

## Impact of NH<sub>4</sub>OH treatment on the ion-exchange and pore characteristics of metakaolin-based geopolymer

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## 1. Variable-temperature $^{129}\text{Xe}$ spectra and fits

$^{129}\text{Xe}$  spectra collected under various temperatures of four samples are shown in Fig. S1.

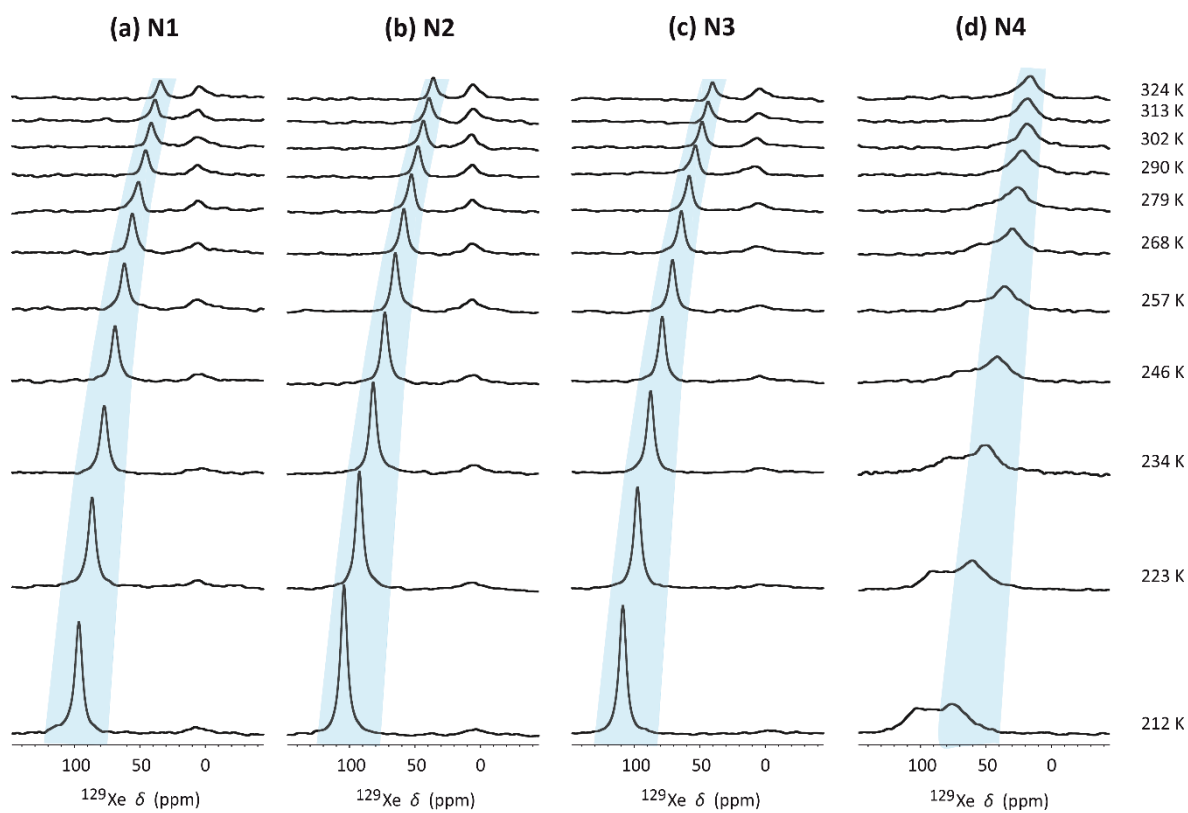


Figure S1. The  $^{129}\text{Xe}$  spectra acquired under variable temperature of (a) N1, (b) N2, (c) N3 and (d) N4.

## 2. $^{129}\text{Xe}$ dynamic data and fits

The  $^{129}\text{Xe}$  dynamic in site IP and BP were detected by measuring exchange rate ( $k$ ), spin-lattice relaxation time ( $T_1$ ) and spin-spin relaxation time ( $T_2$ ). Selective-Inversion-Recovery (IR),<sup>1</sup> IR,<sup>2</sup> and Carr-Purcell-Meiboom-Gill (CPMG)<sup>3</sup> experiments were performed to get these three values, respectively. All of these data were analysed according to the following theories in MATLAB R2017b (Mathworks, Natick, Massachusetts, United States of America).

Based on the previous studies,<sup>1,4,5</sup> when the exchange process takes place between two sites, IP and BP, the time dependence of their magnetizations under  $^{129}\text{Xe}$  selective-IR experiment is given by,

$$\frac{\partial}{\partial t} \begin{pmatrix} M_{IP}(t) - M_{eq,IP} \\ M_{BP}(t) - M_{eq,BP} \end{pmatrix} = - \begin{pmatrix} R_{1,IP} + k_{IP-BP} & -\frac{M_{eq,IP}}{M_{eq,BP}} * k_{IP-BP} \\ -k_{IP-BP} & R_{1,BP} + \frac{M_{eq,IP}}{M_{eq,BP}} * k_{IP-BP} \end{pmatrix} \begin{pmatrix} M_{IP}(t) - M_{eq,IP} \\ M_{BP}(t) - M_{eq,BP} \end{pmatrix}, \quad (S1)$$

where  $M_{IP}(t)$  and  $M_{BP}(t)$  are the magnetizations for two sites at time  $t$ ,  $M_{eq,IP}$  and  $M_{eq,BP}$  are the equilibrium magnetizations,  $R_{1,IP}$  and  $R_{1,BP}$  are the spin-lattice relaxation rates ( $1/T_1$ ) for two sites,  $k_{IP-BP}$  is the exchange rate between two sites.

Then our  $^{129}\text{Xe}$  selective-IR data were fitted using the solution of Eq. S1, which is,

$$\begin{pmatrix} M_{IP}(t) \\ M_{BP}(t) \end{pmatrix} = \begin{pmatrix} M_{eq,IP} \\ M_{eq,BP} \end{pmatrix} + \exp \left[ - \begin{pmatrix} R_{1,IP} + k_{IP-BP} & -\frac{M_{eq,IP}}{M_{eq,BP}} * k_{IP-BP} \\ -k_{IP-BP} & R_{1,BP} + \frac{M_{eq,IP}}{M_{eq,BP}} * k_{IP-BP} \end{pmatrix} t \right] \begin{pmatrix} M_{IP}(0) - M_{eq,IP} \\ M_{BP}(0) - M_{eq,BP} \end{pmatrix}, \quad (S2)$$

ere  $M_{IP}(0)$  and  $M_{BP}(0)$  are the magnetizations at  $t=0$ .

The  $^{129}\text{Xe}$   $T_1$ IR signals intensities of peak IP and BP were derived by Topspin software (Bruker, Rheinstetten, Germany) and then fitted by the following equation, respectively,

$$M(t) = M_{eq} * \left[ 1 - n * \exp \left( \frac{-t}{T_1} \right) \right], \quad (S3)$$

where  $M(t)$  is the magnetization at time  $t$ ,  $M_{eq}$  are the magnetization at equilibrium-state,  $T_1$  is the spin-lattice relaxation time and  $n$  is a free fitting parameter.

The fit equation for  $^{129}\text{Xe}$  CPMG data is,

$$M(t) = M_{eq} * \exp \left( \frac{-t}{T_2} \right), \quad (S4)$$

where  $M(t)$  is the magnetization at time  $t$ ,  $M_{eq}$  are the magnetization at equilibrium-state,  $T_2$  is the spin-spin relaxation time.

The raw signals and fitted signals of three experiments are shown in Fig. S2. The detailed fit results of these signals are shown in Tables S1-3.

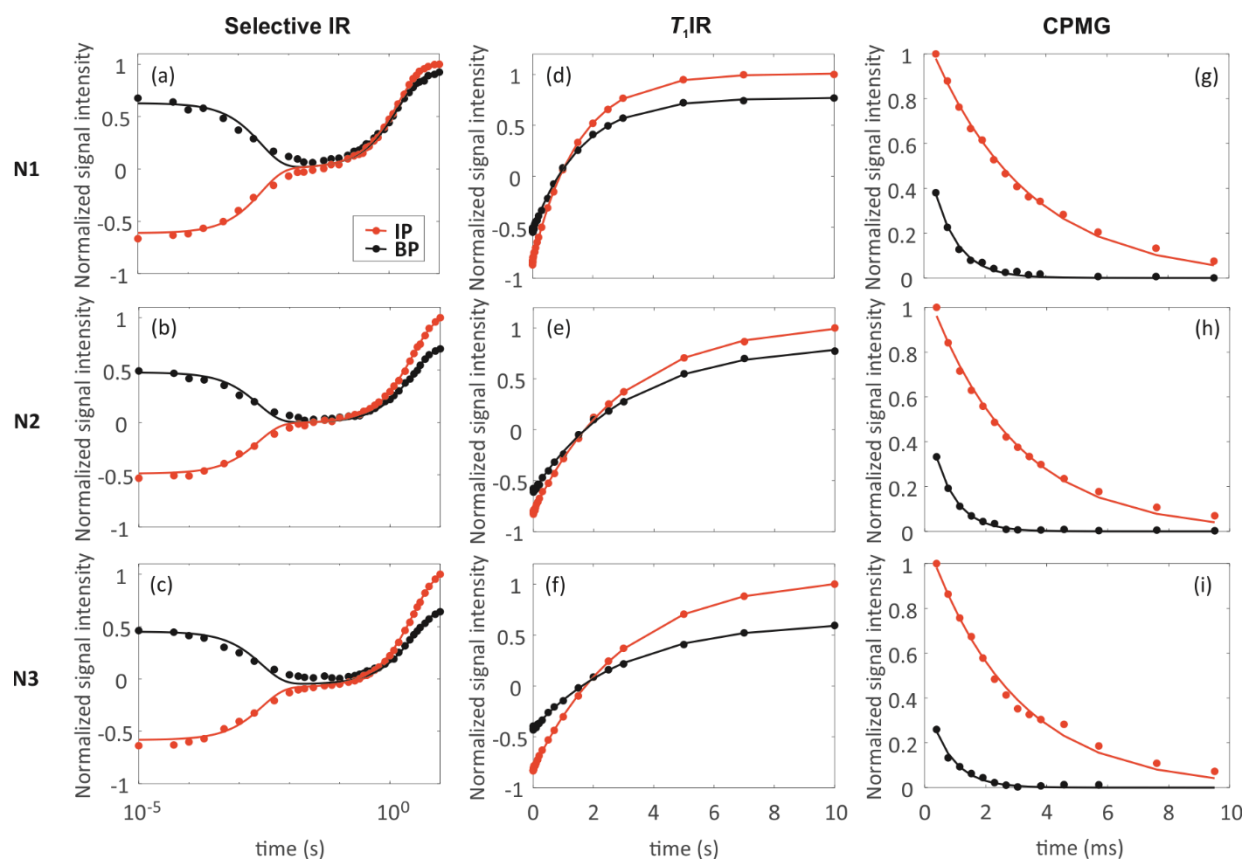


Figure S2. The raw signal (dots) and fitted signal (lines) of (a-c)  $^{129}\text{Xe}$  selective IR, (d-f)  $T_1\text{IR}$  and (g-i) CPMG experiments performed at room temperature of (a, d and g) N1, (b, e and h) N2 and (c, f and i) N3.

Table S1. The fit results of  $^{129}\text{Xe}$  selective-IR data.

SellR	$M_{\text{eq}}$ (IP)	$M_{\text{eq}}$ (BP)	$R_1$ (IP)	$R_1$ (BP)	$k_{\text{IP-BP}}$	$M_0$ (IP)	$M_0$ (BP)	resnorm
<b>N 1</b>	$0.99 \pm 0.03$	$0.91 \pm 0.03$	$1.33\text{e-}3 \pm 4.79$	$1.39 \pm 5.25$	<b><math>181.67 \pm 30.81</math></b>	$-0.61 \pm 0.03$	$0.63 \pm 0.03$	0.075
<b>N 2</b>	$1.01 \pm 0.03$	$0.73 \pm 0.03$	$4.41\text{e-}3 \pm 3.48$	$0.85 \pm 4.86$	<b><math>239.77 \pm 36.23</math></b>	$-0.49 \pm 0.02$	$0.48 \pm 0.02$	0.037
<b>N 3</b>	$1.02 \pm 0.05$	$0.67 \pm 0.04$	$4.84\text{e-}3 \pm 3.57$	$0.87 \pm 5.46$	<b><math>209.22 \pm 40.75</math></b>	$-0.58 \pm 0.03$	$0.45 \pm 0.03$	0.066

Table S2. The fit results of  $^{129}\text{Xe}$   $T_1$ IR data.

$T_1$ IR	N 1		N 2		N 3	
	IP	BP	IP	BP	IP	BP
$M_{\text{eq}}$	$1.01 \pm 0.02$	$0.77 \pm 0.02$	$1.06 \pm 0.03$	$0.85 \pm 0.04$	$1.07 \pm 0.02$	$0.63 \pm 0.02$
$T_1$ (s)	<b><math>1.50 \pm 0.04</math></b>	<b><math>1.60 \pm 0.07</math></b>	<b><math>2.99 \pm 0.10</math></b>	<b><math>3.18 \pm 0.18</math></b>	<b><math>3.05 \pm 0.07</math></b>	<b><math>3.15 \pm 0.16</math></b>
n	$1.83 \pm 0.02$	$1.70 \pm 0.02$	$1.76 \pm 0.02$	$1.71 \pm 0.03$	$1.77 \pm 0.02$	$1.67 \pm 0.03$
resnorm	0.004	0.005	0.003	0.006	0.002	0.002

Table S3. The fit results of  $^{129}\text{Xe}$  CPMG data.

CPMG	N 1		N 2		N 3	
	IP	BP	IP	BP	IP	BP
$M_{\text{eq}}$	$1.11 \pm 0.04$	$0.61 \pm 0.05$	$1.10 \pm 0.04$	$0.57 \pm 0.03$	$1.12 \pm 0.06$	$0.42 \pm 0.05$
$T_2$ (ms)	<b><math>3.20 \pm 0.17</math></b>	<b><math>0.79 \pm 0.08</math></b>	<b><math>2.88 \pm 0.18</math></b>	<b><math>0.72 \pm 0.04</math></b>	<b><math>2.89 \pm 0.23</math></b>	<b><math>0.75 \pm 0.10</math></b>
resnorm	0.004	0.001	0.005	4e-4	0.009	0.001

### 3. <sup>1</sup>H cryoporometry data and fits

The signal intensity at each temperature step ( $S(X)$ ) is dependent on the inverse temperature,  $X=1000/T$ . Their relationship agrees with the following equation,<sup>6</sup>

$$S(X) = \sum_{i=1}^n \frac{S_{0i}}{2} \left[ 1 - \operatorname{erf} \left\{ \frac{X - X_{ci}}{\sqrt{2}\sigma_i} \right\} \right], \quad (S5)$$

where  $n$  is the number of phase transitions,  $S_{0i}$ ,  $X_{ci}$  and  $\sigma_i$  are the signal intensity, the inverse transition temperature and the width of the temperature distribution curve of phase  $i$ , respectively.

Our signal intensity data as a function of  $X$  are shown in Fig. S3. The data were fitted with Eq. S5, assumed with containing three components ( $n=3$ ), S, B and V, by using CFTOOL installed in MATLAB R2017b (Mathworks, Natick, Massachusetts, United States of America). The detailed fit results are shown in Tab. S5.

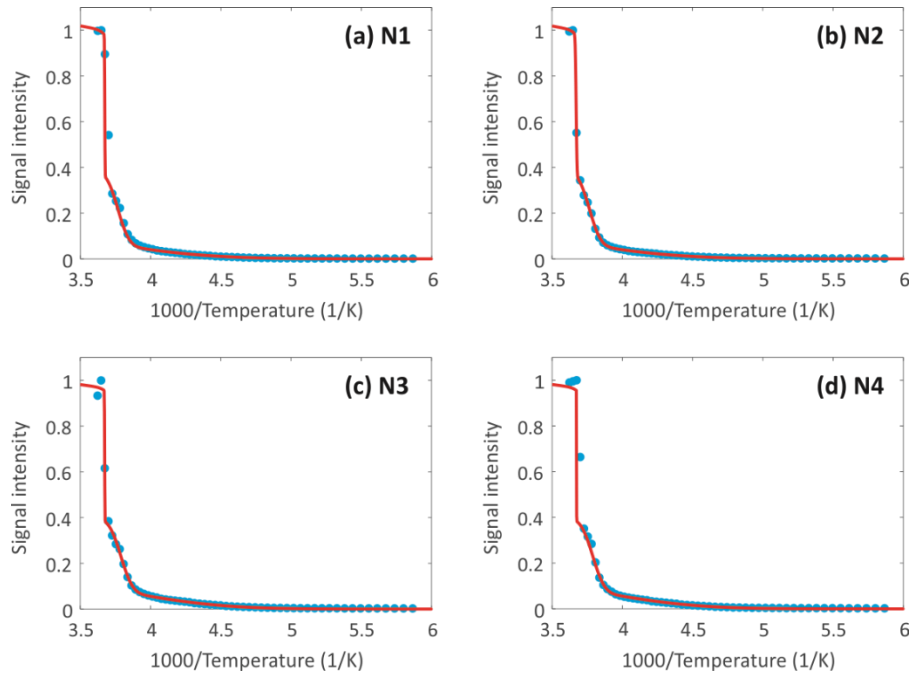


Figure S3. The raw (dots) signal intensity at each temperature step and the fitted (lines) signal intensity as a function of 1000/temperature of (a) N1, (b) N2, (c) N3 and (d) N4.

Table S4. Fit results of cryoporometry data.

		$S_0/2$	$X_c$	$\sqrt{2}\sigma$	$R^2$
<b>N1</b>	<b>S</b>	$0.15 \pm 0.03$	$3.78 \pm 0.01$	$0.09 \pm 0.02$	0.9997
	<b>B</b>	$0.09 \pm 0.25$	$3.50 \pm 1.64$	$0.91 \pm 0.73$	
	<b>BP</b>	$0.31 \pm 0.02$	$3.67 \pm 1e12$	$0.003 \pm 3e12$	
<b>N2</b>	<b>S</b>	$0.15 \pm 0.04$	$3.79 \pm 0.01$	$0.09 \pm 0.03$	0.9983
	<b>B</b>	$0.07 \pm 0.17$	$3.88 \pm 1.45$	$0.73 \pm 0.83$	
	<b>BP</b>	$0.29 \pm 0.01$	$3.67 \pm 1e12$	$0.002 \pm 7e12$	
<b>N3</b>	<b>S</b>	$0.15 \pm 0.04$	$3.79 \pm 0.02$	$0.09 \pm 0.03$	0.9983
	<b>B</b>	$0.07 \pm 0.16$	$3.89 \pm 1.41$	$0.73 \pm 0.81$	
	<b>BP</b>	$0.29 \pm 0.02$	$3.67 \pm 4e11$	$0.006 \pm 2e12$	
<b>N4</b>	<b>S</b>	$0.15 \pm 0.04$	$3.79 \pm 0.01$	$0.09 \pm 0.03$	0.9997
	<b>B</b>	$0.07 \pm 0.16$	$3.89 \pm 1.40$	$0.73 \pm 0.81$	
	<b>BP</b>	$0.29 \pm 0.02$	$3.67 \pm 2e12$	$0.0005 \pm 7e12$	

By plugging the fitted  $S_0/2$ ,  $X_c$  and  $\sqrt{2}\sigma$  into the following equation, the pore size distribution  $dS/da$  as a function of  $a$  were obtained,

$$\frac{dS}{da} = \frac{10^3 k}{\sqrt{2\pi}(aT_{\text{melting}} - k)^2} \sum_{i=1}^n \frac{S_{0i}}{\sigma_i} \times \left[ - \left\{ \frac{10^3 a - X_{ci}(aT_{\text{melting}} - k)}{\sqrt{2}\sigma_i(aT_{\text{melting}} - k)} \right\}^2 \right], \quad (S6)$$

where  $T_{\text{melting}}$  is the melting temperature of bulk water, which is 273 K in our cases,  $k$  is a constant, 89 K\*nm is used here. The  $k$  has the same value as our pervious geopolymer sample<sup>7</sup> we used to evaluate the  $K_0$  in Section S3, and it also agrees with the  $k$  value used for silica gel from one study.<sup>8</sup>

## Reference

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