

# Supporting Information

## **Giant Zn<sub>14</sub> Molecular Building Block in Hydrogen-bonded Network with Permanent Porosity for Gas Uptake**

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## 1 General Remarks

All reagents and solvents were used as purchased from commercial suppliers (Sigma-Aldrich, Fluka, Alfa Aesar, and others) without further purification, if not stated otherwise. Elemental analysis (C, H, N) was performed on Elementar Vario EL elemental analyzer.

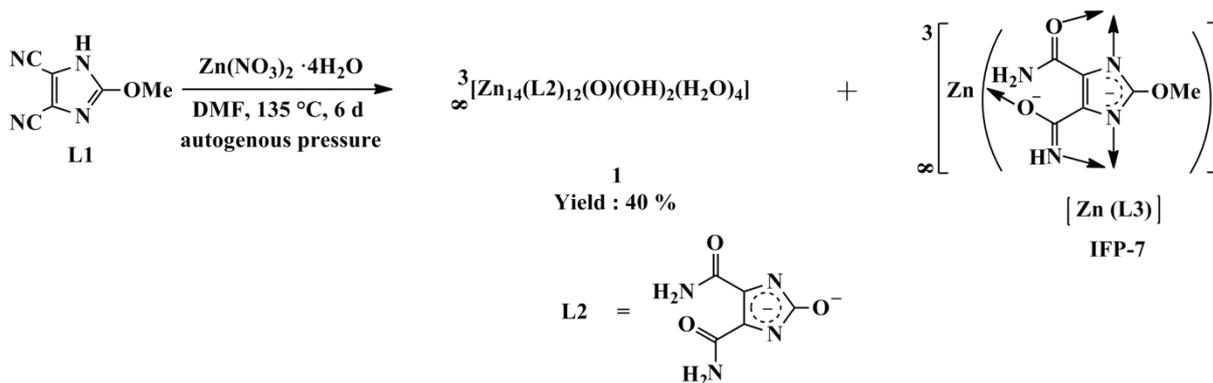
Scanning Electron Micrographs (SEM) image of **1** was taken by Phenom from FEICO.

The linker precursor 4,5-dicyano-2-methoxyimidazole (L1) was synthesized following a published procedure.<sup>1</sup>

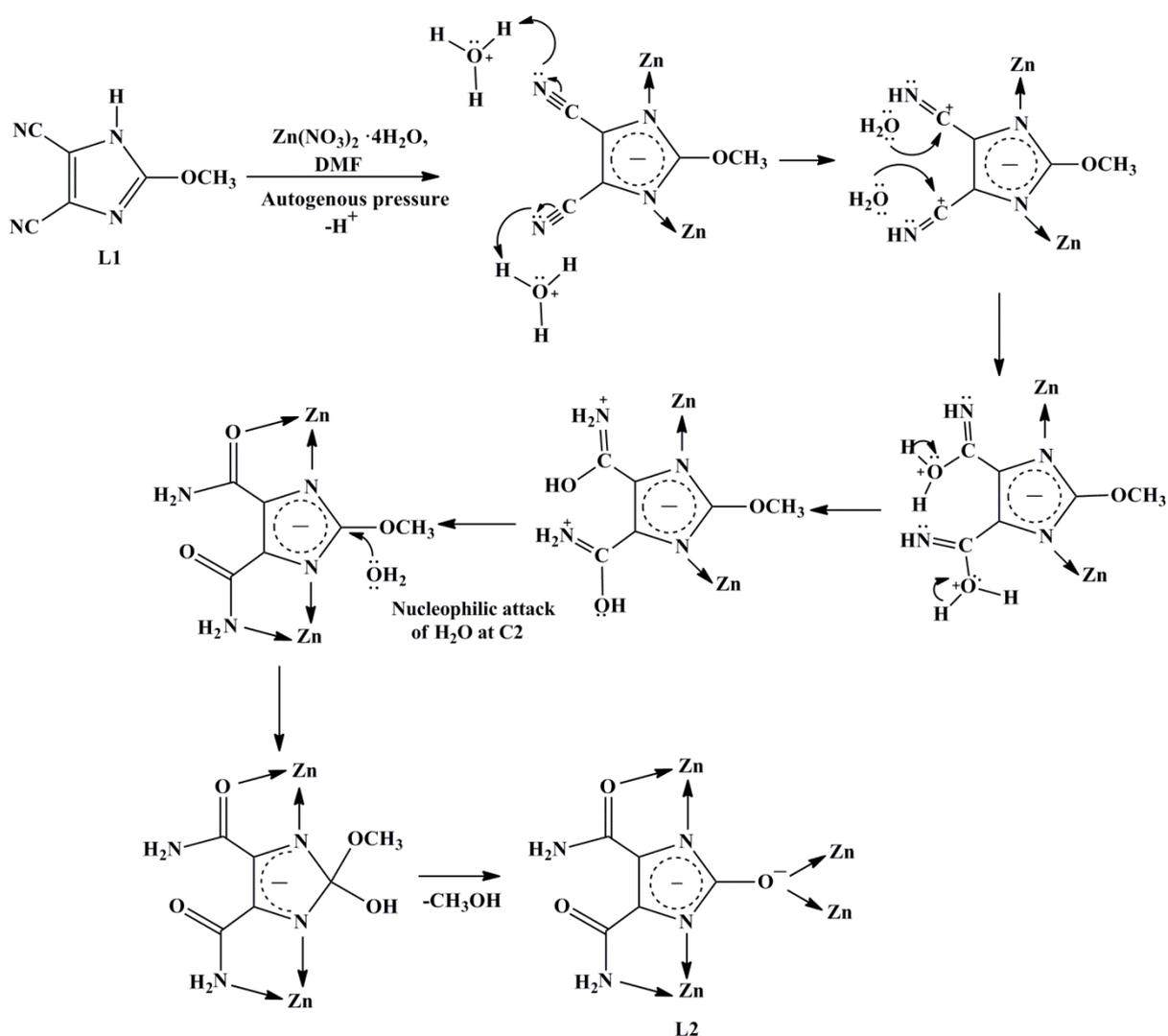
## 2 Experimental Details

Synthesis of  $[\text{Zn}_{14}(\text{L}2)_{12}(\text{O})(\text{OH})_2(\text{H}_2\text{O})_4](\text{DMF})_{18}$  (**1**)

In a sealed tube (Type A, company: Ace) 0.10 g (0.76 mmol) of 4,5-dicyano-2-methoxyimidazole (L1) and  $\text{Zn}(\text{NO}_3)_2 \cdot 4\text{H}_2\text{O}$  (0.17 g, 0.76 mmol) were dissolved in DMF (6 mL). The sealed tube was closed and the mixture was heated at 135 °C for 6 days and was then allowed to cool down to room temperature with 5 °C per hour. Yellow prismatic crystals are formed (~40 wt%) together with a powder material which was identified as the known IFP-7 (Scheme S1, IFP = Imidazolate Framework Potsdam).<sup>2</sup> We optimized the synthetic condition for better yield and X-ray quality crystals. In a modified synthetic procedure the reaction was run for 6 days at 135 °C, instead of 36 h at 130 °C. The yellow crystals were separated by a sieving technique, wherein the crystals were trapped by a mesh (mesh size: 125  $\mu\text{m}$ ) while the powder material (IFP-7) filtered through it.<sup>3</sup> The fine crystalline product was washed with DMF and EtOH and dried in air. Yield: 0.067 g (~40 %) based on L1; <sup>13</sup>C CP-MAS NMR:  $\delta$  =167.1, 164.7, 138.8, 127.7 ppm; elemental analysis:  $[\text{Zn}_{14}(\text{L}2)_{12}(\text{O})(\text{OH})_2(\text{H}_2\text{O})_4] \cdot (\text{DMF})_{18} = \text{C}_{114}\text{H}_{184}\text{N}_{66}\text{O}_{61}\text{Zn}_{14}$ ; Calcd., C 31.33, H 4.24, N 21.15, Found: C 31.57, H 3.82, N 20.69; IR (KBr pellet):  $\nu_{\text{max}} = 3440$  (m), 3355 (m), 3205 (m), 1658 (vs), 1548 (s), 1436 (s), 1280 (m), 1101 (m), 734  $\text{cm}^{-1}$  (m).

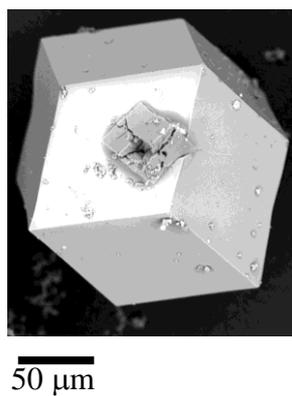


**Scheme S1.** Syntheses of frameworks **1** and IFP-7.



**Scheme S2.** Proposed *in-situ* linker (L2) generation under solvothermal condition.

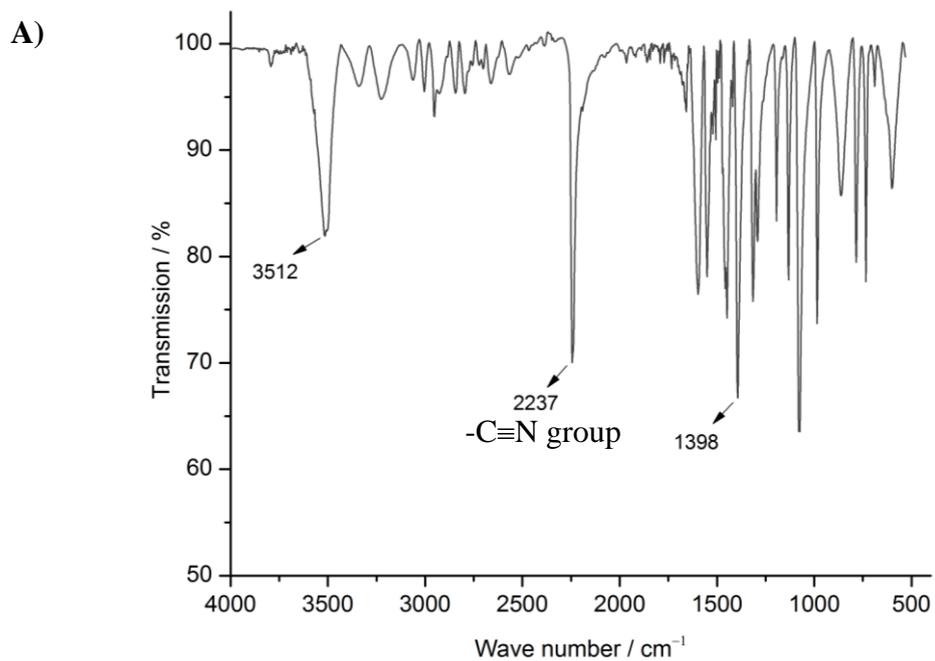
### 3 SEM image

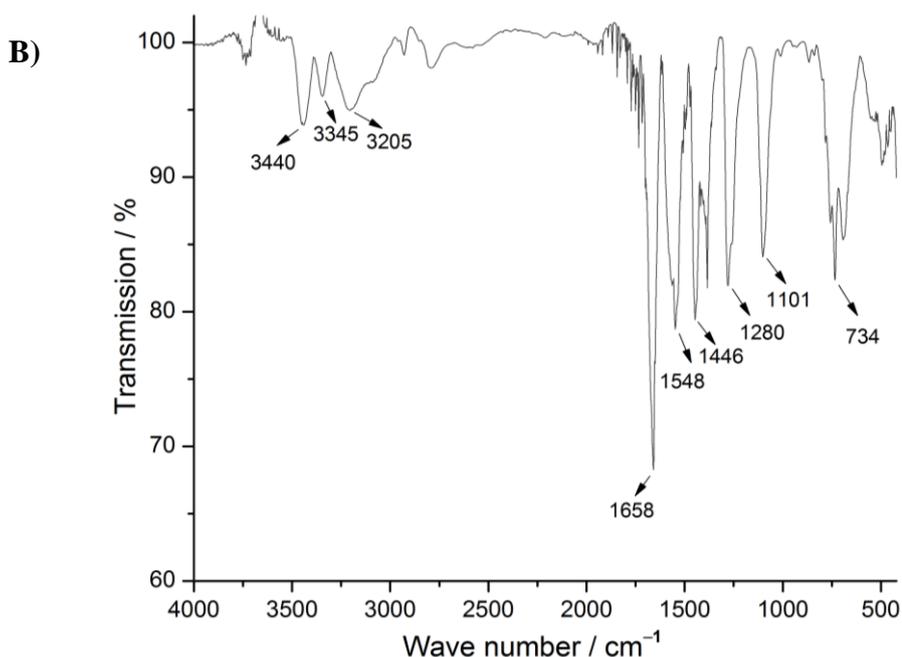


**Figure S1.** Scanning electron micrograph of **1**.

### 4 IR spectra

IR spectra were recorded on FT-IR Nexus from Thermo Nicolet in the region of 4000 – 400  $\text{cm}^{-1}$  using KBr pellets.





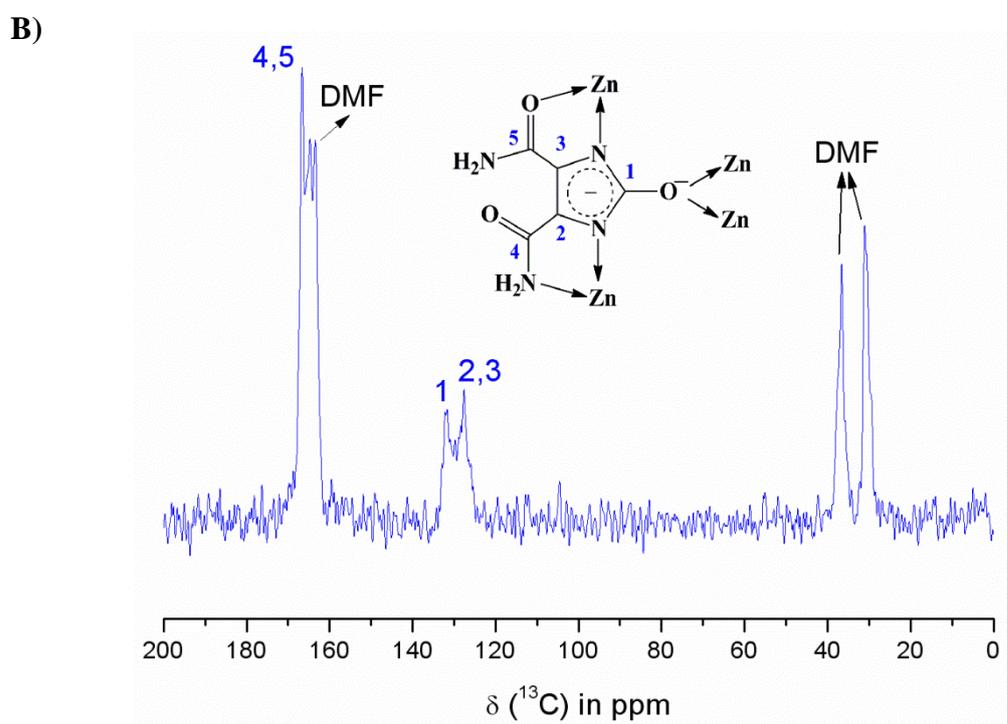
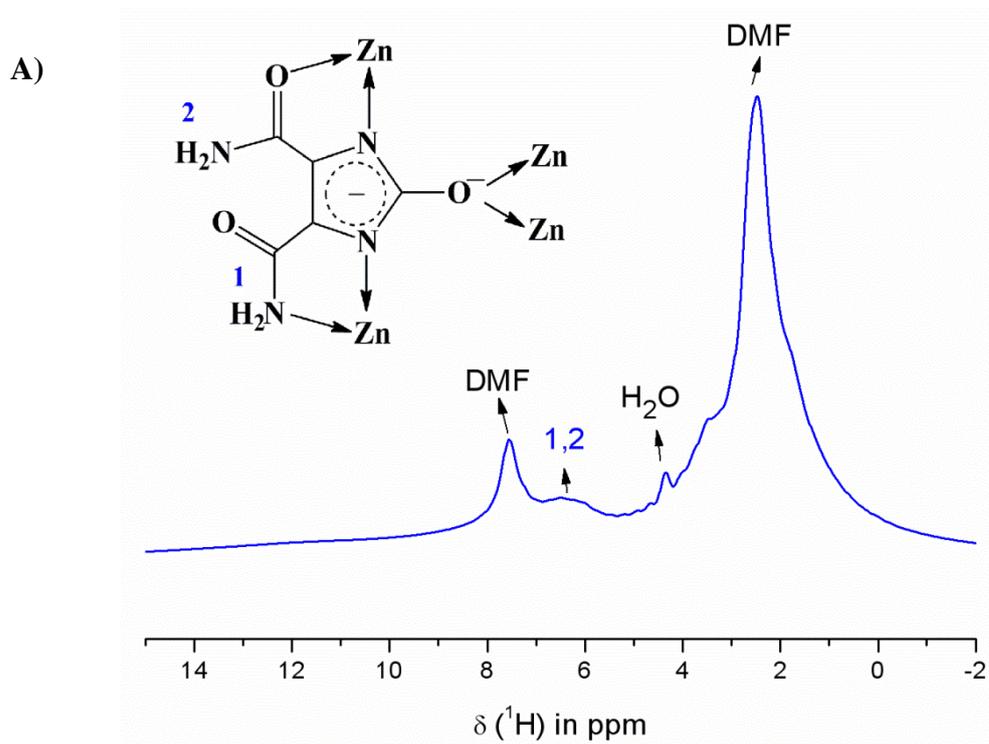
**Figure S2.** IR-spectra of: **A)** 4,5-dicyano-2-methoxyimidazole (L1); **B)** **1** as-synthesized.

## 5 NMR Spectroscopy

### $^1\text{H}$ Magic Angle Spinning (MAS) NMR and $^{13}\text{C}\{^1\text{H}\}$ Cross Polarization (CP) MAS NMR spectra of as-synthesized **1**

$^1\text{H}$  and  $^{13}\text{C}$  Cross Polarization experiments with Magic Angle Sample Spinning ( $^{13}\text{C}$  CPMAS NMR) were performed on a Bruker Avance 600 spectrometer (Bruker Biospin GmbH, Rheinstetten, Germany,  $B_0 = 14.1$  T) operating at a frequency of 150.9 MHz using a double resonant 4 mm MAS (magic angle sample spinning) probe. The spinning frequency was 12.5 kHz. For CP, a ramped (50 % ramp) lock field was used on the  $^1\text{H}$  channel with a CP contact time of 2 ms, a recycle delay of 3 s and 512 scans. The  $90^\circ$   $^1\text{H}$  pulse length was 3.3  $\mu\text{s}$ . High power TPPM decoupling was applied ( $B_1 = 80$  kHz). The  $^{13}\text{C}$  chemical shifts are referenced using the glycine carboxyl group signal ( $\delta(^{13}\text{C}) = 176.4$  ppm).

$^{13}\text{C}$  CP-MAS NMR spectrum is a good evidence of the *in situ* functionalization of cyano groups, as specifically around 165 ppm two peaks appeared for the carbon atoms of the two different amide groups (Figure S3).

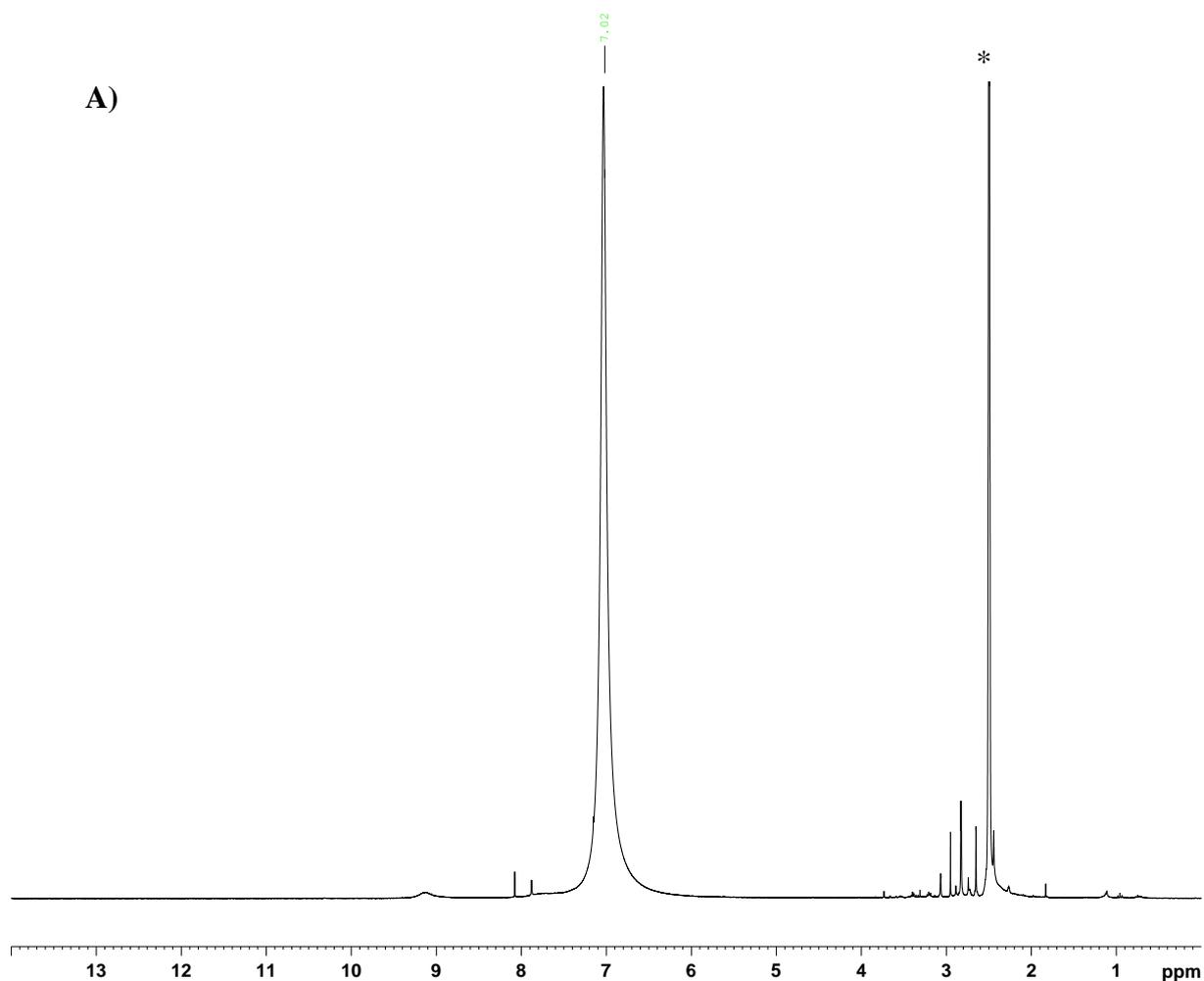


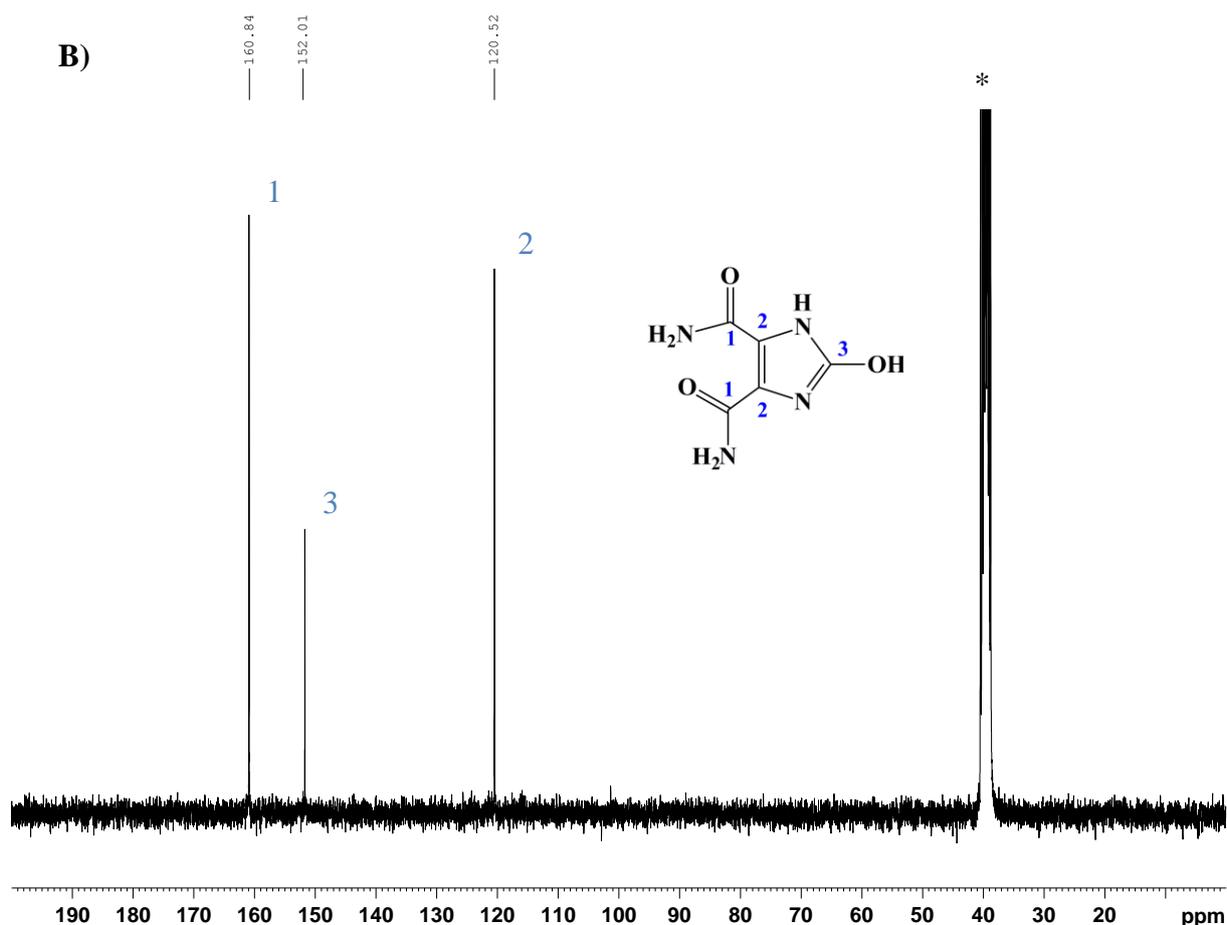
**Figure S3.** <sup>1</sup>H- (A) and <sup>13</sup>C- (B) CPMAS NMR spectra of as-synthesized **1**.

### Liquid-phase $^{13}\text{C}$ NMR spectroscopy of activated **1**

NMR spectra were recorded with a Bruker Avance 300 spectrometer. Resonances for NMR spectra are reported relative to  $\text{Me}_4\text{Si}$  ( $\delta = 0.0$  ppm) and calibrated based on the solvent signal for  $^1\text{H}$  and  $^{13}\text{C}$ .

Liquid-phase  $^{13}\text{C}$  NMR spectroscopy of activated **1** sample (26 mg) was digested in 0.5 mL  $[\text{D}_6]$  DMSO and 0.1 mL DCI (20%)/ $\text{D}_2\text{O}$ .





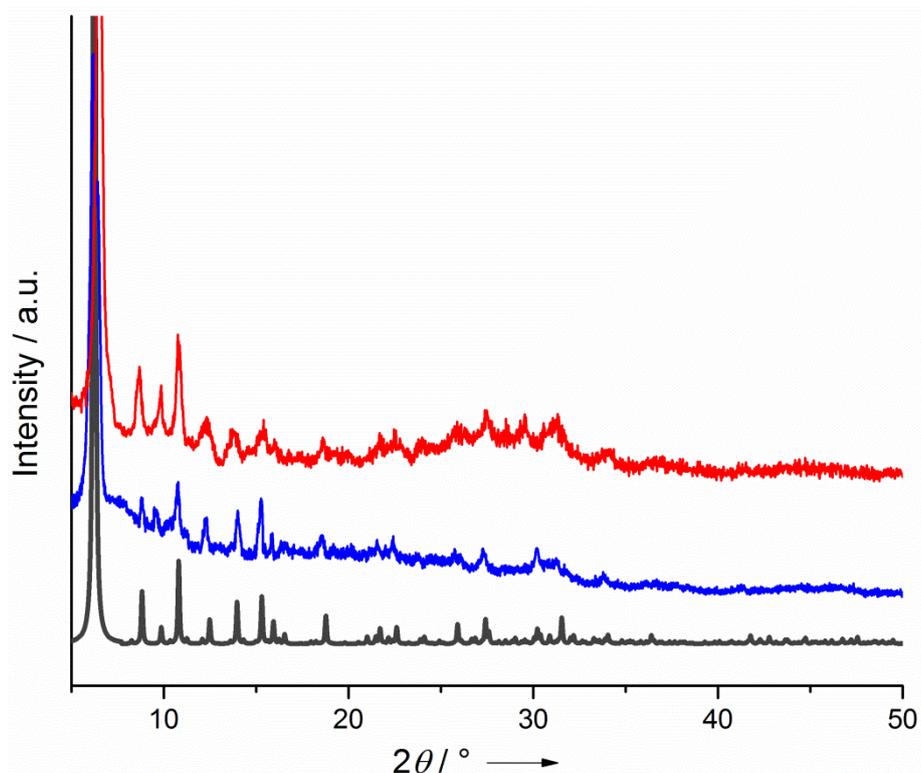
**Figure S4.**  $^1\text{H}$ - (A)  $^{13}\text{C}$ - (B) NMR spectra of a digested activated sample **1** in 0.1 mL DCl (20%)/D<sub>2</sub>O and 0.5 mL [D<sub>6</sub>] DMSO. Solvent signals ([D<sub>6</sub>] DMSO) are marked with an asterisk (\*). At  $^1\text{H}$  NMR spectra 2.6–3.0 ppm, weak triplet peaks are observed due to slow HDO intermolecular rate of exchange.<sup>4</sup>

Activated sample did not show any peaks, related to DMF at around 31, 36 and 162 ppm. Digested sample of activated **1** in DMSO-d<sub>6</sub> and DCl/D<sub>2</sub>O, there is no coordination to zinc. Moreover, olate group and imidazolate can be protonated. That could be a nice comparison with solid state NMR spectra (deprotonated and coordinated state).

## 6 Powder X-ray-diffraction patterns

Powder X-ray diffraction (PXRD) patterns were measured on a Siemens diffractometer D5005 in Bragg-Brentano reflection geometry. The diffractometer was equipped with a copper tube, a scintillation counter, automatic incident- and diffracted-beam soller slits and a graphite secondary monochromator. The generator was set to 40 kV and 40 mA. All measurements were performed with sample rotation. Data were collected digitally from 3° to 70°  $2\theta$  using a step size of 0.02°  $2\theta$  and a count time of 4 seconds per step. The simulated

powder pattern for **1** was calculated using single-crystal X-ray diffraction data and processed by the free Mercury v1.4.2 program provided by the Cambridge Crystallographic Data Centre.



**Figure S5.** Powder X-ray diffraction patterns of **1** (Color: gray = simulated; blue = as-synthesized, red = activated).

## 7 Single crystal X-ray structure determination

The crystal was embedded in perfluoropolyalkylether oil and mounted on a glass fibre. Intensity data were collected at 210 K using a STOE Imaging Plate Diffraction System IPDS-2 with graphite monochromatized  $\text{Mo}_{K\alpha}$  radiation ( $\lambda = 0.71073 \text{ \AA}$ ) at 50 kV and 40 mA (180 frames,  $\Delta\omega=1^\circ$ , exposure time per frame: 1 min). The data were corrected by a numerical absorption correction using the program X-Area (Stoe, 2004) as well as for Lorentz and polarisation effects. The structure was resolved with direct methods using SHELXS-97<sup>5</sup> and refined with full-matrix least-squares on  $F^2$  using the program SHELXL-2013<sup>6</sup> (Sheldrick, 1997). All non-hydrogen atoms were refined anisotropically.

The amide moieties show a high degree of vibrational motion, and attempts to model this behaviour to get better displacement parameters were not fully successful. The hydrogen atoms could not be located from the difference Fourier map and their positions had to be calculated using the AutoCN and HSite options of the Program TOPOS.<sup>7</sup> All hydrogen atoms were involved in the refinement, but their coordinates and temperature factors were fixed.

Therefore the hydrogen bonds can only be characterized by approximate donor-acceptor distances. Residual electron density ( $2.96 \text{ e} \text{ \AA}^3$ ) was found at a threefold axis which could not be assigned to an atom, because the distances to the next oxygen atoms are only  $1.8 \text{ \AA}$ . Even though the data collections were repeated with three other crystals these shortcomings could not rectify. The unit cell contains channels filled with disordered solvent molecules. In spite of several attempts, no chemically reasonable solution could be received for the solvent species in the channels of the crystal material. Very high displacement parameters, high estimates and partial occupancy due to the disorder make it impossible to determine accurate atomic positions for these molecules. Therefore the contribution of the disordered solvent species was subtracted from the structure factor calculations by the SQUEEZE instruction of PLATON. SQUEEZE calculated a solvent-accessible void volume in the unit cell of  $34081 \text{ \AA}^3$  (53 % of the total cell volume), corresponding to 11657 electrons (residual electron density after the last refinement cycle) per unit cell.<sup>8</sup> This number agrees with three molecules of dimethylformamide ( $3 \times 40 \times 96 = 11520$ ). The deposited atom data (*cif*) reflect only the refined cell content.

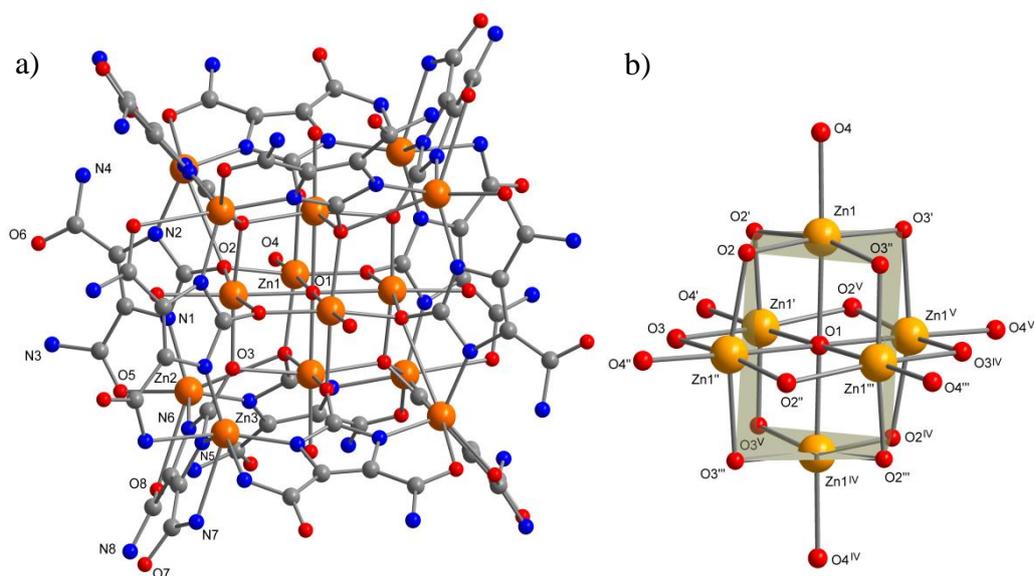
**Table S1.** Crystal Data, Details of Intensity Measurements, and Structure Refinement for  $[\text{Zn}_{14}(\text{L2})_{12}(\text{O})(\text{OH})_2(\text{H}_2\text{O})_4] (\text{DMF})_{18} (= \mathbf{1})$ .

Chemical formula, asymmetric unit	$\text{C}_{10}\text{H}_{9.67}\text{N}_8\text{O}_{7.17}\text{Zn}_{2.33}$
Formula mass	515.80
Crystal system	cubic
Space group	$Ia\bar{3}d$ (No 230)
$a = b = c / \text{\AA}$	40.1873(12)
$\alpha = \beta = \gamma / ^\circ$	90
Unit cell volume / $\text{\AA}^3$	64903(6)
Temperature / K	210(2)
Crystal size / mm	0.50 x 0.45 x 0.40
Z	96
Density (calculated) / $\text{g}\cdot\text{cm}^{-3}$	1.25
Radiation type	MoK $\alpha$
$\mu / \text{mm}^{-1}$	2.10
Theta range / $^\circ$	1.24-25.00
Index ranges	$-23 \leq h \leq 46$ $-37 \leq k \leq 47$ $-47 \leq l \leq 31$
Reflections collected	31851
Independent reflections	4761
$R_{\text{int}}$	0.0653
$R_1 / wR_2 [I > 2\sigma(I)]$	0.0586 / 0.1540
$R_1 / wR_2$ (all data)	0.0870 / 0.1652
Goodness of fit on $F^2$	0.861
Max. diff. peak and hole / $e\cdot\text{\AA}^{-3}$	2.86 / -0.89

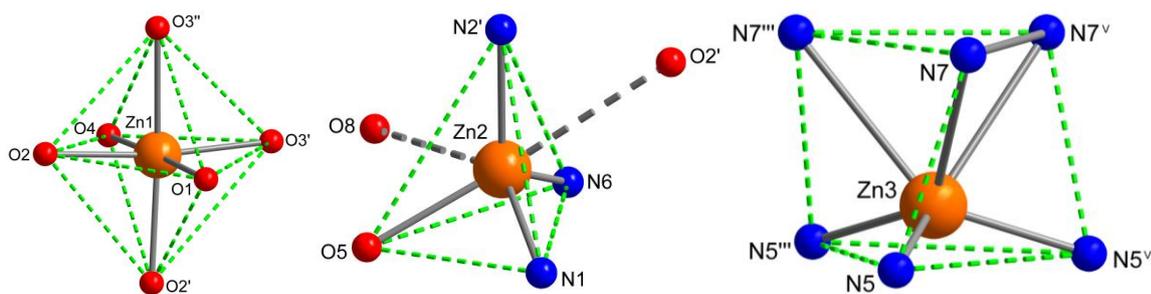
### Details of structural description:

The asymmetric unit contains three different zinc centres (Zn1, Zn2, Zn3), two imidazolate-4,5-diamide-2-olate ligands (L2) and moreover  $\text{O}^{2-}$ ,  $\text{OH}^-$  and  $\text{H}_2\text{O}$ . The MBB consists of 14 zinc atoms (six Zn1, six Zn2, two Zn3) and twelve L2 ligands, one  $\text{O}^{2-}$  (O1), two  $\text{OH}^-$  (O4) anions and four  $\text{H}_2\text{O}$  (O4) molecules (Figure S6a). The oxide ion (O1) is located in the centre of the MBB, surrounded by six Zn1 atoms in an exact octahedral coordination environment (Figure S6b). Each of these six Zn1 atoms is further coordinated by four olate oxygen atoms of four imidazolate ligands (two O2 and two O3) and one O4 forming a octahedral coordination geometry (Figure S7). For electroneutrality we must state that 1/3 of O4 is hydroxide and 2/3 is water. Each of the olate oxygen atoms (O2 and O3) is coordinated to two Zn1 centres; in addition O2 is coordinated to the Zn2 atom. The six Zn2 centres are

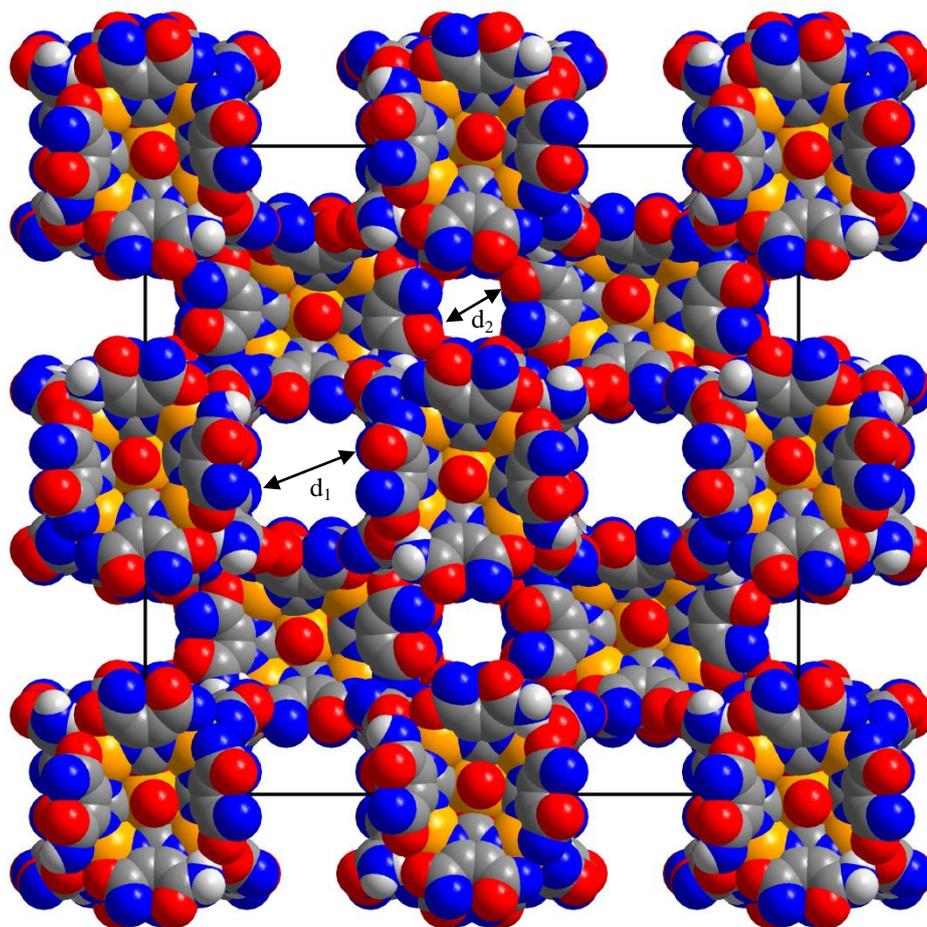
surrounded by three imidazolate nitrogen atoms: N1, N2, N6, and three oxygen atoms: O5, O8 of amide moieties and O2 (olate oxygen atom) to form a twofold face-capped tetrahedron (Figure S7). Both Zn3 atoms are coordinated by six nitrogen atoms from three imidazolate nitrogen atoms (N5 and symmetry related atoms) and three amide nitrogen atoms N7 in a distorted trigonal prismatic fashion (Figure S7).



**Figure S6.** a) Tetradecanuclear zinc the MBB with the three zinc centres (Zn1, Zn2 and Zn3) of **1**, for symmetry operators see legend of Table S4; b) central core of MBB (hydrogen atoms were omitted for clarity; for symmetry code - see legend of Table S4).



**Figure S7.** The three zinc centres (Zn1, Zn2 and Zn3) of **1** in the MBB, for symmetry operators see legend of Table S4.



**Figure S8.** Space filling model for **1**, having two types of channels (approximate diameter,  $d_1 = 6.0 \text{ \AA}$  and  $d_2 = 3.9 \text{ \AA}$ , orange Zn, blue N, red O, dark gray C, light gray H).

**Table S2.** Donor-Acceptor distances of the hydrogen -bonds [ $\text{\AA}$ ] for **1**.

	D...A
N3 - H3A...O6	2.664(9)
N8 - H8B...O7	2.74(2)
N3 - H3B...O6 <sup>VII</sup>	2.790(9)
N4 - H4A...O5 <sup>II</sup>	3.26(1)
N4 - H4A...O8 <sup>II</sup>	2.87(1)
N4 - H4B...O5 <sup>VI</sup>	3.006(8)
N4 - H4B...O8 <sup>VI</sup>	3.21(1)

Symmetry codes - see legend of Table S4.

**Table S3.** Selected bond lengths [ $\text{\AA}$ ] for **1**.

Zn1 - O1 <sup>II</sup>	2.1914(7)
Zn1 - O2	2.027(4)
Zn1 - O3 <sup>II</sup>	1.986(5)
Zn1 - O4	2.094(6)
Zn2 - O5	2.081(4)
Zn2 - N1	2.013(5)
Zn2 - N2 <sup>I</sup>	1.996(5)
Zn2 - N6	1.968(5)
Zn3 - N5 <sup>III</sup>	1.981(5)
Zn3 - N7 <sup>V</sup>	2.512(1)

Zn2...O2<sup>I</sup>: 2.819(7)  $\text{\AA}$ ; Zn2...O8: 2.730(7)  $\text{\AA}$

**Table S4.** Selected bond angles [ $^\circ$ ] for **1**.

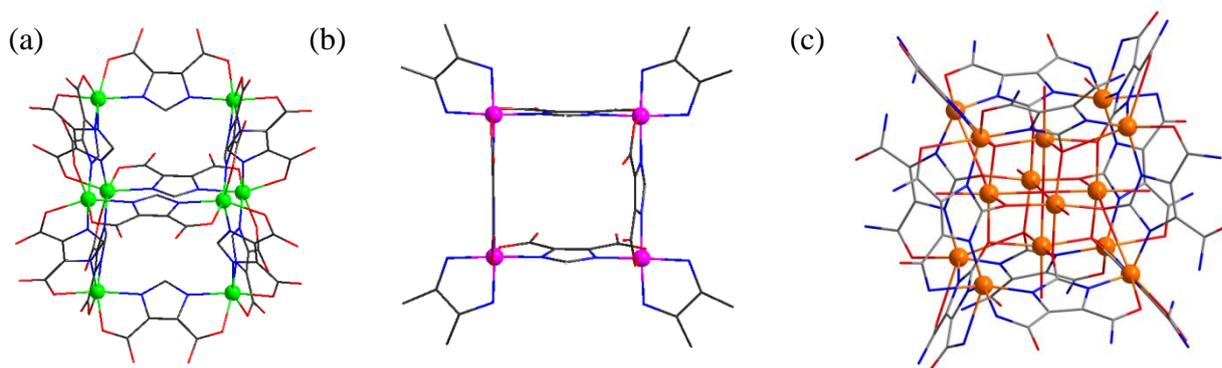
Zn1 - O1 - Zn1 <sup>II</sup>	180.00(4)
Zn1 - O1 - Zn1 <sup>II</sup>	89.72(3)
O3 - Zn1 - O2 <sup>I</sup>	171.0(2)
O3 - Zn1 - O4 <sup>I</sup>	92.1(2)
O2 - Zn1 - O4	92.6(2)
O3 - Zn1 - O1 <sup>I</sup>	89.20(1)
O2 - Zn1 - O1	86.4(1)
O4 - Zn1 - O1	177.65(2)
N6 - Zn2 - N2 <sup>I</sup>	115.2(2)
N6 - Zn2 - N1	114.7(2)
N2 - Zn2 - N1 <sup>I</sup>	113.0(2)
N6 - Zn2 - O5	115.4(2)
N2 - Zn2 - O5 <sup>I</sup>	114.8(2)
N1 - Zn2 - O5	78.83(2)
N5 - Zn3 - N5 <sup>III</sup>	111.40(1)
N5 - Zn3 - N7 <sup>III</sup>	69.6(3)
N7 - Zn3 - N7 <sup>III</sup>	72.4(5)

Symmetry Operators: <sup>I</sup>0.5-z,0.5+x,y-1 <sup>II</sup>-0.5+y,1+z,0.5-x <sup>III</sup>0.5+z,1.5-x,1-y <sup>IV</sup>1-x,2-y,-z  
<sup>V</sup>1.5-y,1-z,x-0.5 <sup>VI</sup>0.75-x,1.25-z,-0.75+y <sup>VII</sup>0.75-x,0.75+z,1.25-y





is to fully hydrolyzed to carboxylate group under such condition. Additionally, *in situ* hydrolysis of the methoxy to the hydroxy group allows the linker to chelate the metal ion by locking it into its position through the formation for Zn<sub>14</sub> metal nodes, compared with others carboxylate based linkers. Moreover, no capping ligand was added to the reaction mixture because free amides of MBB which are directed to vertices are involved with other MBBs via hydrogen-bonds. Different types of nets of various clusters are presented in Table S5.



**Figure S9.** Schematic presentations of (a) In<sub>8</sub> nodes based metal organic cube (MOC)<sup>10a</sup> (b) Co<sub>4</sub> nodes based metal organic square (MOS),<sup>11</sup> formed from 4,5-imidazoledicarboxylate linkers. Each building block is connected through its vertices or edges with other building blocks by intermolecular O–H···O hydrogen bonds, generating supramolecular assembly, (c) tetradecanuclear zinc MBB for **1**.

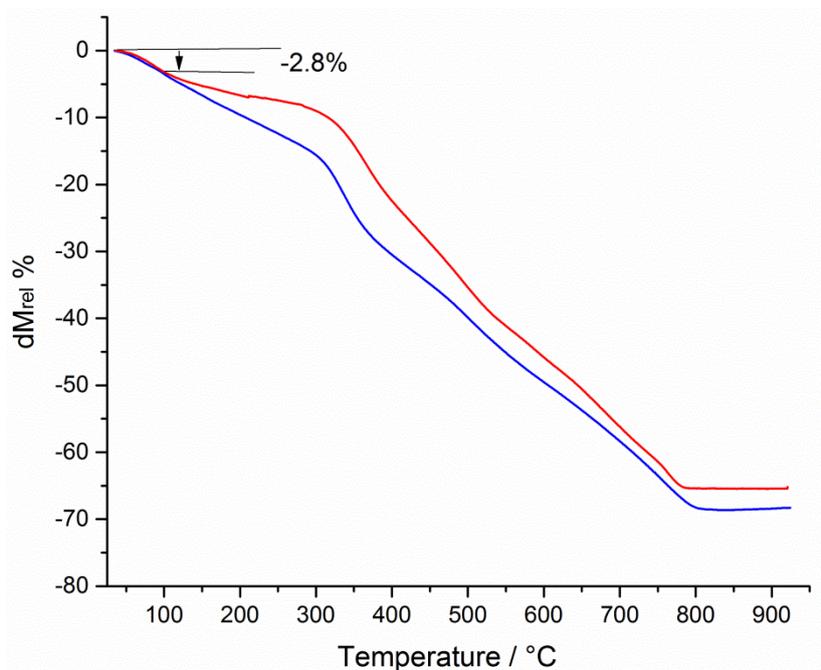
**Table S5.** Comparison of various nodes of different cluster based networks.

Serial No.	Linker name	Nodes	Topology/ nets	Reference
1	imidazolate-4,5-diamide-2-olate	Zn <sub>14</sub>	bcu	this study
2	4,5-imidazoledicarboxylic acid	In <sub>8</sub>	zeolite ACO	<i>J. Am. Chem. Soc.</i> <b>2009</b> , <i>131</i> , 10394.
3	4,5-imidazoledicarboxylic acid	In <sub>8</sub>	zeolite AST	<i>J. Am. Chem. Soc.</i> <b>2009</b> , <i>131</i> , 10394.
4	4,5-imidazoledicarboxylic acid	Co <sub>4</sub>	zeolite gis	<i>J. Am. Chem. Soc.</i> <b>2010</b> , <i>132</i> , 18038.
5	1,2,3-triazole-4,5-dicarboxylic acid	In <sub>4</sub>	zeolite rho	<i>J. Am. Chem. Soc.</i> <b>2010</b> , <i>132</i> , 18038.
6	4,5-imidazoledicarboxylic acid	Zn <sub>8</sub>	zeolite AST	<i>J. Am. Chem. Soc.</i> <b>2009</b> , <i>131</i> , 17753.
7	4,5-imidazoledicarboxylic acid	Cd <sub>8</sub>	zeolite LTA	<i>J. Am. Chem. Soc.</i> <b>2009</b> , <i>131</i> , 17753.
8	4,6-pyrimidicarboxylate	InN <sub>4</sub>	sodalite topology	<i>J. Am. Chem. Soc.</i> <b>2008</b> , <i>130</i> , 3768.
9	2-pyrimidinecarboxylate	CdN <sub>4</sub>	rho zeolite net	<i>J. Am. Chem. Soc.</i> <b>2008</b> , <i>130</i> , 3768.
10	4,5-imidazoledicarboxylic acid	InN <sub>4</sub>	zeolite rho topology	<i>Chem. Commun.</i> <b>2006</b> , 1488.
11	5-tetrazolylisophthalic acid	Cu <sub>12</sub>	(3,24)-connected rht net	<i>J. Am. Chem. Soc.</i> <b>2008</b> , <i>130</i> , 1833.
12	3,5-dicarboxyl-(3',5'-dicarboxylazophenyl) benzene	Ni	12-connected fcu net10	<i>J. Am. Chem. Soc.</i> <b>2008</b> , <i>130</i> , 1560.
13	H4BIPA-TC	Co	NbO -like	<i>J. Am. Chem. Soc.</i> <b>2008</b> , <i>130</i> , 1560.
14	1H-imidazole-4,5-dicarboxylic acid	Cd <sub>4</sub>	4·8 <sup>2</sup> fes	<i>Cryst. Growth Des.</i> <b>2012</b> , <i>12</i> , 1697.
15	2,5-pyridinedicarboxylic acids	4-connected node, <i>cis</i> -InN <sub>2</sub> (CO <sub>2</sub> ) <sub>2</sub>	Kagomé lattice topology	<i>J. Am. Chem. Soc.</i> <b>2005</b> , <i>127</i> , 7266.

## 9 Thermogravimetric (TG) analysis

The TG measurements were performed in a stationary air atmosphere (no purge) from room temperature up to 900 °C using a Linseis thermal analyzer (Linseis, Germany) working in vertical mode. The heating rate was 10 °C/min. The samples were placed in cups of aluminium oxide.

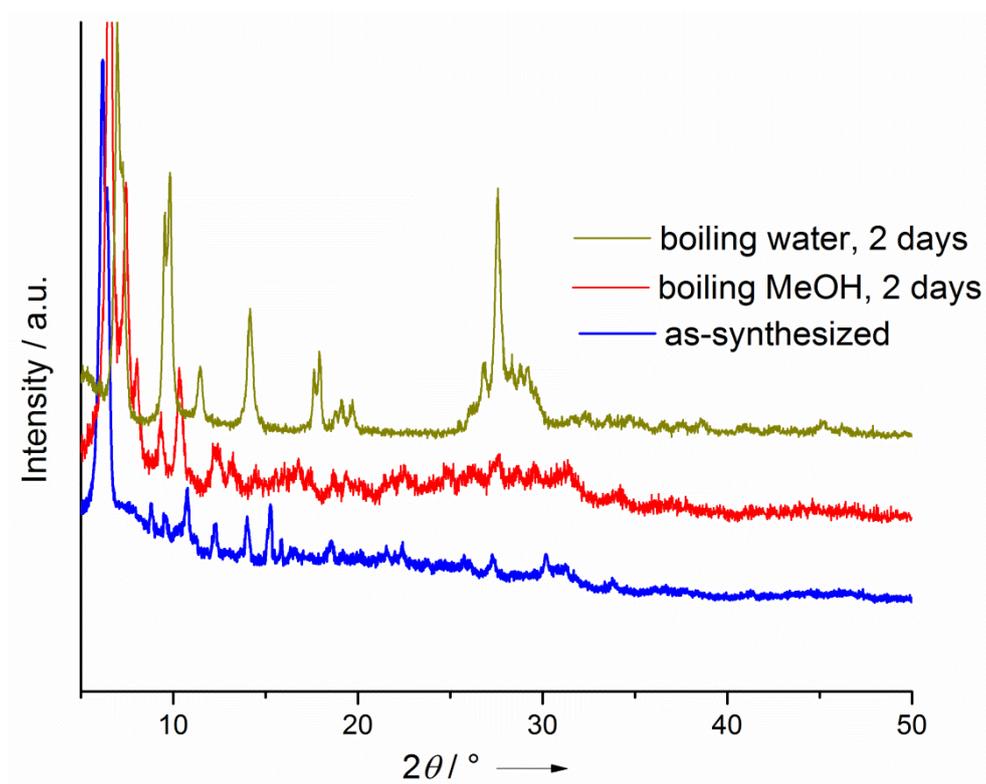
The as-synthesized sample of **1** exhibited a continuous weight loss upon heating. Activated sample showed weight loss of 2.8 % (calculated for loss of 4 H<sub>2</sub>O 2.3 %) at 25–105 °C, corresponding to the release of coordinated water molecules (aqua ligands). No stable plateau was observed.



**Figure S10.** TGA curves for **1** (color: blue = as-synthesized, red = activated).

## 10 Chemical Stability

Material is stable at ambient air and moisture. Moreover, the chemical stability of the material was analyzed by suspending the samples for 2 days in boiling methanol and water. After such extensive treatment, the material is irreversibly transformed to a so-far unknown crystalline phase (Figure S11). Hence, it can be concluded that the non-rigid H-bonded framework is solvothermally unstable above 65 °C.



**Figure S11.** Powder X-ray diffraction profiles of **1** collected during the chemical stability tests in refluxing methanol and water.

## 10 Gas-sorption measurements

The sample was connected to the preparation port of the sorption analyzer and degassed under vacuum until the out gassing rate, i.e., the rate of pressure rise in the temporarily closed manifold with the connected sample tube, was less than 2  $\mu$ Torr/min at the specified temperature 50 °C for 24 h. After weighing, the sample tube was then transferred to the analysis port of the sorption analyzer. All used gases ( $H_2$ , He,  $N_2$ ,  $CO_2$ ,  $CH_4$ ) were of ultra high purity (UHP, grade 5.0, 99.999%) and the STP volumes are given according to the NIST standards (293.15 K, 101.325 kPa). Helium gas was used for the determination of the cold and warm free space of the sample tubes.  $H_2$  and  $N_2$  sorption isotherms were measured at 77 K (liquid nitrogen bath), whereas  $CO_2$  and  $CH_4$  sorption isotherms were measured at  $298\pm 1$  K (passive thermostating) and 273.15 K (ice/deionized water bath). The heat of adsorption values and the DFT calculations (' $N_2$  DFT slit pore' model) were done using the ASAP 2020 v3.05 software.

**Table S6.**  $N_2$  uptake at 77 K and 1 bar for three consecutive runs on the same probe.

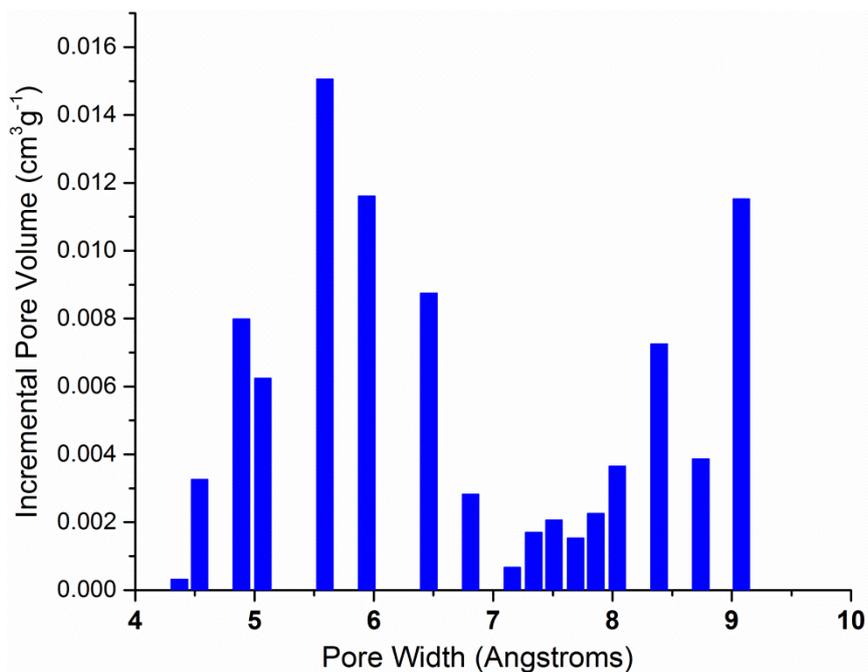
	$S_{BET}$ [ $m^2g^{-1}$ ]	$S_{Langmuir}$ [ $m^2g^{-1}$ ]
1 <sup>st</sup> run	465	565
2 <sup>nd</sup> run	471	570
3 <sup>rd</sup> run	468	567

The hydrogen-bonding interactions at **1** can be ruptured to some extent upon activation, giving even smaller particles or cracks in the crystals. Moreover, PXRD pattern of activated sample showed slight broad peaks. Therefore, we repeated the  $N_2$  sorption measurement two times on the same sample and confirmed the reproducibility of the results (Table S6). Hence, we confirmed that non-rigid H-bonded framework is still active for gas-sorption under such activated condition.

**Table S7.** Comparison of characteristic sorption data of **1** with selected azolate-based hydrogen-bonded supramolecular frameworks (MOC-3 and ZSA-2), hydrogen-bonded porous organic compounds (SOF-1a, HOF-1a, CBDU and TTEB) and ZIFs.

	$SA_{\text{BET}}^2$ [m <sup>2</sup> g <sup>-1</sup> ]	$SA_{\text{Langmuir}}^2$ [m <sup>2</sup> g <sup>-1</sup> ]	V (N <sub>2</sub> ) [cm <sup>3</sup> g <sup>-1</sup> ]	V (CO <sub>2</sub> ) [cm <sup>3</sup> g <sup>-1</sup> ]	wt % (H <sub>2</sub> )	Ref.
<b>1</b>	<b>471</b> <sup>[a]</sup>	<b>570</b> <sup>[a]</sup>	<b>172</b> <sup>[b]</sup>	<b>56</b> <sup>[c]</sup> / <b>38</b> <sup>[d]</sup>	<b>0.85</b> <sup>[b]</sup>	this study
MOC-3	— <sup>[e]</sup>	456	165 <sup>[f]</sup>	— <sup>[d]</sup>	0.75 <sup>[f]</sup>	<i>J. Am. Chem. Soc.</i> <b>2009</b> , <i>131</i> , 10394.
ZSA-2	395	— <sup>[e]</sup>	120 <sup>[l]</sup>	— <sup>[e]</sup>	0.35 <sup>[g]</sup> / 0.95 <sup>[h]</sup>	<i>J. Am. Chem. Soc.</i> <b>2010</b> , <i>132</i> , 18038.
SOF-1a	474 <sup>[i]</sup>	639 <sup>[i]</sup>	143 <sup>[i]</sup>	16 <sup>[d]</sup>	— <sup>[e]</sup>	<i>J. Am. Chem. Soc.</i> <b>2010</b> , <i>132</i> , 14457.
HOF-1a	359 <sup>[j,k]</sup>	609 <sup>[k]</sup>	— <sup>[e]</sup>	225 <sup>[l]</sup>	— <sup>[e]</sup>	<i>J. Am. Chem. Soc.</i> <b>2011</b> , <i>133</i> , 14570.
CBDU	341 <sup>[j]</sup>	— <sup>[e]</sup>	— <sup>[e]</sup>	72 <sup>[m]</sup>	— <sup>[e]</sup>	<i>Chem. Mater.</i> <b>2006</b> , <i>18</i> , 4855.
TTEB	278 <sup>[a]</sup>	— <sup>[e]</sup>	— <sup>[e]</sup>	— <sup>[e]</sup>	3.9 <sup>[n]</sup>	<i>Angew. Chem. Int. Ed.</i> <b>2009</b> , <i>48</i> , 3273.
TTP	— <sup>[d]</sup>	240 <sup>[a]</sup>	— <sup>[d]</sup>	22 <sup>[d]</sup>	—	<i>Angew. Chem. Int. Ed.</i> <b>2005</b> , <i>44</i> , 1816; <i>Microporous Mesoporous Mater.</i> <b>2006</b> , <i>88</i> , 170.
ZIF-95	1050	—	—	19.7 <sup>[d]</sup>	—	<i>Nature</i> <b>2008</b> , <i>453</i> , 207; <i>Acc. Chem. Res.</i> <b>2010</b> , <i>43</i> , 58.
ZIF-100	595	—	—	32.6 <sup>[c]</sup>	—	<i>Nature</i> <b>2008</b> , <i>453</i> , 207; <i>Acc. Chem. Res.</i> <b>2010</b> , <i>43</i> , 58.
ZIF-81	760	—	—	38.2 <sup>[d]</sup>	—	<i>J. Am. Chem. Soc.</i> <b>2009</b> , <i>131</i> , 3875.
ZIF-79	810	—	—	33.5 <sup>[d]</sup>	—	<i>J. Am. Chem. Soc.</i> <b>2009</b> , <i>131</i> , 3875.
ZIF-69	950	—	—	40.6 <sup>[d]</sup>	—	<i>J. Am. Chem. Soc.</i> <b>2009</b> , <i>131</i> , 3875.
ZIF-68	1090	—	—	37.6 <sup>[d]</sup>	—	<i>J. Am. Chem. Soc.</i> <b>2009</b> , <i>131</i> , 3875.

[a] Determined by N<sub>2</sub> sorption at 77 K with data points in the range for  $p/p_0$  between 0.01 and 0.05 for determining BET surface area and between 0.01 and 0.15 for the Langmuir surface area. [b] Measured at 77 K and  $p/p_0 \approx 0.97$ . [c] Measured at 273 K and 1 bar. [d] Measured at 298 K and 1 bar. [e] Not reported. [f] Approx. value was taken from SI. [g] Measured at 298 K and 80 bar. [h] Measured at 77 K and 20 bar. [i] Calculated from N<sub>2</sub> adsorption at 125 K and 1 bar. [j] Determined by CO<sub>2</sub> adsorption at 196 K. [k] Data points  $p/p_0$  between 0.1 and 0.3 and BET C-constant is negative. [l] At 196 K and 1 bar. [m] 196 K, 1 bar at  $p/p_0 = 0.5$ . [n] At 77 K and 10 bar.

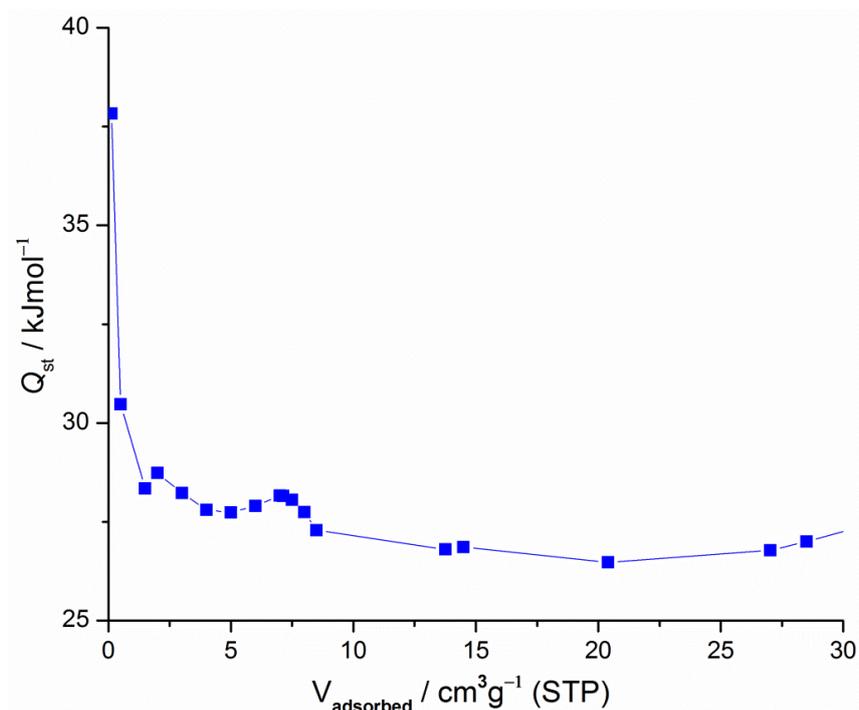


**Figure S12.** The pore size distribution from a CO<sub>2</sub> adsorption isotherm at 273 K, calculated by the NL-DFT with a “CO<sub>2</sub> on carbon based slit-pore” model at 273 K.

### Heat of adsorption

From two adsorption isotherms acquired at different temperatures  $T_1$  and  $T_2$ , the differential heat of adsorption  $\Delta H_{ads,diff}$ , that is isosteric heat of adsorption  $Q_{st}$  can be calculated for any amount of adsorbed substance after determining the required relative pressures  $p_1$  and  $p_2$ . A modified form of the Clausius-Clapeyron equation is used (eq (1))<sup>12</sup>  $\Delta H_{ads,diff}$  was calculated over the whole adsorption range from the 273 K and 298 K isotherms for CO<sub>2</sub>.

$$\Delta H_{ads,diff} = Q_{st} = -R \ln \left( \frac{p_2}{p_1} \right) \frac{T_1 T_2}{T_2 - T_1} \quad (1)$$



**Figure S13.** Isosteric heats of CO<sub>2</sub> adsorption as a function of the adsorbent loading for **1**.

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