- 1 Title: Multi-scale Pore-network Representation of Heterogeneous Carbonate Rocks Tannaz Pak $^{1,\,2,\,\star}$ , Ian B. Butler $^{1,\,2}$ , Sebastian Geiger $^{2,\,3}$ , Marinus I.J. van Dijke $^{2,\,3}$ , Zeyun Jiang $^{2,\,3}$ , Rodrigo Surmas $^{2,\,3}$ 2
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- 13 Key points:
- 14 This work investigates the multi-scale porosity of a carbonate rock.
- 15 We explore representativeness of multi-scale networks by comparing simulated and 16 laboratory measured capillary-pressure vs saturation curves.
- 17 We present a workflow for selecting number and length scales of networks to 18 generate multi-scale pore-networks.

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## Abstract:

A multi-scale network integration approach introduced by Jiang et al. [2013] is used to generate a representative pore-network for a carbonate rock with a pore-size distribution across several orders of magnitude. We predict the macroscopic flow parameters of the rock utilising i) 3D images captured by X-ray computed micro-tomography and ii) pore-network flow simulations. To capture the multi-scale pore-size distribution of the rock we imaged four different rock samples at different resolutions and integrated the data to produce a porenetwork model that combines information at several length-scales that cannot be recovered from a single tomographic image. A workflow for selection of the number and length-scale of the required input networks for the network integration process, as well as fine tuning the model parameters is presented. Mercury injection capillary-pressure data were used to evaluate independently the multi-scale networks. We explore single-scale, two-scale, and three-scale network models and discuss their representativeness by comparing simulated capillary-pressure versus saturation curves with laboratory measurements. We demonstrate that for carbonate rocks with wide pore-size distributions, it may be required to integrate networks extracted from two or three discrete tomographic data sets in order to simulate macroscopic flow parameters.

## 1. Introduction

Pore-scale fluid flow simulation is a useful tool for calculation of the macroscopic transport properties of porous media, specifically for homogeneous systems [*Meakin and Tartakovsky*, 2009; *Joekar-Niasar and Hassanizadeh*, 2012; *Blunt et al.*, 2013; *Wildenschild and Sheppard*, 2013]. For complex porous material realistic representation of the pore structure is essential for successful simulations but has proven to be challenging. High-resolution X-ray computed micro-tomography (μCT) imaging allows detailed study of porous media in 3D (reviews by *Wildenschild and Sheppard* [2013], *Fusseis et al.* [2014], and *Bultreys et al.* [2016]). Reliable application of this technique requires sufficient image quality to allow discrimination of the important features of the sample. Pore-network models (PNM) extracted from binarised μCT images (see review by *Dong* [2007]) are simple representations of pore

structures to simulate flow at a reduced computational cost. Although appealing, PNM predictions often do not agree well with the corresponding laboratory measurements, specifically for complex pore systems such as carbonate rocks [Sorbie and Skauge, 2012]. The trade-off between the sample size and μCT image resolution means that a single μCT volume may not capture the details of a multi-scale pore structure [Hebert et al., 2014]. Capturing a representative image of heterogeneous samples with pore-size distribution (PSD) spanning several orders of magnitude is an outstanding challenge with existing µCT instrumentation. A representative elementary volume (REV) is required that is the smallest sample to which measured properties of the bulk sample can be assigned [Bear, 1972]. Generating two-scale PNM that are representative of complex pore systems has been studied by Biswal et al. [2007]; Biswal et al. [2009] and Ghous et al. [2008], and more recently by Prodanović et al. [2015], Mehmani and Prodanović [2014], and Bultreys et al. [2015]. Jiang et al. [2013] introduced a network integration method to generate PNMs of arbitrarily large volumes that can incorporate multi-scale pore systems extracted [Jiang et al., 2007] from images captured at different resolutions. A detailed PNM extracted from a high resolution µCT volume (fine-network) is combined with a network extracted from a coarse resolution volume (coarse-network). For details of how the nodes/bonds of the differently scaled models are interconnected see Jiang et al. [2013]. The domain of the fine-network is smaller than that of the coarse-network, hence a larger network that is statistically equivalent to the fine-network is generated and then integrated into the coarse-network. The generated multi-scale networks can be heavier than the original fine-network, with an adverse effect on the computational cost of fluid flow simulations. To address this *Jiang et al.* [2013], reduced the number of fine-network elements by a fraction  $(f_F)$  such that  $0 \le f_F \le 1$ . It is also possible to scale the size of the domain of the two-scale network (nesting-domain) with respect to that of

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- 74 the original coarse image using a scaling factor ( $\delta$ ) that is arbitrary but is restricted by the
- 75 available computational power for flow simulations.
- We investigate generation of a representative PNM for a dolomite using *Jiang et al.* [2013]'s
- 77 approach and a quasi-static pore-network flow simulator [Ryazanov et al. [2009]. We
- evaluate Jiang et al. [2013]'s approach by comparing simulated  $P_c(S)$  curves with those
- 79 measured experimentally using mercury injection capillary-pressure (MICP) tests. A
- 80 workflow is proposed for multi-scale network generation to address (i) the choice of sample
- size and, hence, image resolution, and (ii) the tuning of model parameters to reproduce the
- laboratory  $P_c(S)$  curves.

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## 2. Imaging and Mercury Porosimetry

- 84 A sucrosic dolomite (Silurian Dolomite (SD), Thornton formation, Chicago, Illinois, US)
- 85 with pore sizes spanning 3 orders of magnitude was used (Figure 1). Laboratory
- measurements of a 38mm diameter core displayed porosity and permeability of 17% and
- 87 ∼50mD, respectively (Figure 1b).
- We imaged, using  $\mu$ CT, four samples of diameter D=2, 5, 25 and 38mm, at resolutions of 2.3,
- 89 12, 21 and 32.8µm, respectively (Figure 1b). Both the image voxel sizes and the corrected
- 90 image resolutions are reported in Table 1, the correction accounts for the unsharpness caused
- by the X-ray source spot size [Feser et al., 2008]. The values presented throughout the rest of
- 92 the manuscript are the corrected image resolution values. As currently worded, this is still a
- 93 bit unclear. Methods are detailed in the SI and Pak et al. [2013]. Sizes were selected because
- 94 38mm is a standard petrophysical plug, 25mm is the largest sample the MICP could analyse,
- and 5mm and 2mm provide details of the finer porosity. Figure 1c displays the PSD derived
- 96 from these four data volumes (Figure 1b) using a sphere fitting method [Jiang et al., 2007].
- 97 Each volume has captured only a portion of the PSD. In SD recrystallization to dolomite has
- 98 produced a structure in which the pores are lined by the apices and edges of dolomite rhombs

with well sorted crystals sizes. Microscope images (Figure S1.) show that smaller pores are formed in between a few crystals, are uniformly present throughout the sample, suggesting little or no spatial correlation of the micro-porosity. Hence, we uniformly integrate the fine-network in the entire nesting-domain.

MICP measurement was performed on three 25 mm diameter SD core plugs imaged at resolution  $21\mu m$  (Figure S2). MICP provides  $P_c(S)$  curves from which pore-throat size distribution can be extracted [*Ritter and Drake*, 1945]. The MICP-based porosity for these samples is 18.03, 16.03 and 18.1%. The image-based porosities at this resolution are 8.05, 6.03 and 8.12%, respectively, hence the  $21\mu m$  represent a coarse-scale resolution.

## 3. Multi-scale PNM Reconstruction

We explore representing the SD pore structure using multi-scale networks to incorporate features extracted from the four  $\mu$ CT volumes shown in Figure 1b. For 3D renderings of these networks see Figure S3. The physical size of these networks can be found from the image dimensions and voxel resolutions (Table 1). The two coarse-scale networks are globally disconnected while the fine-scale networks are connected with simulated absolute permeability  $k_{abs}$ >0mD. The total (and connected) porosity shows an increasing trend with resolution enhancement, as does the calculated permeability. The two fine-networks display calculated permeability values higher than the laboratory measurement i.e. 50 mD. The small samples do not contain a REV and the calculated permeability values reflect the local connectivity of the pore system at these scales. However, even for the 2.3 $\mu$ m resolution network about 11% of the pore volume is disconnected, reflecting that the resolution is insufficient to capture the smallest pore-throats, or that isolated pores exist in the sample.

## 3.1. Two-scale PNMs

Six possible combinations for generating two-scale networks are 32.8-21μm, 32.8-12μm, 32.8-2.3μm, 21-12μm, 21-2.3μm, and 12-2.3μm, referring to image resolution. Essentially, we need (i) a sufficiently large coarse-network that represents the largest pores, (ii) a fine-network with sufficient resolution that captures the smaller pores, and (iii) a sufficient overlap of the PSD of the two networks to ensure representation of the intermediate-size pores. Hence, 12-2.3μm and 32.8-21μm combinations are not acceptable. In what follows other combination are discussed.

#### 3.1.1. Combination of 21 and 2.3µm PNMs

Figure 2a shows ten networks extracted from sub-domains ( $\delta$ =0.1-1) of the 21µm coarsenetwork. At  $\delta$ =0.4 the domain is sufficiently large to accommodate a cluster of large pores,

hence the network porosity displays a sharp increase, Figure 2b. At  $\delta$ =0.5 the domain accommodates a connected cluster of pores, therefore,  $k_{abs} > 0$ . However, as the sub-domain grows ( $\delta > 0.5$ ) the networks become again disconnected, suggesting that only a locally connected cluster is captured at  $\delta$ =0.5 which is not representative of the connectivity of larger pores at this scale. Figure 2c displays the sensitivity of these two-scale networks to  $f_F$ . Porosity increases linearly with the inclusion of increasing fractions of the fine-network, while the permeability starts to converge for  $f_F > 0.05$ . Thus even a reduced version of a fine-network can provide the required connectivity to two-scale network. The calculated porosity and permeability display convergence at  $\delta > 0.6$  (i.e. total porosity ~6% and  $k_{abs}$ =0). Figure 2d, however, shows that the  $P_c(S)$  curves, calculated using these two-scale PNMs ( $f_F$ =0.01, 0.6< $\delta$ <1), do not converge. The curves display jumps at different saturations depending on the nesting-domain size. Examination of Figure 2a reveals that the coarse-network contains two connected clusters of elements (pores and nodes) that are disconnected from each other. Network integration provides connectivity between these clusters through elements of the fine-network. Cluster 1 is connected to the inlet and can be invaded by the injected phase directly. Cluster 2 is only accessible through cluster 1 using the fine-network connections. The jump occurs at saturation  $S^*$  that is equal to the ratio of the volume of cluster 1 to the total volume of the connected pore space. As  $\delta$  increases, cluster 2 grows more significantly compared to cluster 1, hence  $S^*$  approaches 0.8. Hence, the selected domain from the 21 µm resolution image does not contain a REV. However, the MICP core plugs were approximately 60% larger than the box selected for pore-network simulations, and the MICP-driven  $P_c(S)$  curves for three 25mm diameter SD plugs are very close (Figure S2a), this suggests that the MICP plugs themselves are representative for this rock.

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#### 3.1.2. Combination of 32.8 and 2.3 µm PNMs

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For this combination the two-scale network porosity converges for  $\delta > 0.6$  (Figure 3a) while  $k_{abs}$ =0 for all sub-domains. Further, the simulated  $P_c(S)$  curves for this combination ( $f_F$ =0.01) also converge for  $\delta$ >0.6 (Figure 3b). This indicates that sub-domains of the 38.2µm image with  $\delta > 0.6$  contain a REV of SD. The  $P_c(S)$  curves, however, show a jump at  $S_{air} \sim 0.8$  causing an overestimation of the experimental capillary-pressure values for  $S_{air}$ <0.8. The PSD of the 2.3μm network (Figure 1b) shows a peak at ~6μm, while pores larger than 10μm display less than 20% probability (i.e. volume fraction). Since the coarse-network is globally disconnected, the two-scale networks are connected only through fine-network elements. The initial displacement ( $S_{air} > 0.8$ ) corresponds to the pores accessible from the inlet. At  $S_{air} \sim 0.8$ the pressure needs to increase sufficiently before elements from the fine-network can be invaded. For the converged set of  $P_c(S)$  curves, on Figure 3b, the  $f_F$  is equal to 0.01. Including a larger  $f_F$  suppresses the jump in the capillary-pressure curve but does not totally remove it. The  $P_c(S)$  curve for the  $\delta$ =0.6 and  $f_F$ =0.04 network shows a small jump at  $S_{air}$ =0.8 from  $\sim$ 21kP to  $\sim$ 47kP corresponding to invasion of pore-throats with radius of 40µm to 16µm. However, this size range is insufficiently represented in both 32.8 and 2.3µm networks. In particular, a small  $f_F$  makes a reduced network less representative of the two ends of the PSD. In practice, the difference in the domain sizes of the coarse and fine-networks causes higher  $f_F$  values to make the two-scale network substantially more computationally expensive.

# 3.1.3. Combination of 32.8 and 12µm PNMs

For this combination the calculated  $P_c(S)$  curves show a good match with the laboratory measurements at  $S_{air}>0.04$  (Figure 4). However, the simulations stop at capillary-pressure values around 276kP, due to fine-network resolution limitations. Using a higher resolution

fine-network may not provide significant additional connectivity, but may prove essential for calculation of saturation values.

## 3.2.Three-scale PNMs

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We have demonstrated that a finer length-scale needs to be integrated into the 32.8–12µm combination, or an intermediate length-scale must be added to the 32.8–2.3µm combination. To integrate three PNMs of different length-scales first the two smaller networks are integrated into a two-scale network that is subsequently integrated into the largest scale network. This involves selecting two sets of  $\delta$  and  $f_F$ . As the number of length-scales increases, sensitivity analysis on the effect of  $\delta$  and  $f_F$  becomes increasingly time consuming. Figure 5 compares the closest simulated three-scale network  $P_c(S)$  curve with the laboratory measurement. This three-scale PNM is generated by integration of the 2.3 and 12μm networks ( $\delta$ =1,  $f_F$ =0.004) followed by integration of that network with the 32.8µm network  $(\delta=0.7, f_F=0.006)$ . The nesting-domain sizes for the 2.3 and 12µm networks are close, hence selecting  $\delta$ =1 is feasible. To keep the three-scale network computationally manageable while ensuring representativity, we selected  $\delta$ =0.7 for the second step. Sensitivity analysis was performed on the value of  $f_F$  within the range of 0.001 to 0.1 for the first step and 0.001 to 0.006 for the second step. The calculated  $P_c(S)$  curves show that keeping the first  $f_F$  value at 0.004 enables the inclusion of a sufficient number of elements from the 2.3µm network for simulations to predict the higher capillary-pressures while eliminating artificial jumps in capillary-pressure. We analysed the goodness of fit for the  $P_c(S)$  curves for the single (at 2.3  $\mu$ m and 12  $\mu$ m), twoscale, and three-scale PNMs using the Chi-squared measure. Single-scale networks yield the poorest capillary-pressure predictions while the three-scale network and the two-scale (32.8-12μm) show only minor deviations from the laboratory measurements (Figure 5). Chisquared for the 32.8-12µm network is marginally better that of the 32.8-12-2.3µm network,

since the Chi-squared test only compares model predictions against the laboratory measurements, but does not capture that the model does not predict capillary-pressures at low saturations for the 32.8-12µm combination. Although the calculated absolute permeability values (Table 2) do not accurately match the experimental observation (50mD), the calculated values are close and within the same order of magnitude. The deviation could stem from (i) the fact that the simulations are conducted on a subsection of the acquired µCT images, (ii) errors introduced in noise filtering and subsequent segmentation steps, and (iii) image properties, in particular, the partial volume effect. This can cause features (e.g. pore-throats) smaller than one voxel to appear as large as a voxel in the segmented images. Larger pore-throats display less resistance to the flow and hence a higher absolute permeability is calculated. No laboratory kr(S) measurements were available, hence simulated kr(S) curves for single and multi-scale PNMs are compared in SI. In summary, we show that the parameters involved in the network integration, as well as the number and resolution of the input singlescale networks can have a substantial impact on the shape of the kr(S) curves calculated based on the multi-scale PNMs. Future studies could aim to compare the calculated kr(S)curves with laboratory measurements to establish the effect of network integration on

## 4. Multi-scale Network Generation Workflow

- Figure 6 is a practical guide for selection of length-scales for generating a multi-scale PNM from µCT data and is additionally informed by thin-section imaging and MICP techniques.

  This information assists sample size selection, the choice of length-scales, and resolutions for
- 227 multi-scale network generation.

predicting kr(S) curves.

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## 5. Summary and Conclusions

A network integration approach introduced by *Jiang et al.* [2013] is used to generate representative models for a dolomite with multi-scale porosity. PNMs extracted from  $\mu$ CT images acquired at different length-scales are integrated into single multi-scale PNMs incorporating all the porosity features that exist in the rock. The main contribution of this paper is to present a workflow for effective selection of appropriate length-scales for generation of the multi-scale network. The choice of images to be combined is informed by MICP throat-size distributions. Additionally, there are two free parameters ( $\delta$  and  $f_F$ ) in the model. The integrated networks are used to reproduce  $P_c(S)$  curves, evaluated against MICP laboratory measurements. We demonstrate that  $P_c(S)$  curves calculated based on the two-scale and three-scale PNMs provide significantly closer comparisons to the laboratory measurements compared to a single-scale network model. Future studies need to validate calculated kr(S) curves against experimental measurements to optimise multi-scale network generation for predictions of multiphase fluid behaviour.

## **Acknowledgement:**

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## 310 Nomenclature Table:

C	, , <del>,</del>
S	saturation

S\* the saturation at which the jump occurs

 $S_{air}$  air saturation

 $f_F$  Fraction of fine network

 $\delta$  Nesting-domain fraction

 $P_c(S)$  relative permeability

 $k_r(S)$  relative permeability

*krnw(S)* non-wetting phase relative permeability

*krw(S)* wetting phase relative permeability

 $k_{abs}$  absolute permeability

r radius

 $\chi^2$  Chi-squared

## 311 Tables:

# Table 1: Network parameters for PNM shown in Figure S3.

	oxel Do	main				Total	
	ize	omain	Number of	Number of	Connected	imaged	$k_{abs}$
	um)		Nodes	Bonds	Porosity %	Porosity	(mD)
for spot size	(vo	oxels)				%	

32.8	32	881×901×901	10318	12736	0	3.52	0
21	21	821×860×720	19897	24093	0	6.03	0
12	5	500×500×500	3418	4127	6.15	9.23	75.63
2.3	2	570×450×380	1733	2089	12.72	14.32	99.75

#### Table 2: Network parameters for multi-scale PNMs

Image Resolution (μm)	Number of Nodes	Number of Bonds	Connected Porosity %	Total Porosity %	$k_{abs}$ (mD)	Average coordination number
32.8-12-2.3	13956	22954	3.81	3.96	128.14	3.44
32.8-2.3	439131	534791	3.31	3.36	178.02	3.02
32.8-12	76916	100743	4.28	4.43	90.55	2.94

# **Figure Captions:**

Figure 1: (a) Backscattered SEM image of a polished thin-section of SD in three different magnifications, acquired at SEM facility at Heriot-Watt University (resolution ~0.7μm). The left hand side image represents a tessellated image constructed from 400 fields of view stitched together, (b) Four μCT images of SD rock samples (D=2, 5, 25 and 38mm) captured at respective resolutions of 2.3, 12, 21 and 32.8μm. The scanned samples were not generated sequentially by cutting the same core, (c) PSD derived from the four images shown in (b) using a sphere fitting method [Jiang et al., 2007].

Figure 2: (a) Networks extracted from ten sub-domains with  $\delta$  ranging from 0.1 to 1 for the coarsenetwork (resolution 21 $\mu$ m), blue colour shows connected pink is globally disconnected elements, (b) Static properties of the sub-domain networks shown in (a), the blue box corresponds to the jump in the total porosity captured for  $\delta$ =0.4 while the red box shows the locally connected cluster captured in the sub-domain with  $\delta$ =0.5, (c) Effect of fraction of fine-network ( $f_F$ ) on porosity and permeability of two-scale networks with  $\delta$ =0.7 (see 3D renderings in Figure S4), and (d) Effect of nesting-domain size on the two-scale (21 and 2.3 $\mu$ m,  $f_F$ =0.01)  $P_c(S)$  curves in comparison with laboratory measurements.

Figure 3: (a) The porosity of the sub-domains of the 32.8 $\mu$ m network as a function of the domain fraction, (b) The  $P_c(S)$  curves for the combined 32.8 – 2.3 $\mu$ m networks compared with the MICP laboratory measurements. The 32.8 $\mu$ m network becomes sufficiently representative for  $\delta$ >0.6 displaying converged porosity and  $P_c(S)$  curves.  $\delta$ : Nesting-domain fraction,  $f_F$ : Fraction of fine-network.

334	Figure 4: The $P_c(S)$ curves for the two-scale 32.8–12 $\mu$ m ( $\delta$ =0.7) network for various $f_F$ . The model
335	predictions show a good fit to the experimental data, however, the model does not predict the higher
336	pressure end of the curve.
337	Figure 5: The $P_c(S)$ curves for the three-scale network compared with the laboratory data, two-scale and
338	single-scale simulations. The plot also shows the Chi-squared ( $\chi^2$ ) measure for these networks. The three-
339	scale network model displays the best fit to the MICP data.
340	Figure 6: Multi-scale network generation workflow and tuning.
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Figure 1. Figure

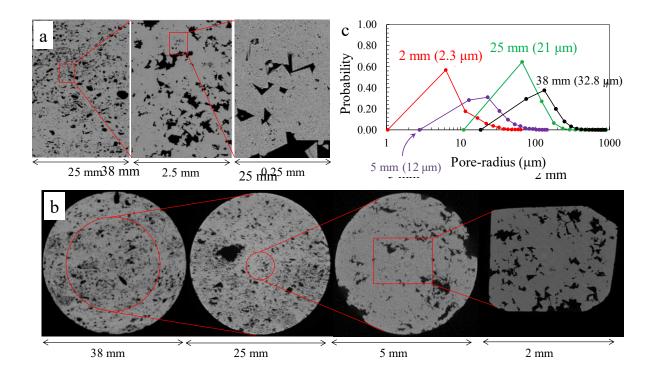


Figure 2. Figure

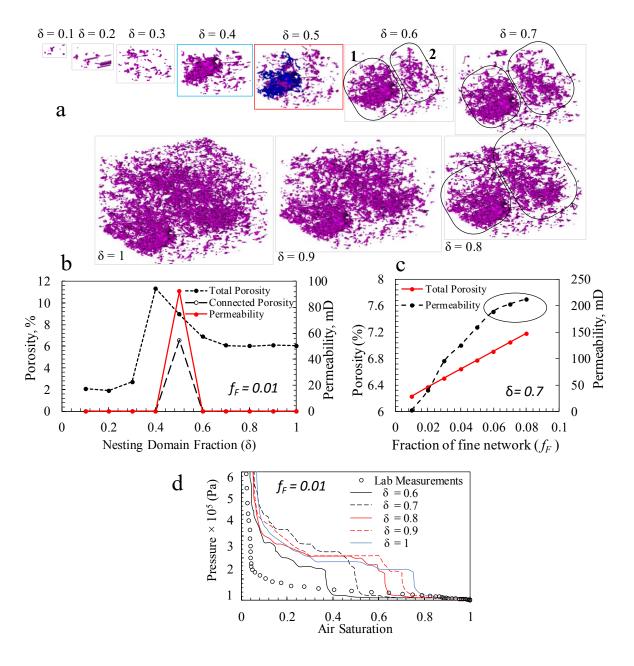


Figure 3. Figure

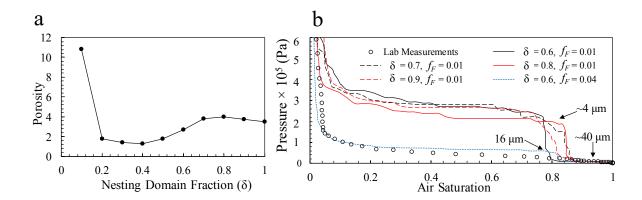


Figure 4. Figure

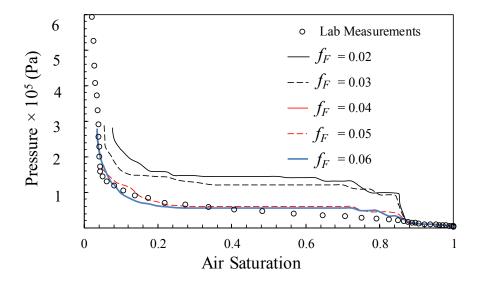


Figure 5. Figure

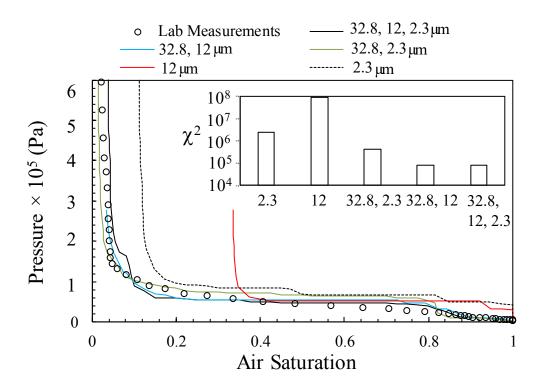
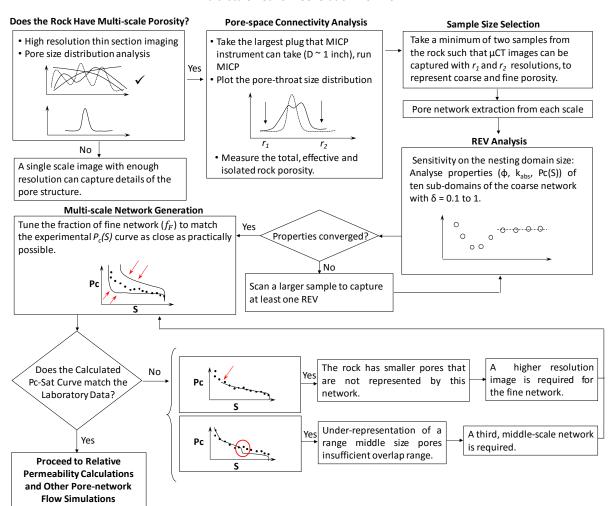


Figure 6. Figure

#### Multi-scale Network Generation Workflow



## Journal of Water Resources Research

# Supporting Information for

## Multi-scale Pore Network Representation of Heterogeneous Carbonate Rocks

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## Contents of this file

- Figures S1 to S5
- Methodological details
- A comparison of simulated kr(S) curves for single or multi-scale pore-networks.

# Introduction

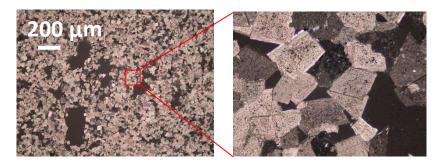


Figure S1: Microscope images for SD showing well sorted crystal sizes and uniform spread of the finer porosity in the rock.

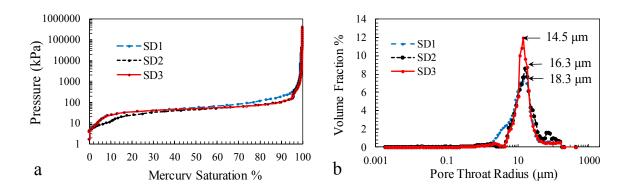


Figure S2: (a)  $P_c(S)$  curve for three SD plugs (D = 25 mm) from which pore-throat size distributions (b) are extracted.

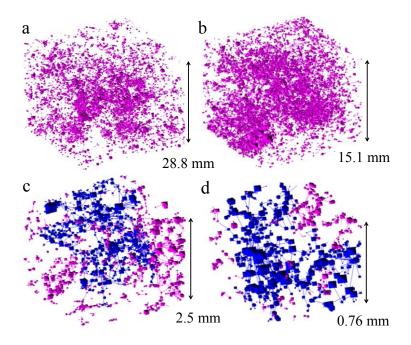


Figure S3: Pore-networks extracted from the four  $\mu$ CT images of the SD samples shown in Figure 3c. Pink and blue indicate the disconnected and connected elements, respectively. See Table 1 for more properties of these four networks.

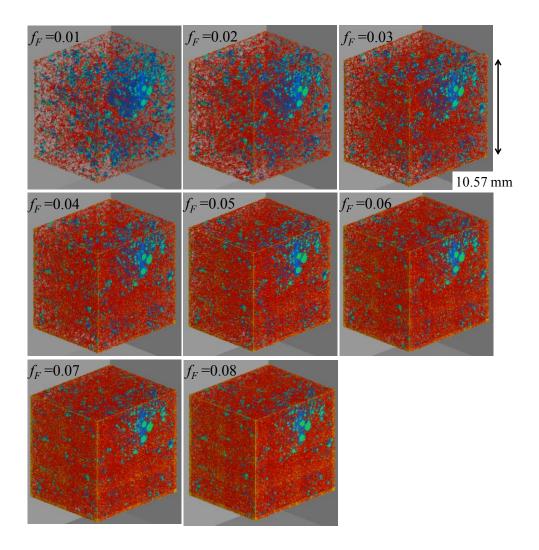


Figure S4: 3D renderings of the two-scale networks with  $\delta = 0.7$  and  $0.01 \le f_F \le 0.08$ . Blue/green shows the coarse-network elements and red shows the fine-network elements.

# **Methodological Details**

X-ray  $\mu$ CT imaging was performed using a  $\mu$ CT instrument built in-house at the University of Edinburgh with a cone-beam configuration. Data were reconstructed using Octopus v 8.5 and processed using Avizo-Fire (v8). Images were filtered using an anisotropic diffusion filter, and segmented using the watershed segmentation method.

# **Simulation of Relative Permeability Curves**

Comparison of experimental kr(S) curves and model predictions for complex (multi-scale) pore structures has not been addressed in many publications (as an example see *Bultreys et al.* [2015]) mainly because of (i) limited availability of experimental data, and (ii) the challenging nature of generating pore-networks that are representative of complex porous media at the scale at which the experiment was performed. What is most commonly reported is the comparison

of simulated kr(S) curves for single or multi-scale pore-networks with a range of the model inputs. For instance,  $Prodanovi\acute{c}$  et al. [2015]. compared the kr(S) curves calculated based on single-scale and two-scale pore-network models and explained the curves display a shift in cross-over point saturation as a result of including micro-porosity. No laboratory kr(S) measurements were available to this research. Therefore, only calculated kr(S) curves based on two-scale and three-scale networks are compared here, (Figure S5). The kr(S) curves are calculated for a drainage process, here we use mercury invading a fully air saturated network. The curve calculated for the mercury/air system can be modified to generate oil/brine curves, by accounting for the difference in the interfacial tension and contact angles.

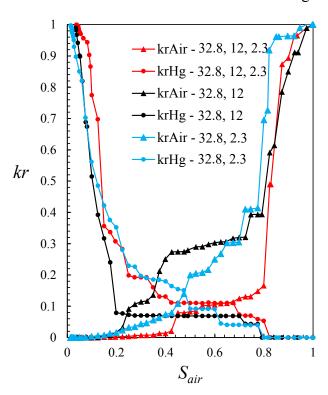


Figure S5: kr(S) curves are calculated for a drainage process, mercury invading a fully air saturated network.

One important parameter to study on relative permeability curves is the saturation where the non-wetting phase (here mercury) breaks through, i.e. the saturation where krnw(S) > 0. The two-scale network models appear to have very close breakthrough saturations ( $S_{air} \sim 0.8$ ) while the three-scale network displays a slightly earlier breakthrough ( $S_{air} \sim 0.82$ ). This is probably due to the higher average coordination number (3.44) of the three-scale network compared to that of the two-scale network ( $\sim$ 3) (Table 2). Coordination number is defined as the number of pore-throats connected to each pore. The irreducible air saturation mostly reflects the proportion of the isolated pores present in each network, which are small

This work suggests that the parameters involved in the network integration, as well as the number and resolution of the input single-scale networks, can have a substantial impact on the shape of the kr(S) curves calculated based on the multi-scale network models. As suggested by *Prodanović et al.* [2015] the saturation of the cross-over point displays considerable shifts for different multi-scale pore-networks discussed in this study.