

Supporting Information for:

NMR-Enhanced Crystallography Aids Open Metal-Organic Framework Discovery Using Solvent-Free Accelerated Aging

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Table S1: Optimized recycle delays for ^1H MAS experiments

Compound	Recycle Delay (s)
1	5
2	5
3	5
4	7.5

Table S2: ^1H - ^{13}C CP/MAS contact times and recycle delays

Compound	Contact Time (ms)	Recycle Delay (s)
1	1	5
2	10	5
3	2	5
4	2	7.5
HMeIm	15	60

Table S3: ^1H - ^{111}Cd CP contact times and recycle delays

Compound	Contact Time (ms)	Recycle Delay (s)
1	8	5
2	17	5
3	7	5
4	5	7.5

Table S4: ^1H - ^{14}N BRAIN-CP/WURST-CPMG experimental parameters

Compound	Recycle Delay (s)	Contact Time (ms)	# of echoes	# of subspectra
1	1	20	40	15
HMeIm	30	20	80	26

Table S5: Selected bond lengths in the Cd-imidazole compounds with known structures

Compound	Bond	Length (Å)
2	Cd-N1	2.215
	Cd-N2	2.228
	Cd-N3	2.260
	Cd-N4	2.226
3	Cd1-N2	2.203
	Cd1-N3	2.198
	Cd1-N5	2.199
	Cd1-N6	2.258
	Cd1-N7	2.167
	Cd2-N1	2.201
	Cd2-N2	2.209
4	Cd-N1	2.401

Table S6: Selected bond angles in the Cd-imidazole compounds with known structures

Compound	Bond	Angle (°)
2	N1-Cd-N2	113.5
	N1-Cd-N3	115.2
	N1-Cd-N4	112.2
	N2-Cd-N3	96.66
	N2-Cd-N4	112.9
	N3-Cd-N4	105.3
3	N2-Cd1-N3	115.3
	N2-Cd1-N5	115.6
	N2-Cd1-N6	80.97
	N2-Cd1-N7	107.4
	N3-Cd1-N5	109.9
	N3-Cd1-N6	126.1
	N3-Cd1-N7	102.0
	N5-Cd1-N6	106.5
	N5-Cd1-N7	105.1
	N6-Cd1-N7	29.33
	N1-Cd2-N2	106.8
	N1-Cd2-N2	117.4
4	N1-Cd-N1	89.769
	N1-Cd-N1	90.231

2. Solid-state NMR data

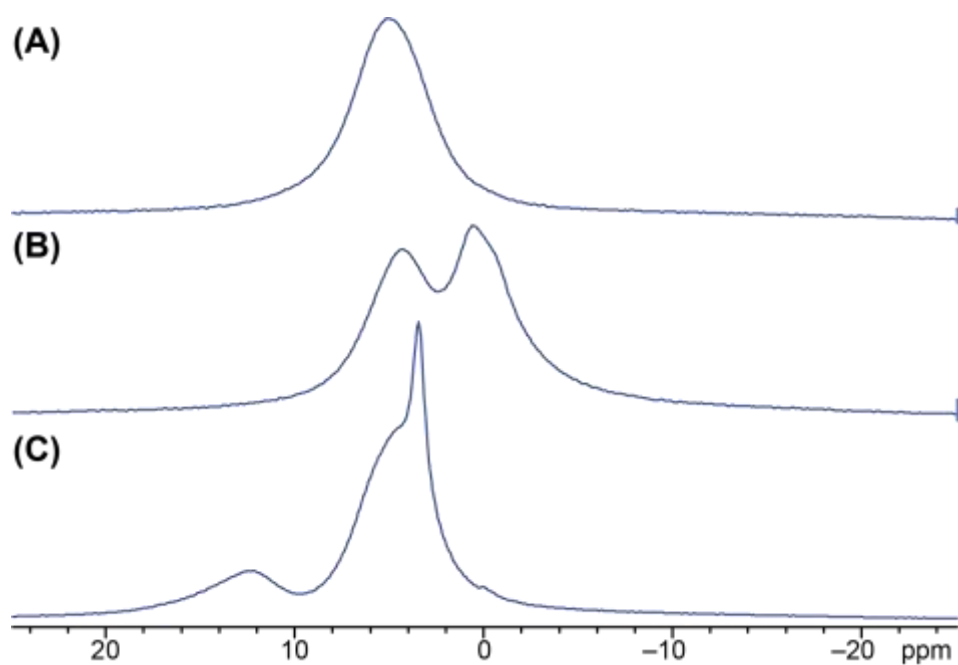


Figure S1: ^1H MAS ($\nu_{\text{rot}} = 16$ kHz) NMR spectra acquired at 9.4 T for (A) framework 2, (B) framework 3, and (C) compound 4.

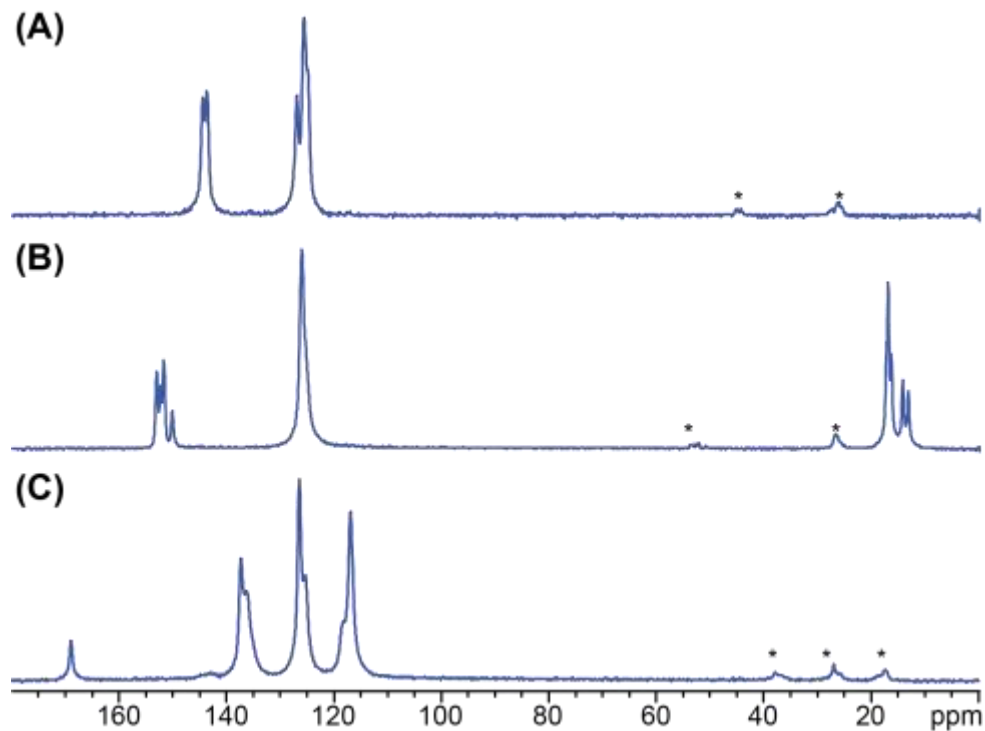


Figure S2: ^1H - ^{13}C CP/MAS ($\nu_{\text{rot}} = 10$ kHz) NMR spectra acquired at 9.4 T of (A) framework 2, (B) framework 3, and (C) compound 4. The asterisks (*) denoted spinning sidebands.

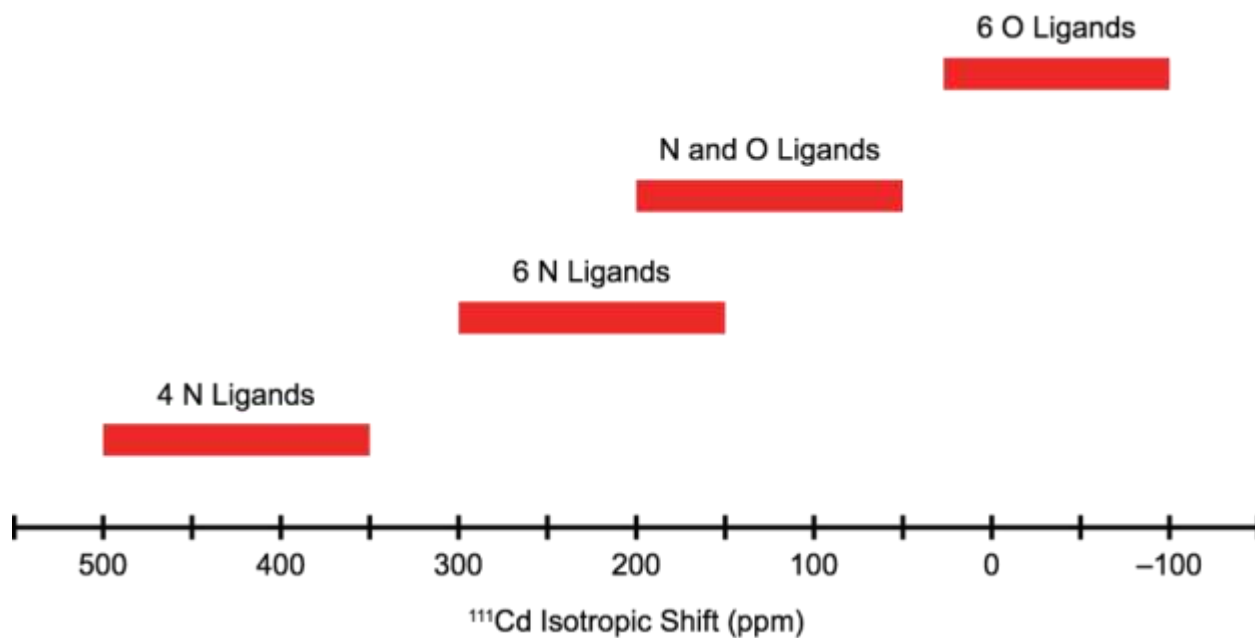


Figure S3: Proposed ^{111}Cd isotropic chemical shift ranges based on the compounds investigated herein and on studies conducted by Ellis *et al.*^[17-24] and Baxter *et al.*^[25]

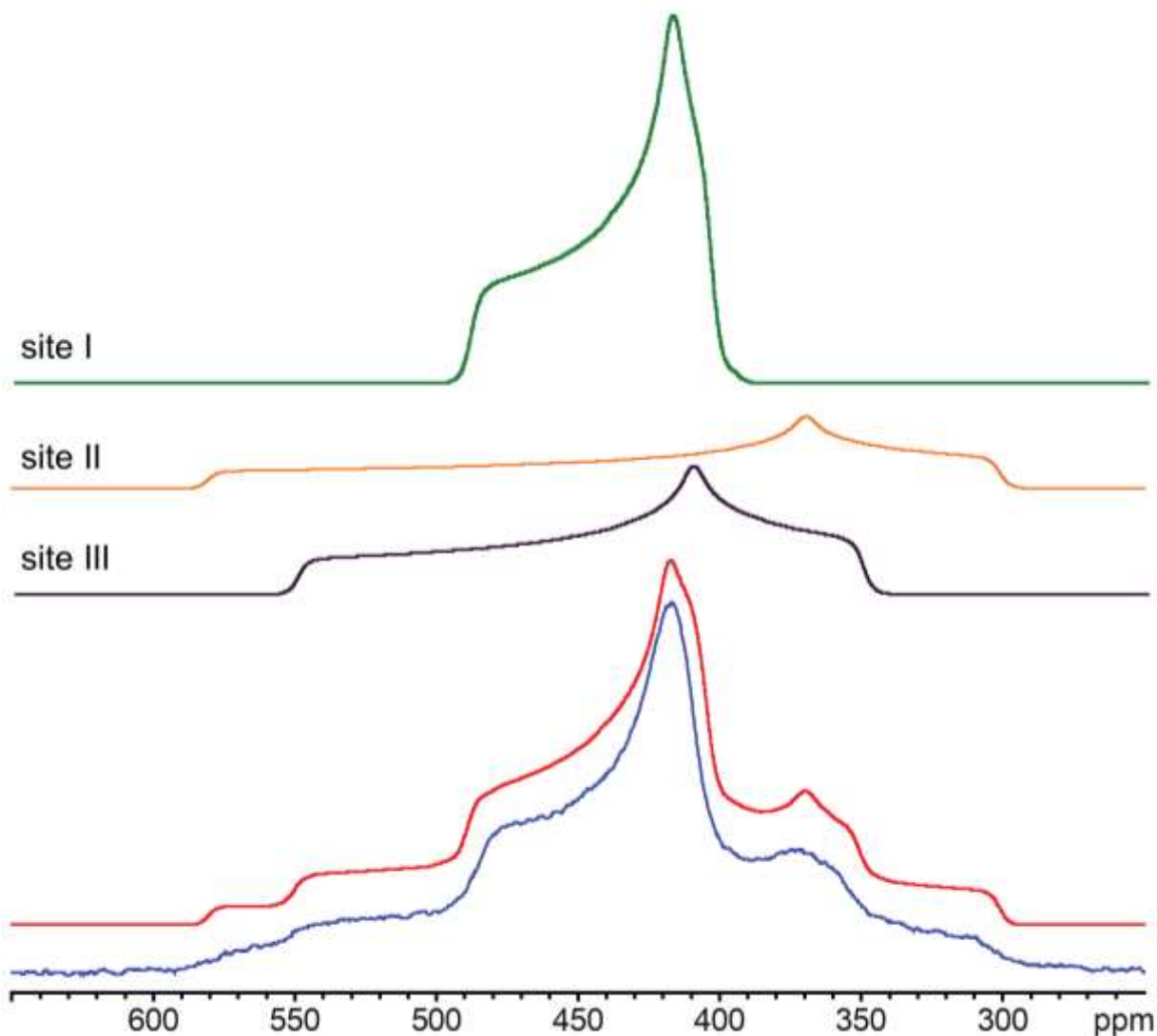


Figure S4: Experimental ^1H - ^{111}Cd CP static spectrum of **2** (blue trace). Simulation of the entire spectrum is shown in the red trace. The simulation is deconvoluted into the individual sites: site I (green trace), site II (orange trace), and site III (black trace). The corresponding ^{111}Cd CS tensor parameters can be found in **Table 1** in the main text.

¹H-¹⁴N BRAIN-CP/WURST-CPMG

The ¹H-¹⁴N static CP NMR spectra for **HMeIm** and **1** are shown in **Figure S5**. Due to the enormous breadth (*ca.* 4.5 MHz) of ¹⁴N patterns, it was necessary to use the broadband adiabatic inversion CP (BRAIN-CP) for polarization transfer,^[3,4] and the wideband uniform-rate smooth-truncation-CPMG (WURST-CPMG) pulse sequence for broadband refocusing.^[5,6,26] The spectrum of bulk **HMeIm** reveals two overlapping Pake-like doublet patterns, corresponding to two distinct nitrogen environments (**Figure S6**). The narrower pattern ($C_Q = 1.95(5)$ MHz, $\eta_Q = 0.44(2)$) is assigned to the nitrogen of the N-H group and the broader pattern ($C_Q = 3.13(5)$, $\eta_Q = 0.21(2)$) to the non-protonated nitrogen atom. The spectrum of **1** exhibits several broad features resulting from multiple overlapping Pake doublet patterns. Some of the features (*i.e.*, horns, shoulders, and feet) in the spectrum of **1** occur at the same frequency as those for bulk **HMeIm**, consistent with **HMeIm** inclusion in the MOF. Furthermore, discontinuities in the spectrum of **1** are not as well defined as those for bulk **HMeIm**, suggesting that guest molecules are disordered in the former. Finally, there are discontinuities in the ¹⁴N spectrum of **1** that do not match those in the spectrum of **HMeIm**; these are assigned to nitrogen atoms of **MeIm**⁻ ligands forming the framework of **1**.

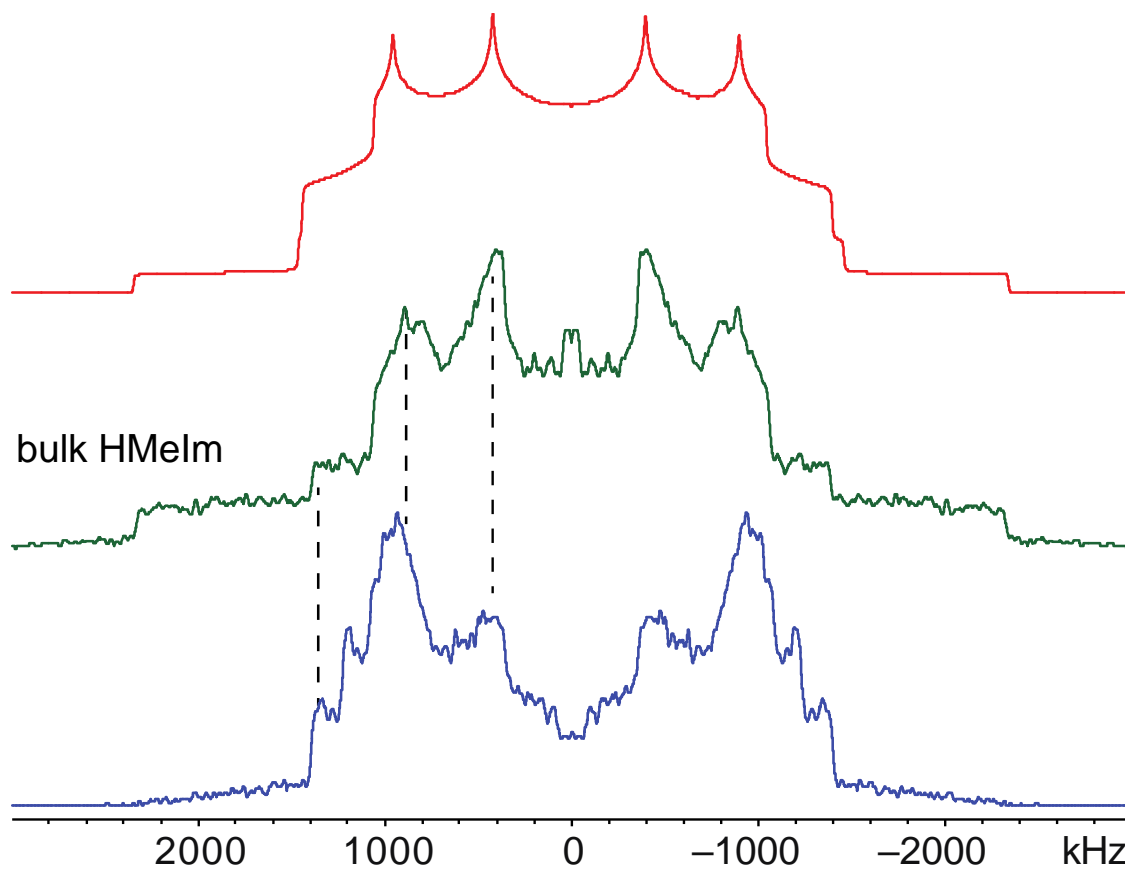


Figure S5: ^1H - ^{14}N BRAIN-CP/WURST-CPMG spectra of HMeIm (green trace) and **1** (blue trace).

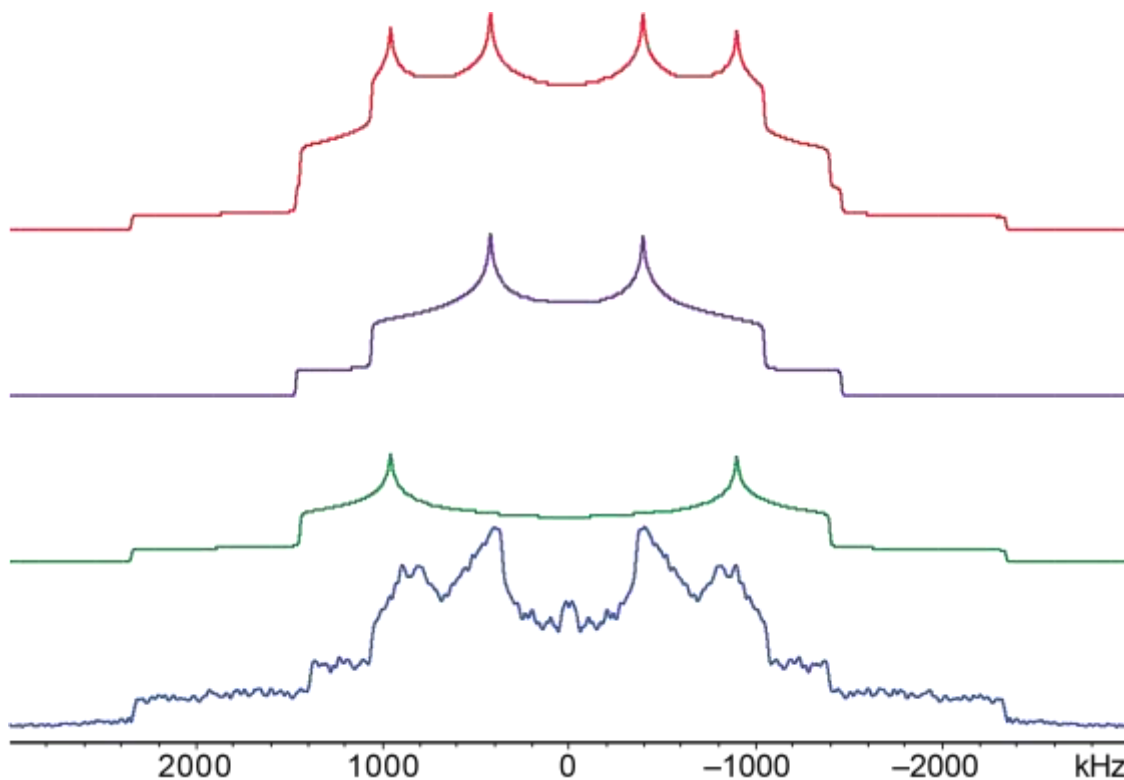


Figure S6: ^1H - ^{14}N BRAIN-CP/WURST-CPMG spectrum of bulk **HMeIm** (blue trace) and simulation of the powder pattern (red trace). A deconvolution of the simulation into the two distinct powder patterns is also shown. The purple trace is the pattern corresponding to the N1 site ($C_Q = 1.95(5)$ MHz, $\eta_Q = 0.44(2)$) and the green trace is the pattern corresponding to the N2 site ($C_Q = 3.13(5)$ MHz, $\eta_Q = 0.21(2)$).

3. Powder X-ray diffraction data

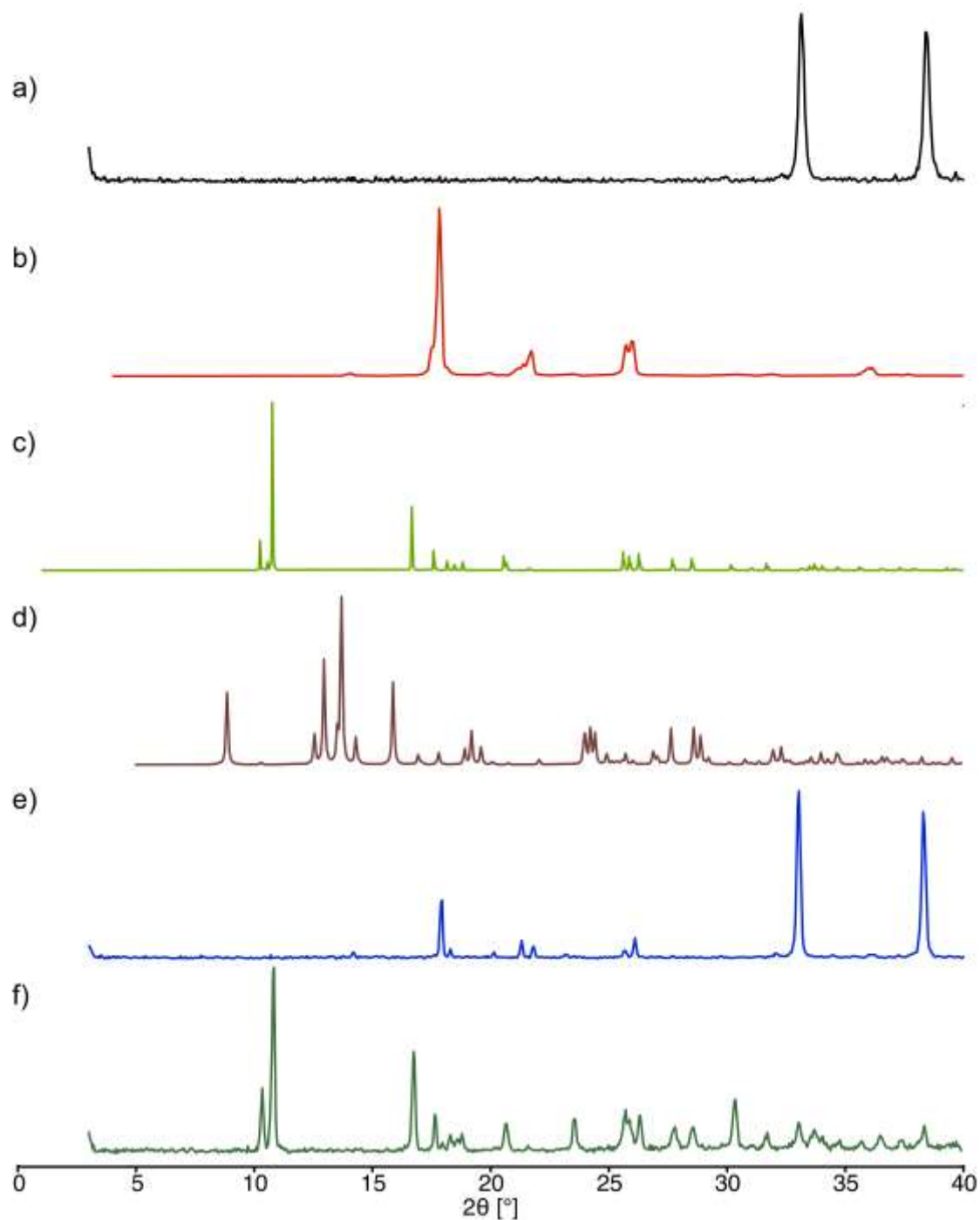


Figure S7: Experimental PXRD patterns of: a) commercial CdO; b) commercial HMeIm; c) simulated PXRD pattern for herein reported compound **1**; d) simulated PXRD pattern of **yqt1**-Cd(MeIm)₂ (CSD code GUPBOJ) (**3**); experimental PXRD patterns of a 1:2 stoichiometric mixture of CdO and HMeIm after: e) pre-milling for 5 min at 30 Hz, and f) aging at 45 C and 100% RH for 12 days.

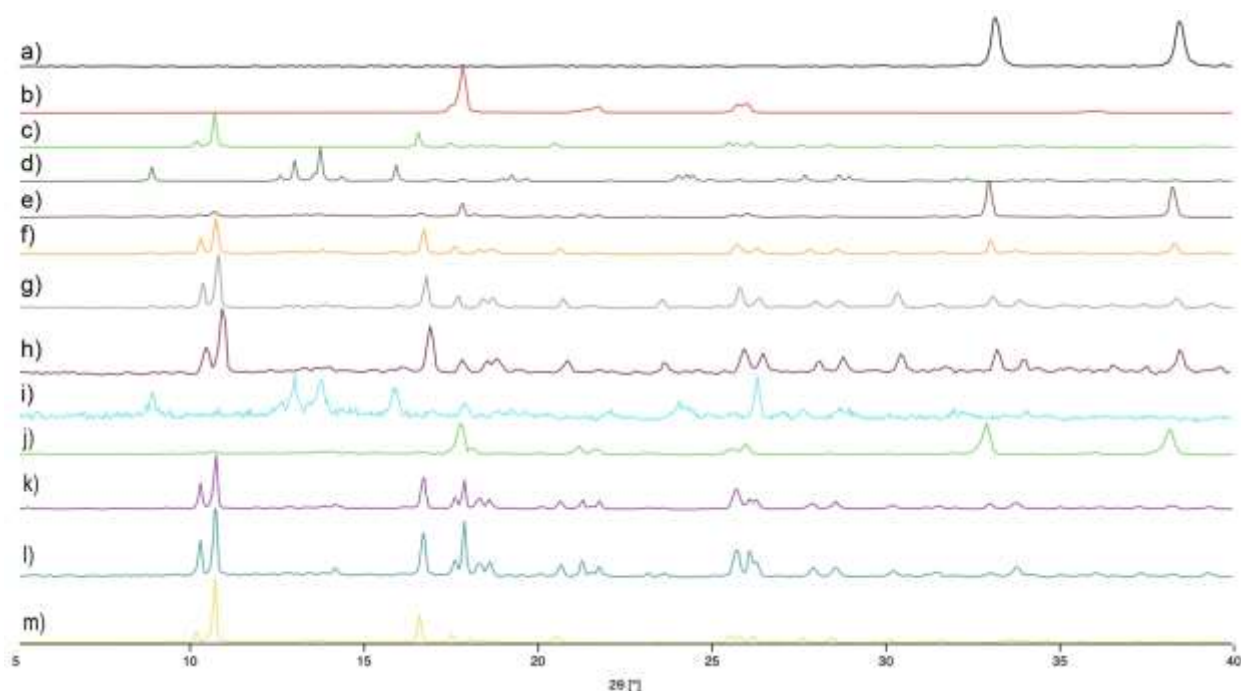


Figure S8: Experimental PXRD patterns of: a) commercial CdO; b) commercial HMeIm; simulated PXRD patterns of: c) **dia**-Cd(MeIm)₂·HMeIm (**1**), and d) **yqt1**-Cd(MeIm)₂ (CSD code GUPBOJ) (**3**); experimental PXRD patterns of a 1:2 stoichiometric mixture of CdO and HMeIm in the presence of NH₄NO₃ after: e) pre-milling for 5 min at 30 Hz; aging at 45 °C and 100% RH for: f) 1 day, g) 7 days, and h) 7 months; i) sample g) after heating to 200 °C; experimental PXRD patterns of a 1:6 stoichiometric mixture of CdO and HMeIm in the presence of NH₄NO₃ after: j) pre-milling for 5 min at 30 Hz; aging at 45 °C and 100% RH for: k) 3 days, and l) 7 days; m) experimental PXRD patterns of a 1:3 stoichiometric mixture of CdO and HMeIm in the presence of NH₄NO₃ after aging at 45 °C and 100% RH for 4 days.

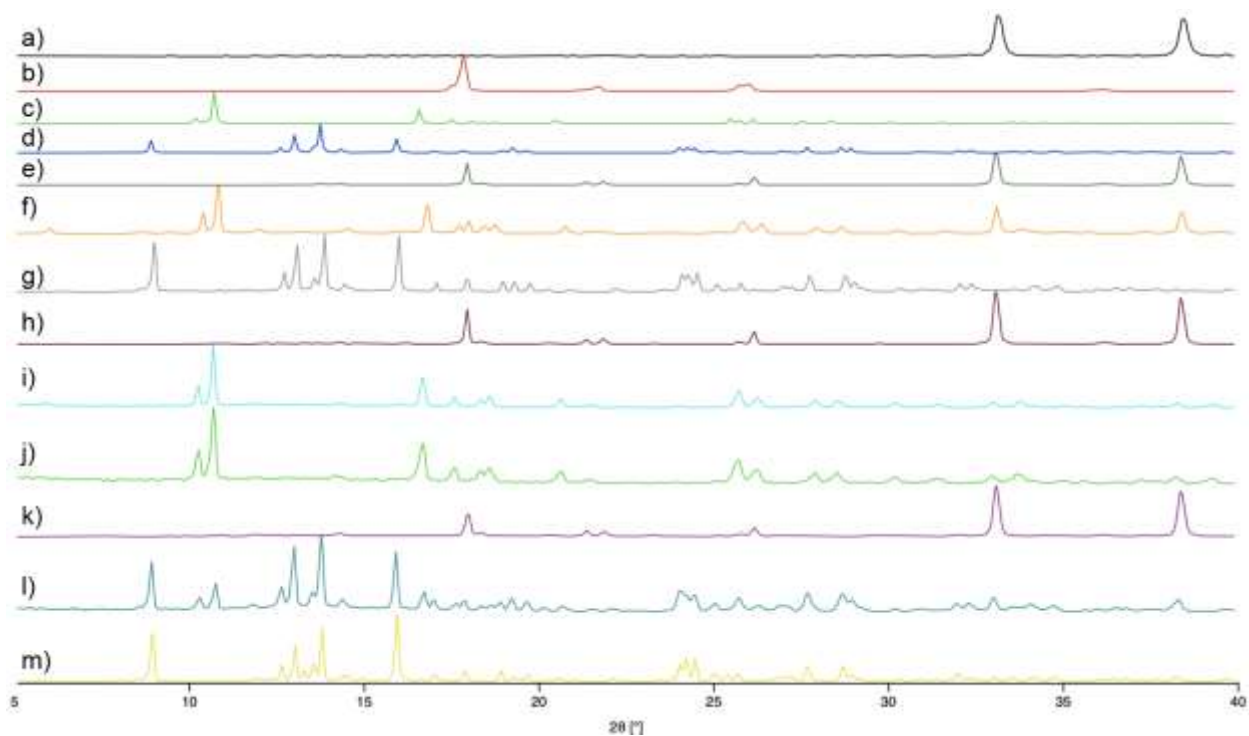


Figure S9: Experimental PXRD patterns of: a) commercial CdO; b) commercial HMeIm; simulated PXRD patterns of: c) **dia**-Cd(MeIm)₂·HMeIm (**1**), and d) **yqt1**-Cd(MeIm)₂ (CSD code GUPBOJ) (**3**); experimental PXRD patterns of a 1:2 stoichiometric mixture of CdO and HMeIm in the presence of (NH₄)₂SO₄ after: e) pre-milling for 5 min at 30 Hz; aging at 45 °C and 100% RH for: f) 1 day, and g) 7 days; experimental PXRD patterns of a 1:3 stoichiometric mixture of CdO and HMeIm in the presence of (NH₄)₂SO₄ after: h) pre-milling for 5 min at 30 Hz; aging at 45 °C and 100% RH for: i) 4 days, and j) 6 days; experimental PXRD patterns of a 1:2 stoichiometric mixture of CdO and HMeIm in the presence of HCafHSO₄ after: k) pre-milling for 5 min at 30 Hz; aging at 45 °C and 100% RH for: l) 1 day, and m) 7 days.

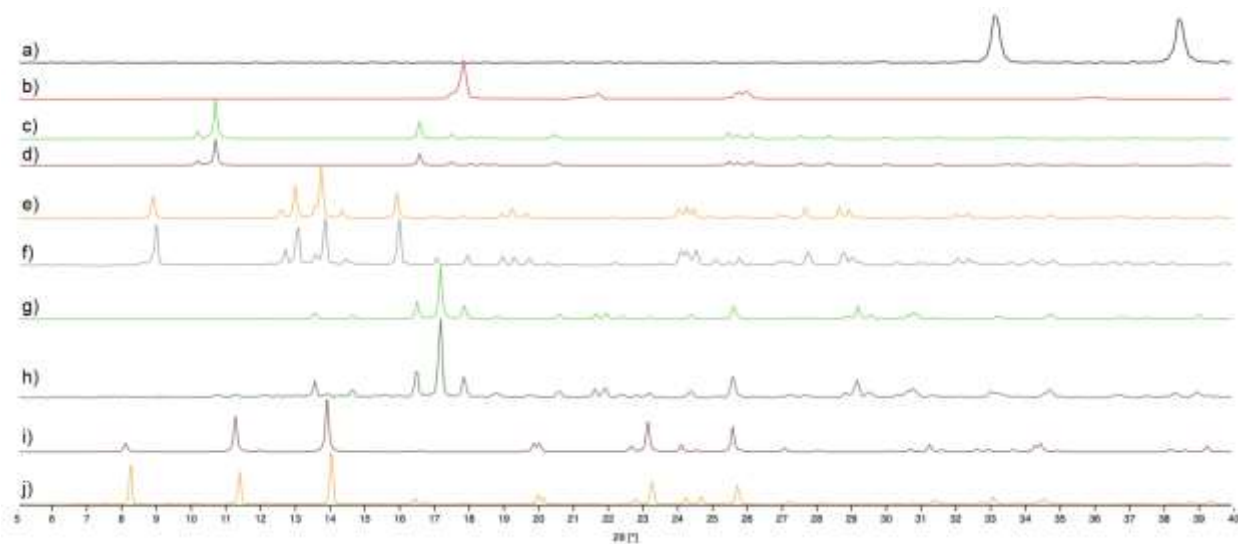


Figure S10: Experimental PXRD patterns of: a) commercial CdO; b) commercial HMeIm; c) simulated, and d) experimental PXRD pattern of **dia**-Cd(MeIm)₂·HMeIm (**1**); e) simulated, and f) experimental PXRD pattern of **yqt1**-Cd(MeIm)₂ (CSD code GUPBOJ) (**3**); g) simulated, and h) experimental PXRD pattern of **dia**-Cd(Im)₂ (CSD code BAYQAU) (**2**); i) simulated, and j) experimental PXRD pattern of Cd(HIm)₆CO₃·3H₂O (CSD code IMIDCP01) (**4**).

4. Thermogravimetric analysis

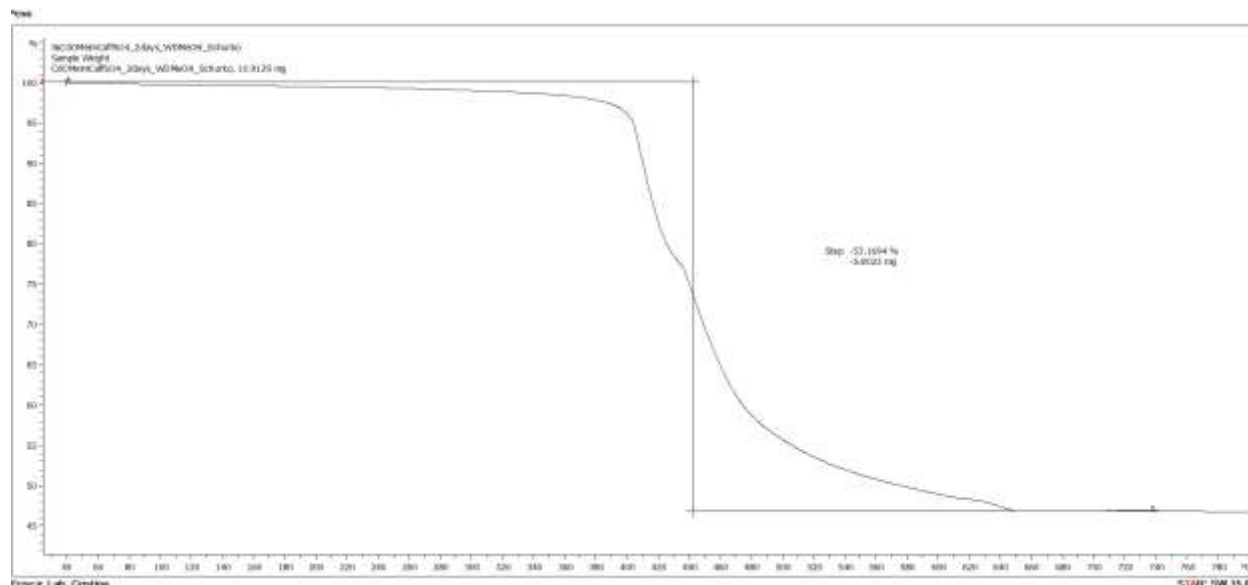


Figure S11: TGA thermogram of **yqt1-Cd(Im)₂** (CSD code GUPBOJ) (**3**) in a dynamic atmosphere of air. The sample was made by aging CdO and HMeIm in a 1:2 ratio at 45 °C and 100% RH in the presence of (HCaf)HSO₄.

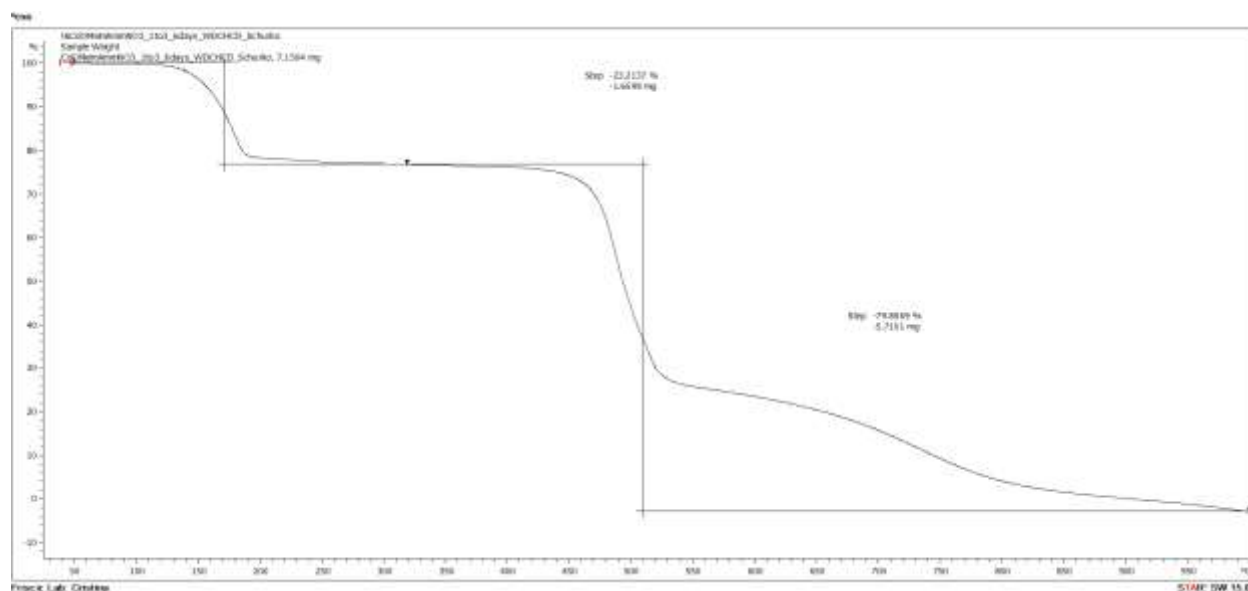


Figure S12: TGA thermogram of **dia-Cd(MeIm)₂·HMeIm** (**1**) in a dynamic atmosphere of N₂. The first step corresponds to the loss of one molecule of unbound HMeIm per Cd²⁺, which accounts for a theoretical value of 23.03% w/w of **1** (experimental value is 23.21%).

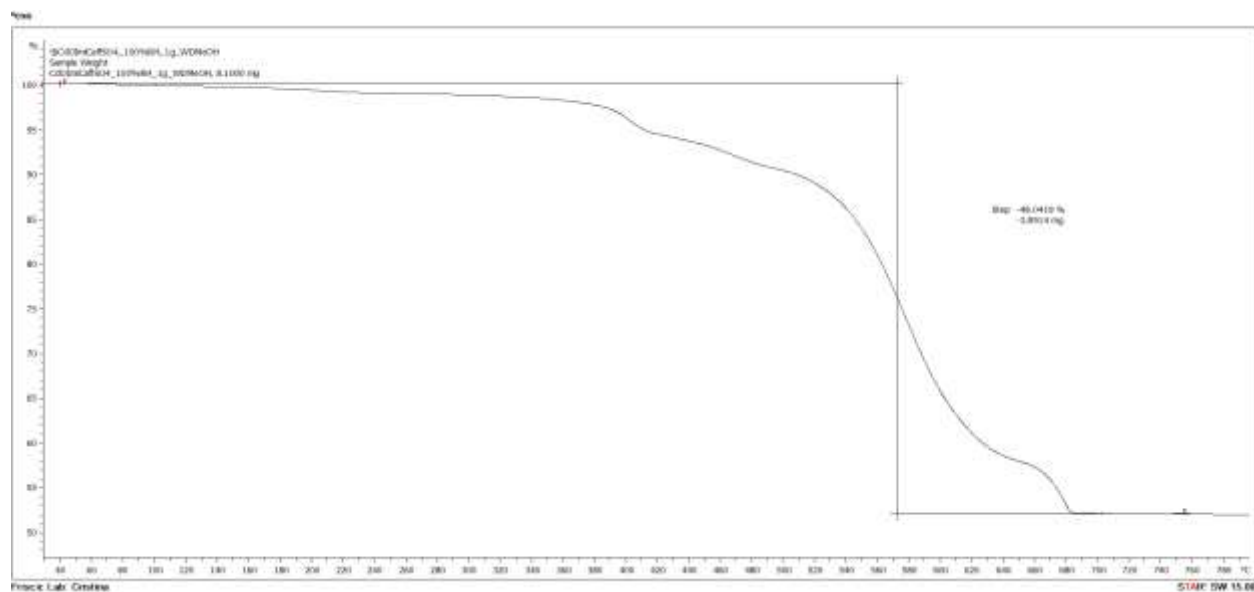


Figure S13: TGA thermogram of **dia-Cd(Im)₂** (CSD code BAYQAU) (2) in a dynamic atmosphere of air.

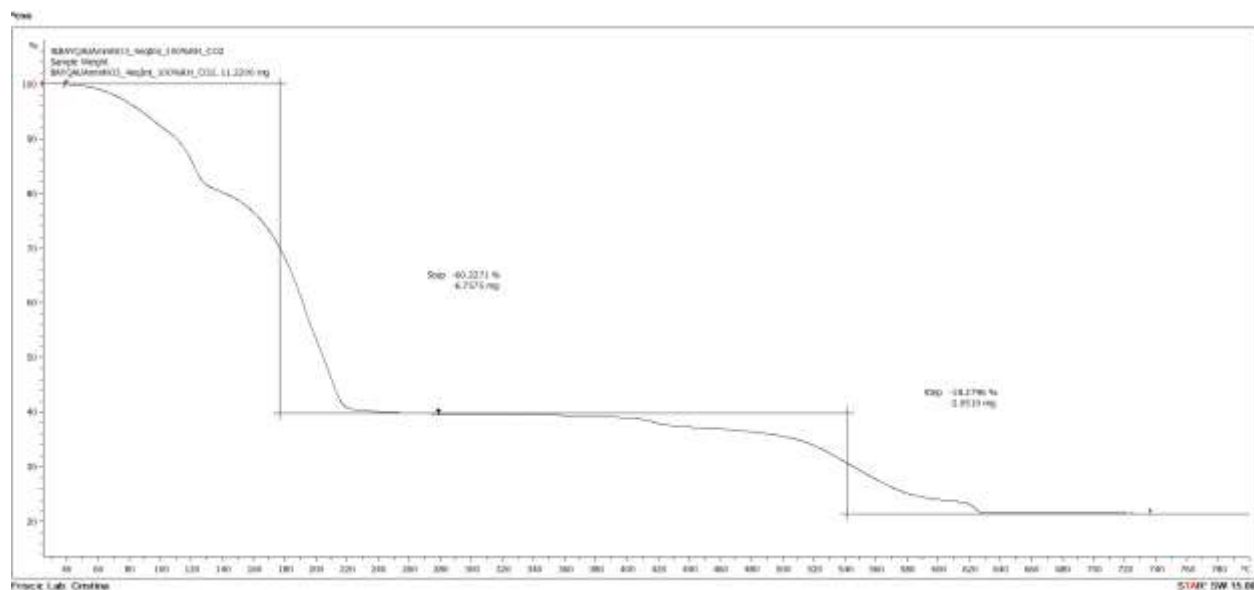


Figure S14: TGA thermogram of **Cd(HIm)₆·3H₂O** (CSD code IMCDP01) (4) in a dynamic atmosphere of air.

4. Fourier-transform infrared (FTIR-ATR) spectroscopy

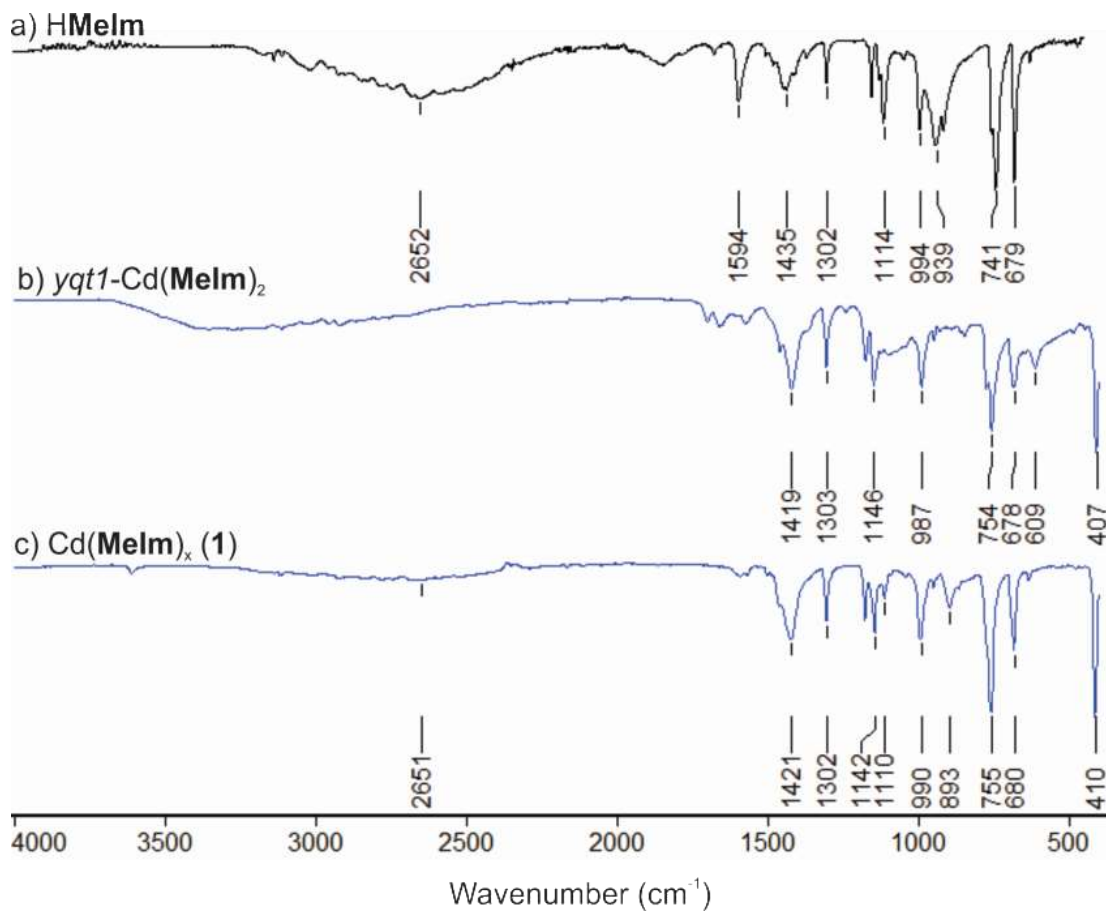


Figure S15: Fourier-transform infrared (FTIR-ATR) spectra of: a) commercial HMeIm; b) yqt1-Cd(MeIm)₂ (CCDC code GUPBOJ) (**3**); and c) dia-Cd(MeIm)₂·HMeIm (**1**).

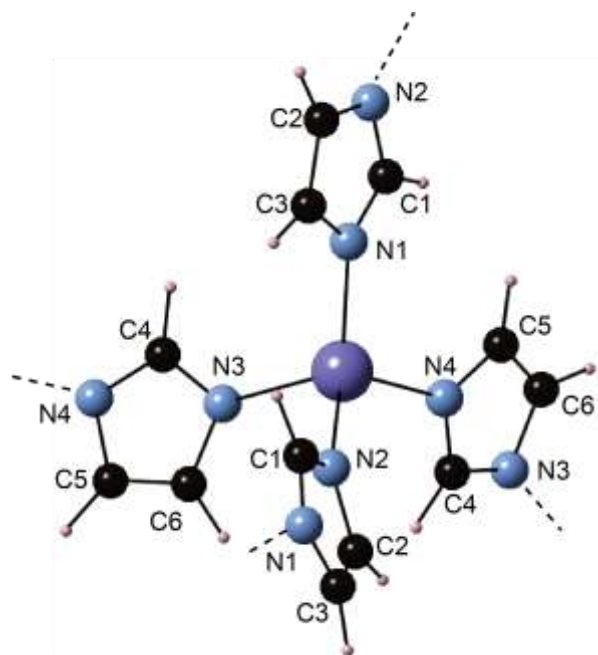


Figure S16: The coordination environment of the Cd atoms in **dia**-Cd[Im]₂ (**2**).

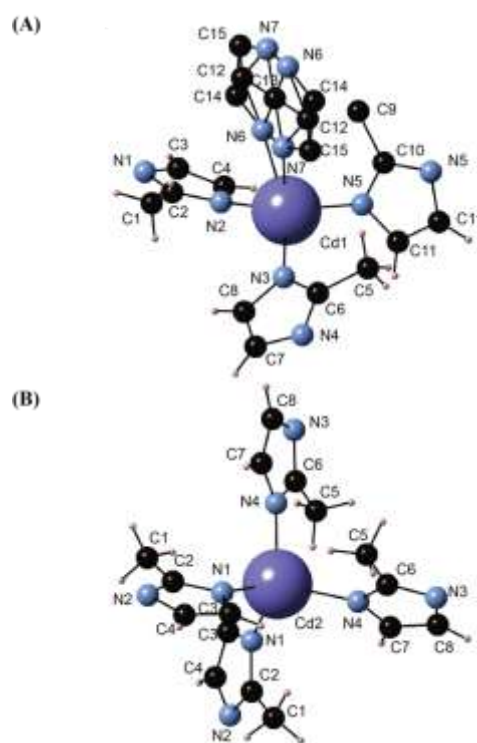


Figure S17: Coordination environment about (A) the Cd1 site, (B) the Cd2 site and atom labeling in the **yqt1**-Cd[MeIm]₂ (**3**) framework. There is an apparent positional disorder of one of the MeIm⁻ ligands at the Cd1 site.

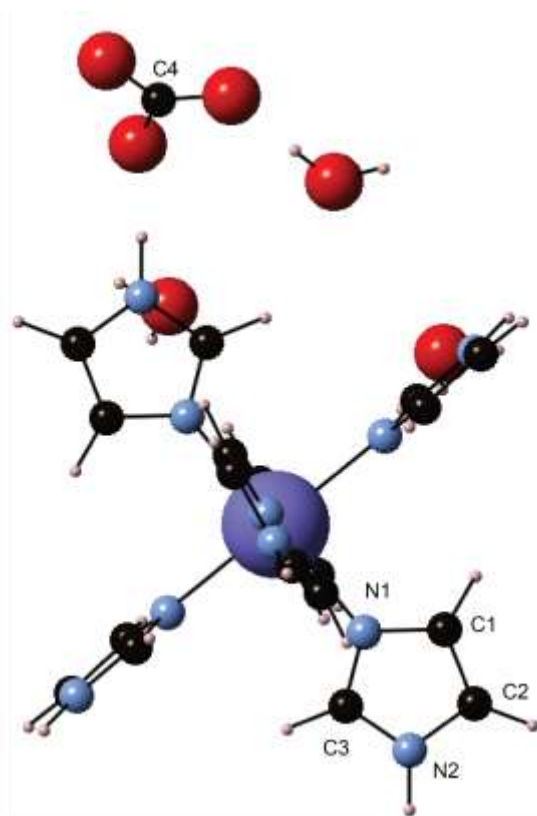


Figure S18: Coordination environment about the Cd atom and atom labelling in the $\text{Cd}[\text{Im}]_6\text{CO}_3 \cdot 3\text{H}_2\text{O}$ (**4**) molecule.

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