Evaluation of Brønsted Acidity and Proton Topology in Zr- and Hf-based Metal– Organic Frameworks Using Potentiometric Acid–Base Titration

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Materials and Methods.

NU-1000,¹ Zr-UiO-66,² Zr-UiO-67(HCl),² Zr-PCN-57,³ and Hf-UiO-66⁴ were synthesized according to methods reported in the literature.

Instrumentation.

Activation of MOF samples was performed on a Micromeritics SmartVacPrep (Micromeritics, Norcross, GA, USA). N₂ adsorption isotherms were collected on a Tristar II 3020 (Micromeritics) at 77 K. All pore size distributions were obtained using a carbon slit pore model with a N₂ kernel (Micromeritics). Powder X-ray diffraction (PXRD) patterns were collected on a Rigaku Smartlab instrument. Thermogravimetric analyses (TGA) were carried out on a TA Instruments Q500 thermogravimetric analyzer with an evolved gas analysis furnace. Samples were heated from 20 °C to 800 °C at a rate of 10 °C/minute under flowing O₂.

MOF Ligands.



Figure S1. Ligands used for a) UiO-66, terephthalic acid, b) UiO-67, biphenyl-4,4'dicarboxylic acid, c) PCN-57, 2',3',5',6'-tetramethyl-terphenyl-4,4"-dicarboxylic acid, d) NU-1000, 1,3,6,8-tetrakis(*p*-benzoic acid)pyrene, and e) MOF-808, benzene-1,3,5tricarboxylic acid. **Synthesis of Hf-UiO-67.** HfCl₄ (120 mg, 0.374 mmol), biphenyl-4,4'-dicarboxylic acid (BPDC) (100 mg, 0.413 mmol), 1.5 g benzoic acid (1.5 g, 12.3 mmol), and *N*,*N*'-dimethylformamide (DMF) (18 mL) were added to an 8-dram vial. The mixture was sonicated for 30 min, then incubated in an oven at 100 °C for 48 h. The resulting white solid was washed with fresh DMF (3 × 20 mL) and then with acetone (3 × 20 mL). After drying the sample in a vacuum oven, the sample was activated on a SmartVacPrep port by heating at 120 °C under vacuum for 12 h. Yield: 65% white microcrystalline powder.

Synthesis of Zr-UiO-67 (Benzoic Acid). The same procedure as Hf-UiO-67 was followed except $ZrCl_4$ (120 mg, 0.515 mmol) was used instead of HfCl₄. Yield: 27% white microcrystalline powder.

Synthesis of Hf-PCN-57. HfOCl₂·8H₂O (120 mg, 0.293 mmol), 2',3',5',6'-tetramethylterphenyl-4,4"-dicarboxylic acid (TPDC-4CH₃) (100 mg, 0.267 mmol), 1.0 g benzoic acid, and *N*,*N*'-dimethylformamide (DMF) (20 mL) were added to an 8-dram vial. The mixture was sonicated for 30 min, then incubated in an oven at 100 °C for 24 h. The sample was washed and activated using the same procedure as that of Hf-UiO-67. Yield: 72% white microcrystalline powder.



Figure S2. N₂ adsorption isotherm for Hf-UiO-67.



Figure S3. N₂ adsorption isotherm for Zr-UiO-67 synthesized with benzoic acid.



Figure S4. N₂ adsorption isotherm for Zr-PCN-57.



Figure S5. N₂ adsorption isotherm for Hf-PCN-57.



Figure S6. N_2 adsorption isotherm for Zr-UiO-66.



Figure S7. N_2 adsorption isotherm for Hf-UiO-66.



Figure S8. N_2 adsorption isotherm for Zr-UiO-67(HCI).





Figure S10. N₂ adsorption isotherm for MOF-808.



Figure S11. Simulated (black, bottom) and experimental (blue, top) PXRD patterns of Hf-UiO-67.



Figure S12. Simulated (black, bottom) and experimental (blue, top) PXRD patterns of Zr-UiO-67 synthesized with benzoic acid.



Figure S13. Simulated (black, bottom) and experimental (blue, top) PXRD patterns of Hf-PCN-57.



Figure S14. Representative crystal structure of UiO-67 showing a missing linker. Zr/Hf, C, and O are shown in green, black, and red, respectively. Hydrogen atoms have been omitted for clarity.



Figure S15. TGA trace of activated Zr-UiO-66.



Figure S16. TGA trace of activated Hf-UiO-66.



Figure S17. TGA trace of activated Zr-UiO-67(HCI) synthesized with hydrochloric acid.



Figure S18. TGA trace of activated Zr-UiO-67 synthesized with benzoic acid.



Figure S19. TGA trace of activated Hf-UiO-67.



Figure S20. TGA trace of activated Zr-PCN-57. N.B.: We do not understand the weight gain observed near 220 °C for Zr-PCN-57; however, we observed similar TGA traces for multiple samples.



Figure S21. TGA trace of activated Hf-PCN-57. N.B.: We do not understand the weight gain observed near 220 °C for Hf-PCN-57; however, we observed similar TGA traces for multiple samples.

MOF	Decompositio (%ZrC	n Starting Point D ₂ /HfO ₂)	Estimated Missing Linkers
	Experimental	Theoretical	(per M ₆ cluster)
Zr-UiO-66	186	222	1.6
Zr-UiO-67(HCl)	242	284	1.3
Zr-UiO-67(BA)	271	284	0.4
Zr-PCN-57	351	391	0.8
Hf-UiO-66	160	171	0.8
Hf-UiO-67(BA)	186	207	1.1
Hf-PCN-57	257	270	0.4

Table S1. Estimated Missing Linkers for UiO-type MOFs from TGA analyses.

Missing linker defects were calculated using the TGA analysis method described by Lillerud et al. 5

PXRD Patterns of MOFs Post-Titration

PXRD assessment of the MOFs in this study titrated to pH = 10.5 show a complete loss of crystallinity. However, typically in the pH range during which node protons appear to be undergoing deprotonation, the MOFs seem to be stable. See below and the manuscript for more details.



Figure S22. PXRD patterns of Zr-UiO-67(BA) parent (black, bottom) and after titration to pH = 8 (blue, top).



Figure S23. PXRD patterns of Zr-UiO-66 parent (black, bottom) and after titration to pH = 8 (blue, top).





Figure S24. PXRD patterns of Zr-UiO-67(HCl) parent (black, bottom) and after titration to pH = 8 (blue, top).



Figure S25. PXRD patterns of NU-1000 parent (black, bottom) and after titration to pH = 9 (blue, top).



2θ (Degrees)

Figure S26. PXRD patterns of MOF-808 parent (black, bottom) and after titration to pH = 9 (blue, top).

Titration Data Zr-UiO-67



Figure S27. Overlay of three acid–base titration curves for Zr-UiO-67 showing reproducibility between samples.



Figure S28. Overlay of three acid–base titration curves for Hf-UiO-67 showing reproducibility between samples.



Figure S29. Acid-base titration curve of Zr-PCN-57 (red) and first derivative curve (blue).

The first derivative curve of Zr-PCN-57 may indicate a second inflection point/equivalence point around pH 7.5 suggesting the presence of some defect sites in the sample; however, due to the poor dispersion of the sample in aqueous solution we are hesitant to draw any conclusions from this data set beyond the pK_a value determined from the equivalence point at 5.4.



Hf-PCN-57

Figure S30. Acid-base titration curve of Hf-PCN-57 (red) and first derivative curve (blue).





Figure S31. Overlay of three acid–base titration curves for Zr-UiO-66 showing reproducibility between samples.



Figure S32. Representative curve fitting for the first derivative of the titration curve of pH as a function of volume of titrant added for Zr-UiO-66: Located peaks (green) (bottom); first derivative data (black circles) and fit trace (blue) (middle); residual plot (red) (top). Peaks were fit as Lorentzian functions.

Multipeak Fit report for Zr-UiO-66: Chi square: 0.0075287 Total fitted points: 220 Multi-peak fit version 2.16 Total Peak Area = 0.20869 +/- 0.0060043 Baseline Type: Constant y0 = 0.0023544+/-0.0010797 Peak 0 Type: Lorentzian Location = +/-0.0055287 1.15 Height = 0.12558 +/-0.0020745 +/-Area = 0.12024 0.0036137 FWHM = 0.60951 +/-0.019437 Fit function parameters Location = 1.15 +/-0.0055287 FWHM = 0.60951 +/-0.019437 Area =0.12024 +/-0.0036137 Peak 1 Type: Lorentzian Location = 2.0103 +/-0.023511 Height = 0.028322 +/-0.0021178 +/-Area = 0.025549 0.0034409 +/-FWHM = 0.57428 0.08251 Fit function parameters +/-Location = 2.0103 0.023511 FWHM = 0.57428 +/-0.08251 Area = 0.025549 +/-0.0034409 Peak 2 Type: Lorentzian Location = 4.5007 +/-0.0084137 Height = 0.069551 +/-0.0020925 Area = 0.06291 +/-0.0033396 FWHM = +/-0.57584 0.032298 Fit function parameters Location = +/-4.5007 0.0084137 FWHM = 0.57584 +/- 0.032298 Area =0.06291 +/- 0.0033396

Table S2. Titrant consumed after the first equivalence point for Zr-UiO-66

	mmoles OH ⁻ consumed between EP1 to EP3
Trial 1	0.333
Trial 2	0.304
Trial 3	0.302
Average	0.313 (± 0.017)

Table S3. Defect calculations for Zr-UiO-66

Missing Linkers	Molecular Formula ^a	MW	mmoles H ⁺ from defects in 0.049 g sample
1	$Zr_6O_4(OH)_4(C_8H_4O_4)_5[(H_2O)(OH)]_2$	1570.01	0.187
2	$Zr_6O_4(OH)_4(C_8H_4O_4)_4[(H_2O)(OH)]_4$	1475.93	0.398
1.50	$Zr_6O_4(OH)_4(C_8H_4O_4)_{4.50}[(H_2O)(OH)]_{3.0}$	1522.97	0.290
1.55	$Zr_6O_4(OH)_4(C_8H_4O_4)_{4.45}[(H_2O)(OH)]_{3.09}$	1518.69	0.299
1.60	$Zr_6O_4(OH)_4(C_8H_4O_4)_{4.40}[(H_2O)(OH)]_{3.20}$	1513.56	0.311
1.67	$Zr_6O_4(OH)_4(C_8H_4O_4)_{4.33}[(H_2O)(OH)]_{3.33}$	1507.29	0.325
1.75	$Zr_6O_4(OH)_4(C_8H_4O_4)_{4.25}[(H_2O)(OH)]_{3.50}$	1499.45	0.343

^aAll missing linkers are assumed to be replaced with 2 H₂O molecules and 2 –OH groups.



Figure S33. Overlay of three acid–base titration curves for Hf-UiO-66 showing reproducibility between samples.



Figure S34. Representative curve fitting for the first derivative of the titration curve of pH as a function of volume of titrant added for Hf-UiO-66: Located peaks (green) (bottom); first derivative data (black circles) and fit trace (blue) (middle); residual plot (red) (top). Peaks were fit as Lorentzian functions.

Multipeak fit report for Hf-UiO-66: Chi square: 0.0061373 Total fitted points: 155 Multi-peak fit version 2.16 Total Peak Area = 0.17198 +/- 0.0046317 Baseline Type: Constant v0 = 0.010397 +/-0.0011316 Peak 0 Type: Lorentzian 0.90188 0.0017373 Location = +/-Height = 0.25346 +/-0.0035568 Area = 0.093242 +/-0.0021403 FWHM = 0.2342 +/-0.0059391 Fit function parameters Location = 0.90188 +/-0.0017373 FWHM = 0.2342 +/-0.0059391 Area =0.093242 +/-0.0021403 Peak 1 Type: Lorentzian Location = 1.3735 +/-0.019753 Height = 0.033173 +/-0.0026263 Area = 0.022545 +/-0.0033673 FWHM = 0.43265 +/-0.069871 Fit function parameters +/-Location = 1.3735 0.019753 FWHM = +/-0.43265 0.069871 Area =0.022545 0.0033673 +/-Peak 2 Type: Lorentzian Location = 3.1761 +/-0.0045788 +/-Height = 0.10617 0.0029212 Area = 0.056192 +/-0.0023521 FWHM = 0.33693 +/-0.01605 Fit function parameters Location = 3.1761 +/-0.0045788 FWHM = 0.33693 +/-0.01605

Area =0.056192 +/- 0.0023521

Table S4. Titrant consumed after the first equivalence point for Hf-UiO-66

	mmoles OH ⁻ consumed between EP1 to EP3
Trial 1	0.252
Trial 2	0.226
Trial 3	0.268
Average	0.249 (±0.021)

Table S5. Defect calculations for Hf-UiO-66

Missing Linkers	Molecular Formula ^a	MW	mmoles H ⁺ from defects in 0.049 g sample
1	$Hf_{6}O_{4}(OH)_{4}(C_{8}H_{4}O_{4})_{5}[(H_{2}O)(OH)]_{2}$	2093.60	0.140
2	Hf ₆ O ₄ (OH) ₄ (C ₈ H ₄ O ₄) ₄ [(H ₂ O)(OH)] ₄	1999.52	0.294
1.55	$Hf_6O_4(OH)_4(C_8H_4O_4)_{4.45}[(H_2O)(OH)]_{3.09}$	2042.28	0.222
1.60	$Hf_6O_4(OH)_4(C_8H_4O_4)_{4.40}[(H_2O)(OH)]_{3.20}$	2037.15	0.231
1.65	$Hf_6O_4(OH)_4(C_8H_4O_4)_{4.35}[(H_2O)(OH)]_{3.30}$	2032.45	0.239
1.70	$Hf_6O_4(OH)_4(C_8H_4O_4)_{4.30}[(H_2O)(OH)]_{3.40}$	2027.74	0.246
1.75	$Hf_6O_4(OH)_4(C_8H_4O_4)_{4.25}[(H_2O)(OH)]_{3.50}$	2023.04	0.254
1.80	$Hf_6O_4(OH)_4(C_8H_4O_4)_{4.20}[(H_2O)(OH)]_{3.60}$	2018.34	0.262
1.85	Hf ₆ O ₄ (OH) ₄ (C ₈ H ₄ O ₄) _{4.15} [(H ₂ O)(OH)] _{3.70}	2013.6	0.270

^aAll missing linkers are assumed to be replaced with 2 H_2O molecules and 2 –OH groups.



Figure S35. Overlay of three acid–base titration curves for NU-1000 showing reproducibility between samples.



Figure S36. Representative curve fittings for the first derivative of the titration curve of pH as a function of volume of titrant added for NU-1000: Located peaks (green) (bottom); first derivative data (black circles) and fit trace (blue) (middle); residual plot (red) (top). Peaks were fit as four Lorentzian functions (left) and three Lorentzian functions (right).

Multipeak fit report for NU-1000 with 4 peaks: Chi square: 0.0026865 Total fitted points: 172 Multi-peak fit version 2.16 Total Peak Area = 0.22374 +/- 0.004041 Baseline Type: Constant y0 = 0+/-0 Peak 0 Type: Lorentzian Location = 1.4513 +/-0.013591 Height = 0.061526 +/-0.00095128 Area = 0.003338 0.12681 +/-FWHM = 1.3121 +/-0.041091 Fit function parameters +/-Location = 1.4513 0.013591 +/-FWHM = 1.3121 0.041091 Area =0.12681 0.003338 +/-Peak 1 Type: Lorentzian Location = 2.1401 +/-0.00069592 Height = 0.31358 +/-0.0031991 Area = 0.057384 +/-0.0011179 FWHM = 0.1165 +/-0.0024424 Fit function parameters Location = 2.1401 +/-0.00069592 +/-FWHM = 0.1165 0.0024424 Area = 0.057384 0.0011179 +/-Peak 2 Type: Lorentzian Location = 2.2713 +/-0.00070087 Height = 0.21853 +/-0.0042816 0.023995 +/-Area = 0.0009536 FWHM = 0.069901 +/-0.0029488 Fit function parameters Location = 2.2713 +/-0.00070087 FWHM = 0.069901 +/-0.0029488 Area = 0.023995 0.0009536 +/-

Peak 3 Type: Lorentzian

Location = Height = Area = FWHM =	2.4977 0.034356 0.015556 0.28826	+/- +/- +/- +/-	0.010343 0.0020581 0.0017403 0.03486			
Fit function Location = FWHM = Area =0.015	parameters 2.4977 0.28826 5556 +/-	+/- +/- 0.001	0.010343 0.03486 7403			
Multipeak fit report Chi square: 0.0082 Total fitted points: Multi-peak fit versio Total Peak Area =	Multipeak fit report for NU-1000 with 3 peaks: Chi square: 0.0082375 Total fitted points: 186 Multi-peak fit version 2.16 Total Peak Area = 0.23603 +/- 0.0051088					
Baseline Type	: Constant					
y0 = 0	+/- 0					
Peak 0 Type	: Lorentzian					
Location = Height = Area = FWHM =	1.5559 0.061113 0.15554 1.6203	+/- +/- +/- +/-	0.023391 0.0014903 0.0045733 0.059643			
Fit function Location = FWHM = Area =0.155	parameters 1.5559 1.6203 554 +/-	+/- +/- 0.004	0.023391 0.059643 I5733			
Peak 1 Type	: Lorentzian					
Location = Height = Area = FWHM =	2.1396 0.30434 0.052134 0.10905	+/- +/- +/- +/-	0.0011479 0.0054959 0.0017519 0.0039493			
Fit function Location = FWHM = Area =0.052	parameters 2.1396 0.10905 2134 +/-	+/- +/- 0.001	0.0011479 0.0039493 7519			

Peak 2 Type: Lorentzian

Location =	2.2716	+/-	0.0013596
Height =	0.21396	+/-	0.0063174
Area =	0.028358	+/-	0.0014543
FWHM =	0.084377	+/-	0.0047285
Fit function p	arameters		
Location =	2.2716	+/-	0.0013596
FWHM =	0.084377	+/-	0.0047285
Area =0.0283	358 +/-	0.0014	4543

Table S6. Total moles titrant consumed for NU-1000

	Total mmoles OH ⁻ consumed to EP3	Total mmoles OH ⁻ consumed to EP4
Trial 1	0.226	0.250
Trial 2	0.216	0.247
Trial 3	0.216	0.237
Average	0.219 (± 0.006)	0.244 (± 0.007)

Table S7. Defect calculations for NU-1000

Occupancy of Mesopores (%)	Molecular Formula	MW	Total mmoles H [⁺] in 0.049 g sample
0	Zr ₆ O ₄ (µ ₃ -OH) ₄ (OH) ₄ (H ₂ O) ₄ (C ₄₄ H ₂₂ O ₈) ₂	2176.76	0.360
20	$\begin{array}{l} [Zr_{6}O_{4}(OH)_{4}(OH)_{4}(H_{2}O)_{4}(C_{44}H_{22}O_{8})_{2}]_{0.80}\\ [(Zr_{6}O_{4}(OH)_{4})_{2}(C_{44}H_{22}O_{8})_{6}]_{0.20}\end{array}$	2827.54	0.252
25	$\begin{array}{l} [Zr_{6}O_{4}(OH)_{4}(OH)_{4}(H_{2}O)_{4}(C_{44}H_{22}O_{8})_{2}]_{0.75}\\ [(Zr_{6}O_{4}(OH)_{4})_{2}(C_{44}H_{22}O_{8})_{6}]_{0.25}\end{array}$	2990.23	0.229
30	$\begin{array}{l} [Zr_6O_4(OH)_4(OH)_4(H_2O)_4(C_{44}H_{22}O_8)_2]_{0.70} \\ [(Zr_6O_4(OH)_4)_2(C_{44}H_{22}O_8)_6]_{0.30} \end{array}$	2949.12	0.211

The first derivative curve of NU-1000 can be fit to three or four Lorentzian peaks to give three or four equivalence points, respectively (Fig. S30). Adding a fourth peak to the fitting does not significantly shift the peak locations of peaks 0–3 (see above). We expect that NU-1000 likely has only three real pK_a values (calculated $pK_{a4} = 9.43 \pm 0.09$), so we calculated defects based on the position of peak 2 (EP3); however, it should be noted that using EP4 in the calculation does not significantly change the occupancy of the mesopores with an estimated value of 20% instead of 25%.⁶



Figure S37. Overlay of three acid–base titration curves for Zr-UiO-67(HCI) showing reproducibility between samples.



Figure S38. Representative curve fitting for the first derivative of the titration curve of pH as a function of volume of titrant added for Zr-UiO-67(HCI): Located peaks (green) (bottom); first derivative data (black circles) and fit trace (blue) (middle); residual plot (red) (top). Peaks were fit as Lorentzian functions. Peak 0 was used for fitting purposes only; calculations were done using Peak 1 as the first equivalence point.

Multipeak fit report for Zr-UiO-67(HCI). Chi square: 0.0041681 Total fitted points: 180 Multi-peak fit version 2.16 Total Peak Area = 0.23759 +/- 0.0080377 Baseline Type: Constant v0 = 0+/-0 Peak 0 Type: Lorentzian Location = 0.93205 +/-0.024521 Height = 0.060447 +/-0.0021793 Area = 0.077706 +/-0.0058788 FWHM = 0.8184 +/-0.052022 Fit function parameters Location = 0.93205 +/-0.024521 FWHM = 0.8184 +/-0.052022 Area = 0.077706 +/-0.0058788 Peak 1 Type: Lorentzian Location = 1.2785 +/-0.0022141 Height = 0.16231 +/-0.0042786 Area = 0.046839 +/-0.0030097 FWHM = 0.18372 +/-0.0096725 Fit function parameters +/-Location = 1.2785 0.0022141 FWHM = +/-0.18372 0.0096725 Area = 0.046839 0.0030097 +/-Peak 2 Type: Lorentzian Location = 1.5512 +/-0.008467 +/-Height = 0.062172 0.0029672 Area = 0.029369 +/-0.002985 FWHM = 0.30073 +/-0.027024 Fit function parameters Location = 1.5512 +/-0.008467 FWHM = 0.30073 +/-0.027024

Area =0.029369 +/- 0.002985

Peak 3 Type: Lorentzian

Location =	2.9567	+/-	0.0099685
Height =	0.029641	+/-	0.0028731
Area =	0.0095518	+/-	0.0012787
FWHM =	0.20515	+/-	0.032995

Fit function parameters				
Location =	2.9567	+/-	0.0099685	
FWHM =	0.20515	+/-	0.032995	
Area =0.009	5518 +/-	0.001	2787	

Peak 4 Type: Lorentzian

Location =	3.9125	+/-	0.015982
Height =	0.045507	+/-	0.0013322
Area =	0.074124	+/-	0.003231
FWHM =	1.0369	+/-	0.059319

Fit function parameters

Location =	3.912	5	+/-	0.015982
FWHM =	1.0369	9	+/-	0.059319
Area =0.074	124	+/-	0.0032	231

	mmoles OH ⁻ consumed between EP1 to EP4
Trial 1	0.268
Trial 2	0.270
Trial 3	0.263
Average	0.266 (± 0.004)

Table S8. Titrant consumed after the first equivalence point for Zr-UiO-67(HCI)

Table S9. Defect calculations for Zr-UiO-67(HCl)

Missing Linkers	Molecular Formula ^a	MW	mmoles H⁺ from defects in 0.049 g sample
1	$Zr_6O_4(OH)_4(C_{14}H_8O_4)_5[(H_2O)(OH)]_2$	1950.51	0.151
2	$Zr_6O_4(OH)_4(C_{14}H_8O_4)_4[(H_2O)(OH)]_4$	1780.33	0.330
1.60	$Zr_6O_4(OH)_4(C_{14}H_8O_4)_{4.40}[(H_2O)(OH)]_{3.20}$	1848.40	0.254
1.65	$Zr_6O_4(OH)_4(C_{14}H_8O_4)_{4.35}[(H_2O)(OH)]_{3.30}$	1839.89	0.264
1.70	$Zr_6O_4(OH)_4(C_{14}H_8O_4)_{4.30}[(H_2O)(OH)]_{3.40}$	1831.38	0.273
1.75	$Zr_6O_4(OH)_4(C_{14}H_8O_4)_{4.25}[(H_2O)(OH)]_{3.50}$	1822.88	0.282
1.80	$Zr_6O_4(OH)_4(C_{14}H_8O_4)_{4.20}[(H_2O)(OH)]_{3.60}$	1814.37	0.292

^aAll missing linkers are assumed to be replaced with 2 H_2O molecules and 2 –OH groups.

Note that if the third equivalence point is used for the missing linker calculation for Zr-UiO-67(HCl) instead of the fourth apparent equivalence point, 0.168 ± 0.002 mmoles of OH⁻ is consumed, which corresponds to approximately 1.10 missing linkers (0.167 mmoles H⁺ from defects in 0.049 g sample).



Figure S39. Overlay of three acid–base titration curves for activated MOF-808 showing reproducibility between samples.



Figure S40. Representative curve fitting for the first derivative of the titration curve of pH as a function of volume of titrant added for MOF-808: Located peaks (green) (bottom); first derivative data (black circles) and fit trace (blue) (middle); residual plot (red) (top). Peaks were fit as Lorentzian functions.

Multipeak fit report for MOF-808 Chi square: 0.0054865 Total fitted points: 247 Multi-peak fit version 2.16 Total Peak Area = 0.23749 +/- 0.0038647 Baseline Type: Constant v0 = 0+/-0 Peak 0 Type: Lorentzian Location = 1.7495 +/-0.013917 Height = 0.059156 +/-0.0010558 Area = 0.13776 +/-0.0031872 FWHM = 1.4825 +/-0.046555 Fit function parameters Location = 1.7495 +/-0.013917 FWHM = 1.4825 +/-0.046555 Area =0.13776 +/-0.0031872 Peak 1 Type: Lorentzian Location = 2.8676 +/-0.0011336 Height = 0.18951 +/-0.0036161 Area = 0.035379 +/-0.00081817 FWHM = 0.11885 +/-0.0035296 Fit function parameters +/-Location = 2.8676 0.0011336 +/-FWHM = 0.11885 0.0035296 Area = 0.035379 +/-0.00081817 Peak 2 Type: Lorentzian Location = 3.4083 +/-0.0044165 0.059309 +/-Height = 0.0029664 Area = 0.016479 +/-0.00092191 FWHM = 0.17688 +/-0.013238 Fit function parameters Location = 3.4083 +/-0.0044165 FWHM = 0.17688 +/-0.013238

Area =0.016479 +/- 0.00092191

Peak 3 Type: Lorentzian

Location =	5.5089	+/-	0.012687
Height =	0.041943	+/-	0.001473
Area =	0.047867	+/-	0.0018049
FWHM =	0.72653	+/-	0.03812

Fit function parameters

Location =	5.508	9	+/-	0.012687
FWHM =	0.726	53	+/-	0.03812
Area =0.0478	367	+/-	0.0018	3049

	mmoles OH ⁻ consumed
Trial 1	0.541
Trial 2	0.580
Trial 3	0.551
Average	0.557 (±0.020)

Table S10. Total moles titrant consumed for MOF-808

Table S11. Proton quantification calculation for MOF-808

Molecular Formula	MW	Total mmoles H ⁺ expected in 0.049 g sample
Zr ₆ O ₄ (µ ₃ -OH) ₄ (OH) ₆ (H ₂ O) ₆ (C ₉ H ₃ O ₆) ₂	1303.73	0.827

 $(0.557/0.827) \times 100\% = 67\%$ of mmoles H⁺ expected

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