

SUPPORTING INFORMATION

NLDFT Pore Size Distribution in Amorphous Microporous Materials

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SIMULATED STRUCTURES

The majority of materials presented in this study were synthesized *in silico* using *Polymatic*^{1,2} (except for OMIMs) followed by a compression/decompression scheme³ for obtaining final structures. The force field abbreviation are as following: TraPPE-UA = Transferable Potentials for Phase Equilibria United Atom,^{4–6} GAFF = Generalized Amber Force Field,⁷ and PCFF = Polymer Consistent Force Field.⁸

Table S1 - Summary of microporous materials used in this work.

Name	Description	Composition	Number of monomers	Box length	Non-bonded force field	Bonded force field
PIM-1	Polymer of intrinsic microporosity ⁹	100% PIM-1	100 PIM-1 monomers	$43.5 \pm 0.2 \text{ \AA}$	TraPPE-UA	GAFF
HCP1	Hypercrosslinked polymer ¹⁰ - Poly(vinylbenzyl chloride- <i>co</i> -divinylbenzene)	98% vinylbenzyl chloride (VBC), 2% divinylbenzene (DVB)	196 VBC monomers, 4 DVB monomers	$37.4 \pm 0.3 \text{ \AA}$	TraPPE-UA	PCFF
HCP2	Hypercrosslinked polymer ¹¹ - Poly(styrene- <i>co</i> -vinylbenzyl chloride)	25% styrene (STR), 75% vinylbenzyl chloride (VBC)	50 STR monomers, 150 VBC monomers	$37.1 \pm 0.5 \text{ \AA}$	TraPPE-UA	PCFF
HCP3	Hypercrosslinked polymer ¹¹ - Poly(vinylbenzyl chloride)	100% vinylbenzyl chloride (VBC)	200 VBC monomers	$37.8 \pm 0.3 \text{ \AA}$	TraPPE-UA	PCFF
PY1	Conjugated microporous polymer ¹²	100% 1,3,6,8-tetrabromopyrene (A_4)	200 A_4 monomers	$55.6 \pm 0.4 \text{ \AA}$	TraPPE-UA	PCFF
PY2	Conjugated microporous polymer ¹²	75% 1,3,6,8-tetrabromopyrene (A_4), 25% 1,3-dibromo-7- <i>tert</i> -butylopyrene (B_2)	150 A_4 monomers, 50 B_2 monomers	$43.1 \pm 0.8 \text{ \AA}$	TraPPE-UA	PCFF
OS-DVB1	0% crosslinked polyolefin ¹³	81% octene, 19% styrene	162 octene monomers, 38 styrene monomers	$35.5 \pm 0.1 \text{ \AA}$	PCFF	PCFF

Table S1 - Summary of microporous materials used in this work. (Continued)

Name	Description	Composition	Number of monomers	Box length	Non-bonded force field	Bonded force field
OS-DVB2	100% crosslinked polyolefin ¹³	4.6% styrene, 95.4% crosslinked unit	9 styrene monomer, 191 crosslinked unit	$35.3 \pm 0.2 \text{ \AA}$	PCFF	PCFF
DCX	Microporous polymer network ¹⁴ - Poly(dichloroxylene)	100% dichloroxylene	176 dichloroxylene monomers	$32.4 \pm 0.3 \text{ \AA}$	PCFF	PCFF
OMIM1	Organic molecules of intrinsic microporosity with 1,4 fragments ¹⁵	100% OMIM 1+4	100 OMIM 1+4 monomers	$61.0 \pm 0.2 \text{ \AA}$	PCFF	PCFF
OMIM2	Organic molecules of intrinsic microporosity with 4,8 fragments ¹⁵	100% OMIM 4+8	100 OMIM 4+8 monomers	$75.9 \pm 0.7 \text{ \AA}$	PCFF	PCFF

Table S2 - Regularization parameter used in this work.

Name	λ for NLDFT	λ for SSNLDFT
PIM-1	1.00	5.00
HCP1	0.75	4.50
HCP2	0.75	4.50
HCP3	1.00	4.50
PY1	0.75	5.25
PY2	0.25	5.00
OS-DVB1	0.75	5.25
OS-DVB2	0.00	4.75
DCX	0.25	4.75
OMIM1	0.00	4.50
OMIM2	0.25	4.50

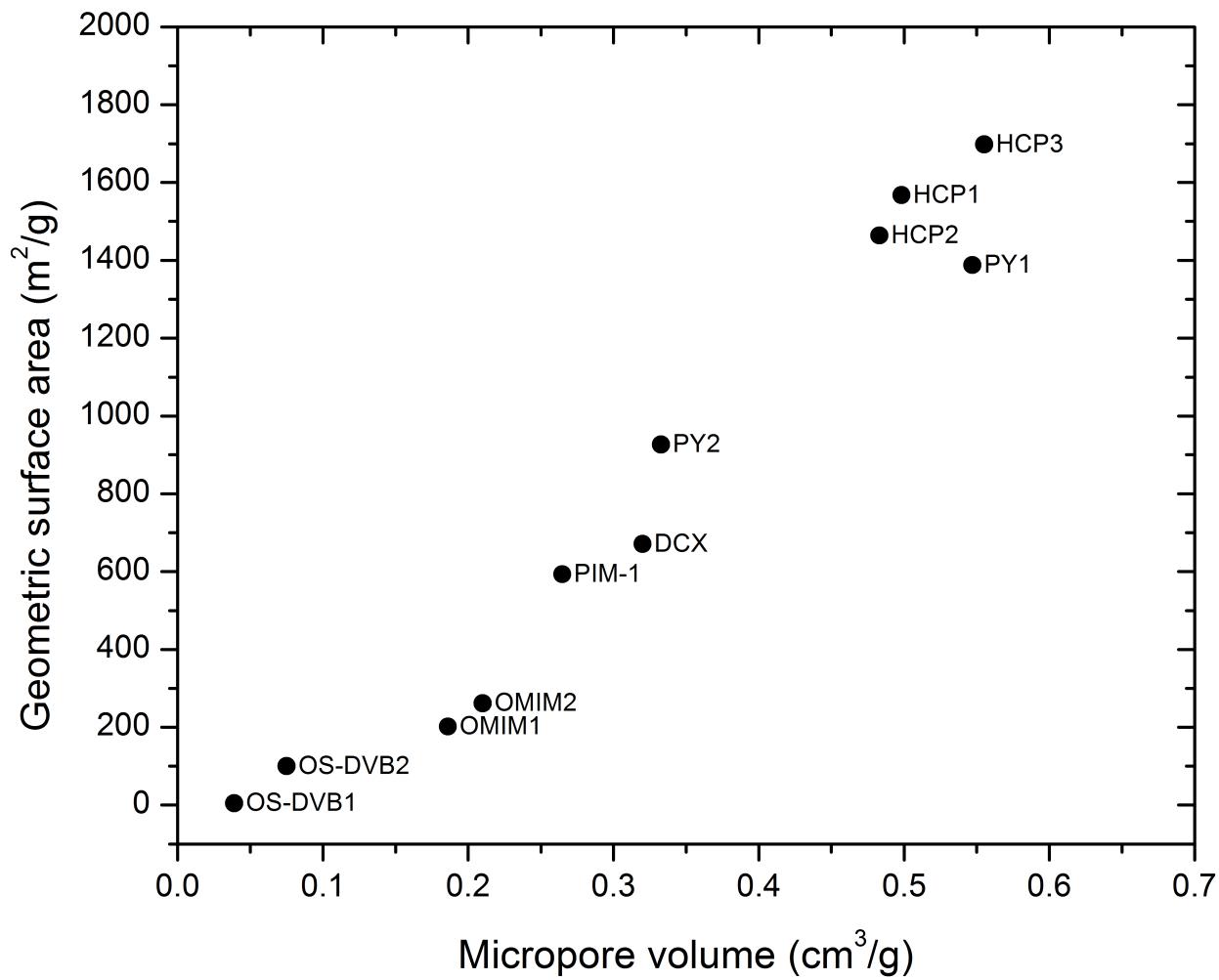


Figure S1 - The range of geometric surface area and micropore volume of amorphous materials investigated in this study.

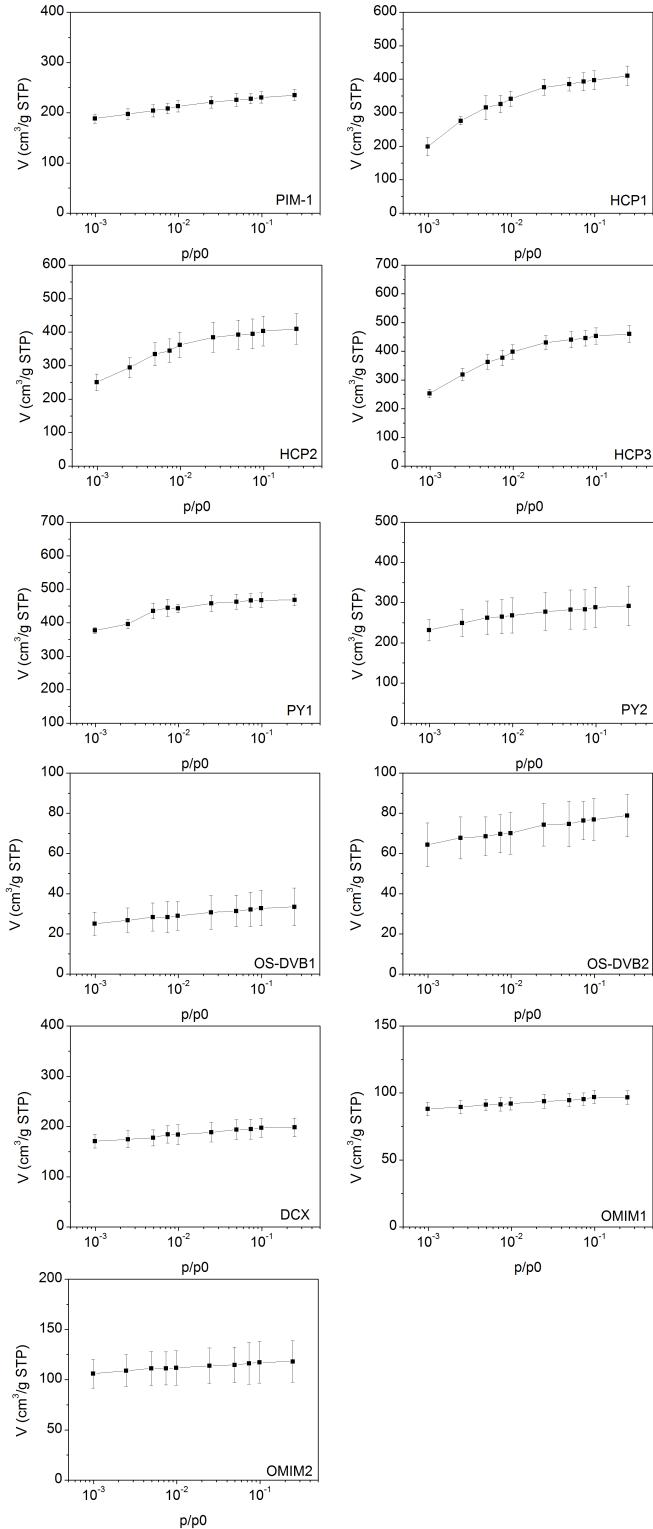


Figure S2 - N_2 adsorption isotherms at 77 K obtained from isobaric-isothermal Gibbs ensemble Monte Carlo (NPT-GEMC) simulations.

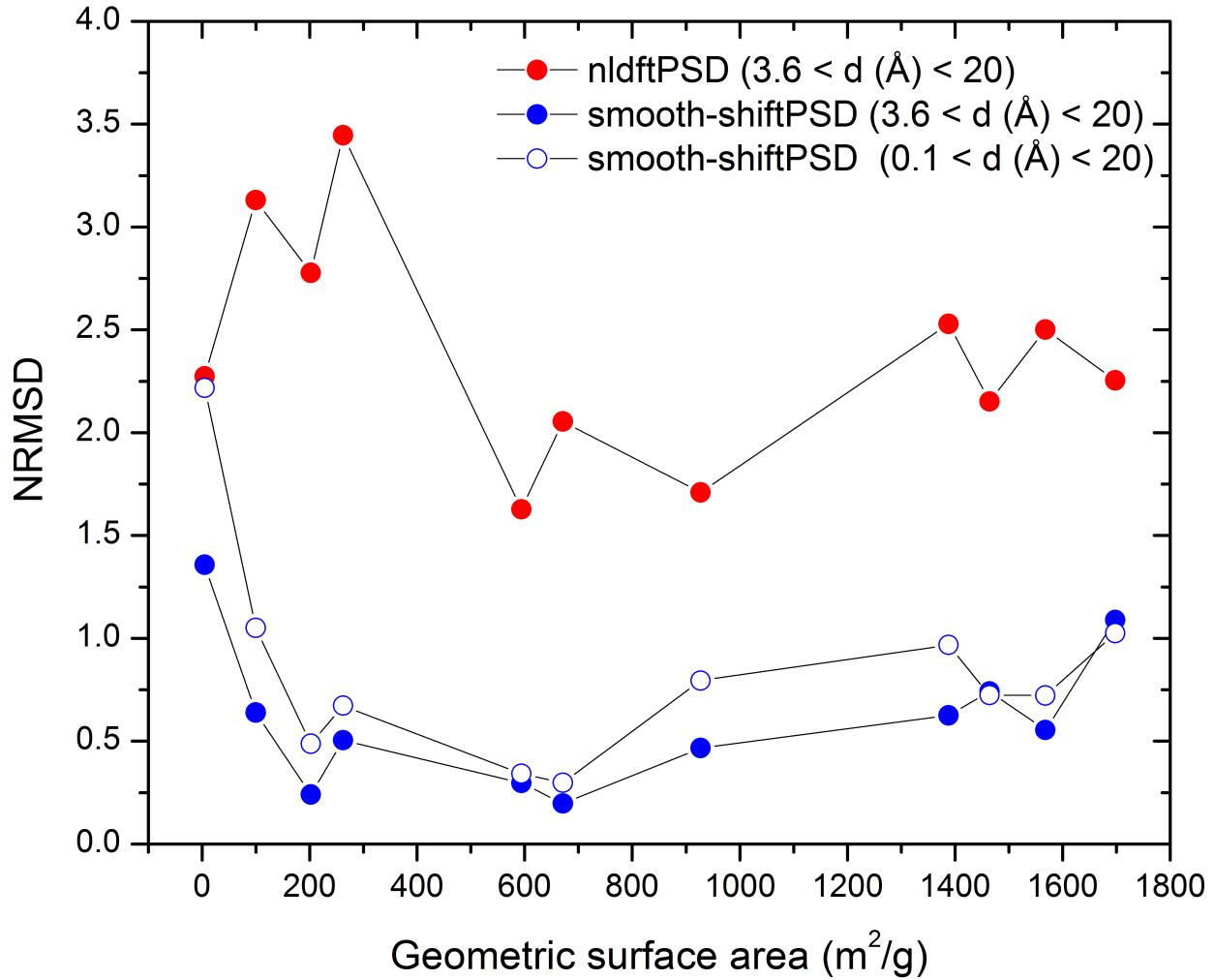


Figure S3 - Normalized root-mean-square deviation (NRMSD) as a function of the geometric surface area. NRMSD measured the deviation from the exact geometric pore size distribution. NRMSD was normalized by the mean of geometric PSD specifically for each material (Eq. S1).

$$NRMSD = \frac{RMSD}{\bar{y}} = \sqrt{\frac{\sum (PSD_i - PSD_{geo,i})^2}{n}} \quad (\text{Eq. S1})$$

In this equation, RMSD is defined by summation of the difference squared between PSD of either NLDFT or smooth-shift and geometric PSD at the same diameter. The mean value of geometric PSD, \overline{PSD}_{geo} , is chosen as the mean value of measurement, \bar{y} , which normalizes RMSD. The normalization is required in order to compare materials with different PSD scales. The analysis includes evenly spaced 100 data points ($n=100$) using the range (d) indicated in Figure S3.

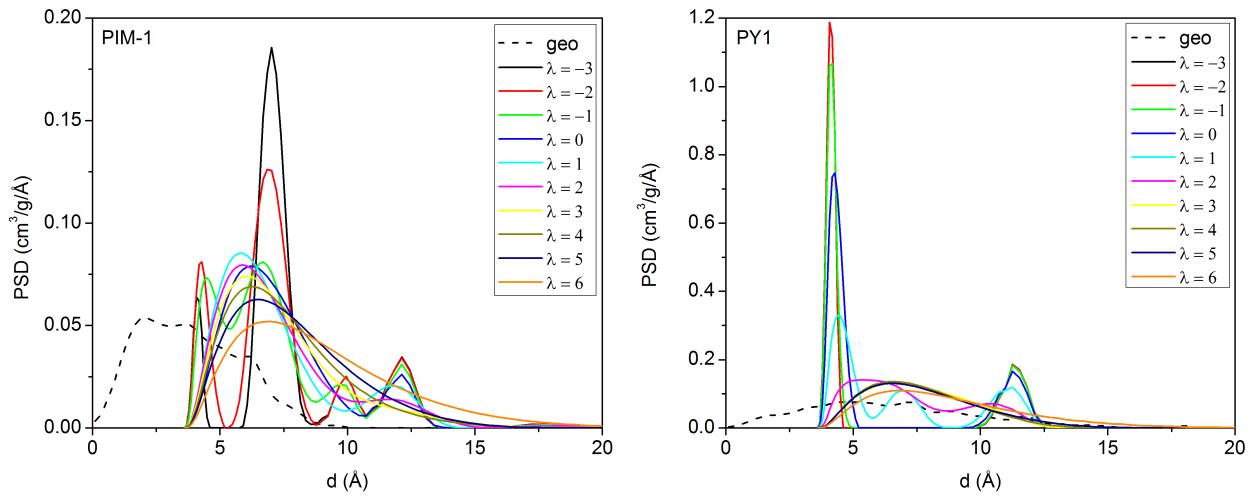


Figure S4 - NLDFT PSDs for PIM-1 (left) and PY1 (right) as a function of the regularization parameter, λ .

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