivalent diameter'? This information can be added to the explanation of the watershed algorithm.
4. Can you provide more information on the kernel density estimation function? How does it work and which information can you obtain? It is not clear in the text why you use this function.

5. On figure 1 it is difficult to judge the performance of the different image processing algorithms. Would it be possible to add a magnification? If it is not possible to add it for all the steps, at least a magnified view of the original image and the water-shed segmentation should be added (panels A and I).
6. In figure 2C the axes are in pixels and the pore area is given in μm^2 . Please change the pixels to μm .
7. General remark: there are no scale bars in the images.

References

1. Mickel W, Münster S, Jawerth LM, Vader DA, et al.: Robust pore size analysis of filamentous networks from three-dimensional confocal microscopy. *Biophys J.* 2008; **95** (12): 6072-80 [PubMed Abstract](#) | [Publisher Full Text](#)
2. Molteni M, Magatti D, Cardinali B, Rocco M, et al.: Fast two-dimensional bubble analysis of biopolymer filamentous networks pore size from confocal microscopy thin data stacks. *Biophys J.* 2013; **104** (5): 1160-9 [PubMed Abstract](#) | [Publisher Full Text](#)
3. Münster S, Fabry B: A simplified implementation of the bubble analysis of biopolymer network pores. *Biophys J.* 2013; **104** (12): 2774-5 [PubMed Abstract](#) | [Publisher Full Text](#)
4. Molteni M, Magatti D, Cardinali B, Rocco M, et al.: Response to. *Biophys J.* 2013; **104** (12): 2776-7 [PubMed Abstract](#) | [Publisher Full Text](#)

Is the work clearly and accurately presented and does it cite the current literature?

Partly

Is the study design appropriate and is the work technically sound?

Yes

Are sufficient details of methods and analysis provided to allow replication by others?

Yes

If applicable, is the statistical analysis and its interpretation appropriate?

Partly

Are all the source data underlying the results available to ensure full reproducibility?

Yes

Are the conclusions drawn adequately supported by the results?

Yes

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: (fluorescence) microscopy, biomaterials, nanotechnology

I confirm that I have read this submission and believe that I have an appropriate level of

expertise to confirm that it is of an acceptable scientific standard.

Reviewer Report 20 April 2021

<https://doi.org/10.5256/f1000research.55764.r83172>

© 2021 Dušková-Smrčková M. This is an open access peer review report distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.



Miroslava Dušková-Smrčková 

Institute of Macromolecular Chemistry, Academy of Sciences of the Czech Republic, Prague, Czech Republic

I have reviewed the authors' answers to my points and I liked them and I agree with the modifications/explanations they added. I have no more points to be addressed and recommend the paper for indexing in its current state.

Is the work clearly and accurately presented and does it cite the current literature?

Yes

Is the study design appropriate and is the work technically sound?

Yes

Are sufficient details of methods and analysis provided to allow replication by others?

Yes

If applicable, is the statistical analysis and its interpretation appropriate?

Yes

Are all the source data underlying the results available to ensure full reproducibility?

Yes

Are the conclusions drawn adequately supported by the results?

Yes

Competing Interests: No competing interests were disclosed.

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

Version 1

Reviewer Report 02 March 2021

<https://doi.org/10.5256/f1000research.30250.r79780>

© 2021 Dušková-Smrčková M. This is an open access peer review report distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.



Miroslava Dušková-Smrčková 

Institute of Macromolecular Chemistry, Academy of Sciences of the Czech Republic, Prague, Czech Republic

It brings a useful piece of computer image analyses software development applicable to quantification of images of heterogeneous materials, namely of 2D projection of pores/distinguishable distributed objects – such as bubbles, particles etc.

First, a brief generic remark. Presently, a number of image analysis software packages for materials science, commercial as well as open-source, for complex scientific image processing are available. However, it is an increasing challenge for the end user to understand the routine behind the image processing and 2D or 3D structure rendering and sometimes the description of the certain function under “Help” does not offer much possibility to grasp the principle.

The presented work is particularly concentrated on the analysis of microscopy images of macroporous hydrogels where the advancement of image processing such as the proposed method is still needed: the reason is that we need to investigate relations between hydrogel morphology, microstructure and hydrogel macroscopic properties, mainly mechanical behavior and its effects in biological application – because not everything is known and there are new types of hydrogels of plethora structures being introduced by chemists.

On top of that, besides the scientific paper containing detailed explanation of itself, Jamshidi and Falamaki made their script (written in Python) available to scientific community – that I found really useful. With more users, we can expect evaluation of high number of samples and possibly building up the software package for more versatile gel characterization. Because of detailed explanation of the hydrogel image processing, the work offers possibility to other interested researchers on the border areas between hydrogel (materials) design and image processing to learn and extend the work. Therefore, I consider publishing this paper with F1000 Research valuable.

Before indexing the manuscript, I still propose some modification and clarification of few points of discussion – mainly to make the paper speaking to hydrogel designers and chemists who are most of the time users or advanced users but do not have the ambition to be software developers.

1. A very important point to clarify is the very first sentence of the Abstract. The authors say: “This work presents an image processing procedure for characterization of porosity and heterogeneity of fully hydrated hydrogels based on the analysis of cryogenic scanning electron microscopy (cryo-SEM) images.” Let’s make this clear: analysis of cryo-SEM images may but may not reveal the structure as in hydrated state. When a fully hydrated hydrogel sample is processed for cryogenic EM, it goes through an inevitable freezing step. The morphology of hydrated sample most likely will be altered depending on the conditions of the freezing step but also depending on the particular sample. Freezing, whatever rapid, will cause change in volume and shape of existing pores¹, possibly will cause formation of

new pores, or will cause some failure, cracks, distortion of the gel phase. Or it will just cause isotropic shrinking of the hydrogel volume. We can't exactly predict the change. This means that whatever structure of gel the cryo-SEM reveals, it may not correspond to the structure in fully hydrated – meaning swollen in water to equilibrium under given conditions – hydrogel. Only in rather special cases, for example in less swollen and mechanically stronger gels, the gel matrix can resist changes, or in the case of macroscopic gels with arrangement of connected spheres (see Fig. 4 of paper by Kalasova et al in DOI 10.1109/TIM.2020.2995232²) it can just isotropically contract (as shown in already cited work (<https://doi.org/10.3390/polym12030578>)³). This all said in the above paragraph does not disrate the impact of the developed image processing – as it can be well applied to images obtained by other methods, too. This point indeed, should be taken into account in the introductory part – pg. 3, where the text again reads: “This article presents an image analysis method to characterise hydrogel structure heterogeneity and porosity based on the cryogenic scanning electron microscopy (cryo-SEM) images of fully hydrated hydrogels.” I propose to modify the formulation similarly as: “This article presents an image analysis method to characterise hydrogel structure heterogeneity and porosity resulting from the treatment of fully hydrated hydrogels – plunge-freezing of hydrated samples done to acquire their cryogenic scanning electron microscopy (cryo-SEM) images.”

2. Several times authors speak about “hydrogel fibers” – I wish to bring up that the form of hydrogel is not fibrous, the gel forms rather thin walls separating the pores in 3D space.
3. There are background and foreground pores mentioned – pls add some schematics what that exactly means and where the reader can see the background vs. the foreground pores on the presented figures.
4. What does the “OTSU” abbreviation stand for and what is OTSU threshold mentioned in part “Morphological transformation”? (abbreviation not explained)
5. The reference where from the Figure 1 was adopted should be given also in the Figure 1 legend.
6. The size of the figures and their resolution even in supplied pptx files is both quite low. Especially the small size of printed letters in the images is critical and must be increased (and improved resolution): legend, axis descriptions, numbers within the Figure 2 are not readable. There are two images both designated Fig.1/ B.
7. The actual results shown on Figure 2 – mean pore size, pore size distribution ...is not discussed in numbers. It should also be compared with the original evaluation given in the source article. To determine the size of pores should be the target of the presented method.
8. It would be of interest to discuss the existing limits of the presented method: for example how the method will tackle anisotropic pores, either regularly distributed or clustered or even oriented? – Will the method detect connecting pores, etc? Indeed, this can be the future direction but even then will be worthwhile mentioning.
9. It would be of interest to adopt the method to evaluate porosity: the void volume fraction in the sample.

10. The thickness of the gel wall separating pores is also of interest and should be also analysed by IA package.

Last but not least my comment of Conclusion:

The material depicted on Figures 1 and 2 should not be called fibrous network – the pores in it look like those in the sponge for washing dishes: pores surrounded by thinner or thicker hydrogel walls of thickness typically 10^0 - 10^1 μm . Indeed, the molecular basis of this material is macromolecular network, that means that all building units molecules are connected to one molecule of a 3D macromolecular network spanning the whole sample space but this molecular nature of the samples indeed is far below resolution of currently available visualization methods.

References

1. Podhorská B, Vetrík M, Chylíková-Krumbholcová E, Kománková L, et al.: Revealing the True Morphological Structure of Macroporous Soft Hydrogels for Tissue Engineering. *Applied Sciences*. 2020; **10** (19). [Publisher Full Text](#)
2. Kalasova D, Brinek A, Slouf M, Duskova-Smrckova M, et al.: Wide-Cone Angle Phase-Contrast X-Ray Computed Tomography of Synthetic Polymer Materials. *IEEE Transactions on Instrumentation and Measurement*. 2020; **69** (11): 8910-8918 [Publisher Full Text](#)
3. Kaberova Z, Karpushkin E, Nevoralová M, Vetrík M, et al.: Microscopic Structure of Swollen Hydrogels by Scanning Electron and Light Microscopies: Artifacts and Reality. *Polymers (Basel)*. 2020; **12** (3). [PubMed Abstract](#) | [Publisher Full Text](#)

Is the work clearly and accurately presented and does it cite the current literature?

Yes

Is the study design appropriate and is the work technically sound?

Yes

Are sufficient details of methods and analysis provided to allow replication by others?

Yes

If applicable, is the statistical analysis and its interpretation appropriate?

Partly

Are all the source data underlying the results available to ensure full reproducibility?

Yes

Are the conclusions drawn adequately supported by the results?

Yes

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: macromolecular chemistry

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard, however I have significant reservations, as outlined above.

Author Response 29 Mar 2021

Cavus Falamaki, Amirkabir University of Technology, P.O. Box 15875-4413, Tehran, Iran

Author's Respond to Reviewer:

Authors would like to sincerely thank reviewer not only for their constructive comments and scientific clarification that improves the scientific quality of the paper, but also for their suggestions that can be used to further develop the current piece of work.

All modifications made in the manuscript are shown with **.

Reply to reviewers' comments are shown with #.

It brings a useful piece of computer image analyses software development applicable to quantification of images of heterogeneous materials, namely of 2D projection of pores/distinguishable distributed objects – such as bubbles, particles etc.

First, a brief generic remark. Presently, a number of image analysis software packages for materials science, commercial as well as open-source, for complex scientific image processing are available. However, it is an increasing challenge for the end user to understand the routine behind the image processing and 2D or 3D structure rendering and sometimes the description of the certain function under “Help” does not offer much possibility to grasp the principle.

The presented work is particularly concentrated on the analysis of microscopy images of macroporous hydrogels where the advancement of image processing such as the proposed method is still needed: the reason is that we need to investigate relations between hydrogel morphology, microstructure and hydrogel macroscopic properties, mainly mechanical behavior and its effects in biological application – because not everything is known and there are new types of hydrogels of plethora structures being introduced by chemists.

On top of that, besides the scientific paper containing detailed explanation of itself, Jamshidi and Falamaki made their script (written in Python) available to scientific community – that I found really useful. With more users, we can expect evaluation of high number of samples and possibly building up the software package for more versatile gel characterization. Because of detailed explanation of the hydrogel image processing, the work offers possibility to other interested researchers on the border areas between hydrogel (materials) design and image processing to learn and extend the work. Therefore, I consider publishing this paper with F1000 Research valuable.

Before indexing the manuscript, I still propose some modification and clarification of few points of discussion – mainly to make the paper speaking to hydrogel designers and chemists who are most of the time users or advanced users but do not have the ambition to be software developers.

- A very important point to clarify is the very first sentence of the Abstract. The authors say: “This work presents an image processing procedure for characterization of porosity and heterogeneity of fully hydrated hydrogels based on the analysis of cryogenic scanning electron microscopy (cryo-SEM) images.” Let's make this clear:

analysis of cryo-SEM images may but may not reveal the structure as in hydrated state. When a fully hydrated hydrogel sample is processed for cryogenic EM, it goes through an inevitable freezing step. The morphology of hydrated sample most likely will be altered depending on the conditions of the freezing step but also depending on the particular sample. Freezing, whatever rapid, will cause change in volume and shape of existing pores¹, possibly will cause formation of new pores, or will cause some failure, cracks, distortion of the gel phase. Or it will just cause isotropic shrinking of the hydrogel volume. We can't exactly predict the change. This means that whatever structure of gel the cryo-SEM reveals, it may not correspond to the structure in fully hydrated – meaning swollen in water to equilibrium under given conditions – hydrogel. Only in rather special cases, for example in less swollen and mechanically stronger gels, the gel matrix can resist changes, or in the case of macroscopic gels with arrangement of connected spheres (see Fig. 4 of paper by Kalasova et al in DOI 10.1109/TIM.2020.2995232²) it can just isotropically contract (as shown in already cited work (<https://doi.org/10.3390/polym120305783>)). This all said in the above paragraph does not disrate the impact of the developed image processing – as it can be well applied to images obtained by other methods, too. This point indeed, should be taken into account in the introductory part – pg. 3, where the text again reads: “This article presents an image analysis method to characterise hydrogel structure heterogeneity and porosity based on the cryogenic scanning electron microscopy (cryo-SEM) images of fully hydrated hydrogels.” I propose to modify the formulation similarly as: “This article presents an image analysis method to characterise hydrogel structure heterogeneity and porosity resulting from the treatment of fully hydrated hydrogels – plunge-freezing of hydrated samples done to acquire their cryogenic scanning electron microscopy (cryo-SEM) images.”

The authors agree and admit that in the paper it should be clarified that there are some artifacts associated with the imaging methods specially cryo-SEM and proper referencing should be made.

The abstract is revised accordingly, and some clarifications is added to the introduction part to cover the reviewer's points:

**This work presents an image processing procedure for characterization of porosity and heterogeneity of hydrogels network mainly based on the analysis of cryogenic scanning electron microscopy (cryo-SEM) images and can be extended to any other type of microscopy images of hydrogel porous network.

#The following paragraph is modified in the introduction part:

**This article presents an image analysis method applied to characterise hydrogel structure heterogeneity and porosity resulting from the treatment of fully hydrated hydrogels during plunge-freezing to acquire their cryogenic scanning electron microscopy (cryo-SEM) images. It should be mentioned that during the process of sample preparation for cryo-SEM microscopy, the sample undergoes a probable morphological alteration, furthermore the process itself might introduce artifacts according to the nature of the material and swelling rate. Readers are encouraged to refer to the studies specialized on the imaging of different hydrogel network by Kalasova et. al¹ and Pradny et. al².

- Several times authors speak about “hydrogel fibers” – I wish to bring up that the form of hydrogel is not fibrous, the gel forms rather thin walls separating the pores in 3D space.

Authors acknowledge the point, in the manuscript wherever the term “fibre” is used, replaced with “hydrogel walls”.

- There are background and foreground pores mentioned – pls add some schematics what that exactly means and where the reader can see the background vs. the foreground pores on the presented figures.

Authors agree that there is a need to clarify the terms used to explain the method. The image of the hydrogel is first segmented into two categories of the hydrogel walls and pores. Because in this analysis we aim to detect the pores, therefore the pores segment is foreground (value 1 – white), and the walls are considered as background (value 0- black). Accordingly, the figure 1 has been edited and a small schematic showing foreground and background is added to the figure 1.

Following section has been corrected also for more clarification in “Thresholding” section and also a schematic is added to Figure 1:

**Since in most of the cryo-SEM images there is uneven illumination, the adaptive threshold technique was undertaken to segment the background (hydrogel walls, value=0 black) and foreground pores (pores, value=1 white).

**Figure 1. Image processing algorithm of sample cryo-SEM micrograph of glycidyl methacrylate hydrogel crosslinked with 0.3 mol% (ethylene glycol) dimethacrylate. Images on the left show the output images of each step in the processing algorithm and the relevant flowchart is illustrated on the right to distinguish foreground (pores) from background (hydrogel walls). The source image adapted from Kaberova *et al.*³ with permission.

- What does the “OTSU” abbreviation stand for and what is OTSU threshold mentioned in part “Morphological transformation”? (abbreviation not explained)

There was a typo here, Otsu’s method is one of the methods used for image binarization or thresholding. The following sentence in under “Morphological transformation” is edited:

**To validate the method, the screenshot images of a hydrogel network from a previously published work where the corresponding pore diameters have been reported, were analysed with the proposed method using Otsu’s thresholding method and setting 0.05 max watershed threshold.

in the corresponding python script under thresholding section, both methods of binarization (Otsu’s and Adaptive are provided), user can comment each line and use what is best to process their image. We have used Otsu’s for validation, and adaptive method is suggested for cryo-EM images.

- The reference where from the Figure 1 was adopted should be given also in the Figure 1 legend.

The comment has been considered and the legend for figure 1 is revised as bellow:

*Figure 1. Image processing algorithm of sample cryo-SEM micrograph of glycidyl methacrylate hydrogel crosslinked with 0.3 mol% (ethylene glycol) dimethacrylate. Images on the left show the output images of each step in the processing algorithm and the relevant flowchart is illustrated on the right to distinguish foreground (pores) from background (hydrogel walls). The source image adapted from Kaberova *et al.*³ with permission.

- The size of the figures and their resolution even in supplied pptx files is both quite

low. Especially the small size of printed letters in the images is critical and must be increased (and improved resolution): legend, axis descriptions, numbers within the Figure 2 are not readable. There are two images both designated Fig.1/ B.

#The comment considered, the resolution of the figures increased and high-quality images in actual sizes has been provided in power point files as attachment.

- The actual results shown on Figure 2 – mean pore size, pore size distribution ...is not discussed in numbers. It should also be compared with the original evaluation given in the source article. To determine the size of pores should be the target of the presented method.

Authors consider the comment; however, it should be noted that for this paper we have analysed only one image (Figure 1. e) of the source article by Kaberova et. al³. This image is related to sample G0/0.3 in the source article which the pore size is reported between 2-40 μm (we assume that these values are based on equivalent diameter) for this sample according to Table 1. Using the current method, the average equivalent diameter of the detected pores is equal to 9.45 pixels or 12.36 μm (pixel to microns scale of the image = 500 μm / 382 pixel). The actual pore size distribution is updated in Figure 2, from this image it is apparent that the measured pore size distribution by this method sit between 2 – 36 μm .

the correction applied to the Results section of the manuscript:

* Moreover, the average equivalent diameter of the pores of the analysed source image (adopted from Kaberova et. al³) was equal to 12.36 μm with the corresponding pore size distribution range of 2-36 μm . The comparison of the results with the range reported by Kaberova et al.³ (2–40 μm) shows a good agreement and reliability of the presented method as it is shown in Figure 2B.

- It would be of interest to discuss the existing limits of the presented method: for example how the method will tackle anisotropic pores, either regularly distributed or clustered or even oriented? – Will the method detect connecting pores, etc? Indeed, this can be the future direction but even then will be worthwhile mentioning.

This is a very interesting point picked up by reviewer. Authors clarify that the current method does not classify the hydrogel network as “clustered” or “distributed” and we have not done any statistical analysis to focus on quantification of anisotropic properties of the hydrogel network. However, structural and cluster analysis of the gel network is the topic of the next paper currently developing by one of the team members. The result and relevant script of the next work will be available to scientific community after publishing.

For the detection of the connecting pores, we have applied watershed transform which is a well-established method to separate touching objects, however authors admit that there are more advanced and complex methods/ pipelines to segment the pores or hydrogel walls. The might be of interest to apply a recently developed methods that combines the classic “watershed” segmentation with “deep learning”⁴ to increase the level of accuracy and performance of the segmentation method of the analysis package.

The following sentences under Conclusion part is revised in the manuscript:

*The application of this method can be extended to structural analysis of any other porous network. Furthermore, it is worth mentioning that applying more sophisticated segmentation methods⁴ to combine classical transformation with deep learning models in the future works might improve the accuracy and performance of the method to distinguish touching pores and separate hydrogel walls. It would be of interest to adopt the method to

evaluate porosity: the void volume fraction in the sample.

Authors acknowledge the reviewer's point. The void fraction calculation has appended to the script and it is printed in the console when the script run has ended (open images should be closed). We consider 2D void fraction = total areal of pores / image area.

Script new version (R2 is uploaded on github accordingly).

[https://github.com/niliou/Hydrogel-pore-](https://github.com/niliou/Hydrogel-pore-size/blob/main/Hydrogel%20Pore%20Size%20Distribution%20-%20R2%20Final.py)

[size/blob/main/Hydrogel%20Pore%20Size%20Distribution%20-%20R2%20Final.py](https://github.com/niliou/Hydrogel-pore-size/blob/main/Hydrogel%20Pore%20Size%20Distribution%20-%20R2%20Final.py)

- The thickness of the gel wall separating pores is also of interest and should be also analysed by IA package.

We admit the comment, the present method can also be used to measure wall thickness of the gel. The only change to be made is the background and fore ground that should be reversed (fore ground is of interest = white) instead of pores we need to bring walls to the front or white. Another point is that for thresholding some values should be tweaked that is commented in the corresponding section in the script. Therefore, by inverting the image look up table the wall thickness can be measured. For the source image we have:

should be noted that in the results file the minimum axis value should be considered as closet parameter describing hydrogel walls. The value of the average wall thickness for the source image is equal to 4.79 pixels or 6.26 μm .

The following clarification has been made to the Image processing section:

*To detect and measure the thickness of the hydrogel walls, the look up table of the binary image should be inversed to bring hydrogel walls in foreground (white).

#The relevant hints are also commented in the script.

#To invert the look-up table of the image in the first part, Then, line 130 should be uncommented and used for hydrogel wall thickness measurement, the input image in the command line (130) could be any of the filtered images that we have included in the script, so the end user can choose the best filter that gives the best results, in the above we have chosen "DoGNN" image which is filtered image by Difference of Gaussian method. In the script we have several filters (lines 45-84) + bandpass filter lines (103-104) that mainly focuses on edge feature detection. We have commented in the script describing what filters are applied in each section. So, user can choose which one of these filters can extract the wall features the best. Line 144 also can be commented in case "hydrogel wall thickness" is being measured.

Last but not least my comment of Conclusion:

The material depicted on Figures 1 and 2 should not be called fibrous network – the pores in it look like those in the sponge for washing dishes: pores surrounded by thinner or thicker hydrogel walls of thickness typically 100-101 μm . Indeed, the molecular basis of this material is macromolecular network, that means that all building units molecules are connected to one molecule of a 3D macromolecular network spanning the whole sample space but this molecular nature of the samples indeed is far below resolution of currently available visualization methods.

Authors appreciate reviewer's attention and their explanation to improve the scientific value of the work as well as authors understanding of the nature of hydrogel network.

Anywhere in the manuscript the term "fibrous" is revised and replaced.

1. Kalasová D, Břínek A, Šlouf M, Dušková-Smrčková M, Zikmund T, Patáková Z, et al. Wide-Cone Angle Phase-Contrast X-Ray Computed Tomography of Synthetic Polymer Materials. *IEEE Transactions on Instrumentation and Measurement*. 2020; **69**(11):8910-8.
2. Přádny M, Dušková-Smrčková M, Dušek K, Janoušková O, Sadakbayeva Z, Šlouf M, et al. Macroporous 2-hydroxyethyl methacrylate hydrogels of dual porosity for cell cultivation: morphology, swelling, permeability, and mechanical behavior. 2014; **21**(11):1-12.
3. Kaberova Z, Karpushkin E, Nevalová M, Vetrík M, Šlouf M, Dušková-Smrčková M. Microscopic Structure of Swollen Hydrogels by Scanning Electron and Light Microscopies: Artifacts and Reality. *Polymers*. 2020; **12**(3):578.
4. Bai M, Urtasun R, editors. Deep watershed transform for instance segmentation. 2017.

Competing Interests: No competing interests were disclosed.

The benefits of publishing with F1000Research:

- Your article is published within days, with no editorial bias
- You can publish traditional articles, null/negative results, case reports, data notes and more
- The peer review process is transparent and collaborative
- Your article is indexed in PubMed after passing peer review
- Dedicated customer support at every stage

For pre-submission enquiries, contact research@f1000.com

F1000Research