Supplementary information

3-(4-Methylthiophenyl)acetylacetone – Ups and Downs of Flexibility in the Synthesis of Mixed Metal-Organic Frameworks. Ditopic Bridging of Hard and Soft Cations and Site-Specific Desolvation

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S1 Refinement and Crystal Structure Details

Intensity data were collected with a Bruker D8 goniometer equipped with an APEX CCD area detector and an Incoatec microsource (Mo-K_{α} radiation, $\lambda = 0.71073$ Å, multilayer optics) at 100(2) K (Oxford Cryostream 700 instrument, Oxfordshire, UK). Data were integrated with SAINT¹ and corrected for absorption by multi-scan methods;² for this purpose, redundancies between 5 and 12 were achieved. The structures were solved by intrinsic phasing³ and refined by full matrix least squares procedures based on F^2 , as implemented in SHELXL-13.³ Hydrogen atoms were treated as riding with C–H = 0.98 Å for CH₃ groups, C–H = 0.95 Å for aryl-hydrogen and C–H = 1.00 Å for alkyl-hydrogen bonds. Their isotropic displacement parameters were constrained to U_{iso} (H) = 1.5 U_{eq} (C) for methyl groups or U_{iso} (H) = 1.2 U_{eq} (C) otherwise. The position of OH hydrogen atoms were refined freely.

S1.1 4

The position of the hydroxyl hydrogen atom H2A was refined freely. An ORTEP plot of the crystal structure is shown below.



Fig. S1 ORTEP plot of 4 drawn at 70% probability.

S1.2 5

The position of the hydroxyl hydrogen atom H2 was refined freely.

S1.3 6a, 6b and 6c

A satisfactory structure model was obtained for **6a**; the CIF has been deposited with the CCDC and is available electronically. The isotypic solids **6b** and **6c** are affected by additional disorder (see below) and only described for comparison.

No structure model based on geometrically reasonable local density maxima could be derived for the co-crystallized disordered solvent molecules around the $\bar{4}$ inversion axis in **6**. For **6a** SQUEEZE⁴ identified solvent accessible voids with a total volume of 216 Å³ and a content of 52 electrons and was used to correct the low-order reflections for the solvent contribution.

Although **6a-c** are isotypic (Fig. S2), additional disorder in **6b** and **6c** precludes refinement of the well-ordered structure model even for the non-solvent part. In these structures the cavity around the $\overline{4}$ inversion axis does not only contain solvent molecules but also one of the three ligands, namely the one incorporating S3, is disordered with the minority conformer tilted towards the cavity with low site occupancies (7.8(3)% for **6b**, 10.4(3)% for **6c**). This conformation would imply prohibitively short

intermolecular contacts to symmetry equivalent molecules and can therefore only be adopted by one molecule in the vicinity of the $\overline{4}$ site (Fig. S2), with the others necessarily in majority conformation. As the remaining cavity is still filled with solvent molecules and the two aspects of disorder are cooperative, no satisfactory correction with the SQUEEZE algorithm can be achieved. Both approaches neglecting one of these two closely associated aspects of disorder either in the framework or the solvent part do not meet our requirements for a convincing structure model.



	6a	6b	6c
a,b / Å	17.703(4)	17.8027(13)	17.826(3)
c / Å	46.013(8)	46.166(3)	45.877(8)
V / Å ³	14420(7)	14632(2)	14578(6)

Fig. S2 (left) ORTEP plots of an excerpt of the crystal structure of **6c** drawn at 50% probability showing the two possible conformations of one of the ligand molecules. d(C36B-C36Ba) = 0.75(4) Å. (right) Table listing the lattice parameters of **6a**, **6b** and **6c**. Symmetry operator: a = 2 - x, 1/2 - y, z.

S1.4 7c, 7d and 7e

A satisfactory structure model was derived for **7e**; the CIF has been deposited with the CCDC and is available electronically. The isotypic solids **7c** and **7d** are only described for comparison.

Fig. S3 depicts the disorder that occurs in the structure type 7. 7c and 7d were treated by assigning split positions A and B to the atoms associated with methyl group C12 and to two alternative orientations A and B for the acetone molecule associated with O13. Only homonymous orientations subtend reasonable intermolecular contacts: the disorder was therefore treated as coupled, with site occupancies in matching orientations constrained to the same value and the sum of the occupancies constrained to unity.

7e on the other hand exhibits a more complex disorder involving a second acetone molecule. In 7e the acacSMe ligands associated with S2 and S6 exhibit additional disorder and were thus assigned split positions **B** and **B**'. The disorder model was restricted to the PhSMe moieties, as the difference between the alternative orientations of the acetylacetone moieties is too small to be modeled. To retain reasonable intermolecular contacts the occupancy of the additional acetone molecule associated with O14 has to be treated as coupled to the split position **B**' of the PhSMe moieties; these occupancies were therefore constrained to the same value (Fig. S3). Furthermore the acetone molecule associated with O14 is only compatible with the **B** orientation of C12. This disorder was, however, not directly treated as coupled using constraints, as it is still chemically sound as long as the occupancy of the acetone associated with O14 is lower than the site occupancy of C12B.



Fig. S3 Cooperative disorder in 7e. The three alternative orientations are depicted in picture A, B and C with their respective occupancy χ . Symmetry operator: a = 1 - x, 1 - y, 1 - z.

Despite the different disorder patterns the three compounds essentially have the same unit cell (Tab. S1).

Table S1 Lattice parameters of 7c-e.	
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	7c	7d	7e
a / Å	14.589(4)	14.6243(9)	14.603(4)
b / Å	25.540(7)	25.7787(15)	25.521(7)
c / Å	20.309(6)	20.1194(12)	20.434(6)
β / Å	100.318(4)	100.652(2)	100.830(4)
V / Å ³	7445(4)	7454.2(8)	7480(4)

Even a restrained model for the co-crystallized acetone molecules remained unsatisfactory. They were assigned the geometry of a published crystal structure⁵ and treated as rigid fragments in the refinement, with a common isotropic displacement parameter for all non-hydrogen atoms in all acetone residues.

S1.5 8a

The structure contains considerable residual electron density that suggests a disorder for the acac-SMePh ligand associated with S1. Tentative refinement of a disordered model resulted in a low occupancy of ca. 0.1 for the minority conformer and led to convergence problems; this more complicated model was therefore discarded.

S1.6 Thermal removal of CHCl₃ from 8c

In order to investigate the partial removal of a $CHCl_3$ molecule from **8c**, a single crystal was fixed with two-component adhesive to a glass fiber. Diffraction data were collected at 100 K with intervening heating cycles at 360 K. During these cycles the crystal was heated to 320 K with a ramp of 360 K h⁻¹ and subsequently to 360 K at $120 K h^{-1}$. The temperature of 360 K was then kept for 15 min, followed by fast cooling to 100 K. For each data set thus obtained the diffraction data were integrated to the same threshold of 0.83 Åand the refinement was conducted with the same number of variables. For the last two data sets not all non-hydrogen atoms could be assigned acceptable anisotropic displacement parameters. Occupancy refinement of **CLF2** was conducted with fixed displacement parameters taken from the data collection on the pristine crystal prior to the first heating cycle.

S1.7 8f

The occupancy of the alternative trivalent cations Al1 and Ga1 sharing the same site was refined while fractional coordinates and displacement parameters were constrained to be equal and the sum of the site occupancies was constrained to unity. Systematic data truncation had been suggested as a tool to judge the reliability of such refined occupancies.⁶ In addition to our refinement with all data, test refinements excluding 20% of the reflections with (I) the lowest resolution and (II) the highest resolution were conducted. Tab. S2 shows that the refined occupancies do not vary significantly.

Table S2 Occupation of Ga1 in 8f in dependance of data truncation.

	(I)	(II)	(III)
resolution of excluded reflections	> 1.25 Å	< 0.79 Å	-
number of unique reflections	9825	9825	12305
occupation of Ga1	0.450(9)	0.447(3)	0.447(3)

Table S3 Coordination sphere of **8b**, **8e** and **8f** in comparison. Listed are the mean M^{III} -O bond lengths \bar{d} and the mean square root of the difference of the ADP component in the direction of the respective bond $\sqrt{\Delta_{ADP}}$.

	8b	8e	8f
<i>ā</i> (M ^Ⅲ −O) / Å	1.878(9)	1.944(12)	1.910(15)
$\sqrt{\Delta_{\text{ADP}}} \text{ (M}^{\text{III}} \text{-O)} / \text{\AA}$	0.032(13)	0.031(13)	0.056(8)
$\sqrt{\Delta_{\text{ADP}}}$ (C–O) / Å	0.048(14)	0.004(3)	0.058(16)

S2 Structure type 8 data

parameters	8a	8b	8c	8d	8e	8f
d(Ag1-09)	2.646(2)	2.649(2)	2.6566(5)	2.664(6)	2.644(3)	2.649(2)
$d(Ag1 - O10^{a})$	2.797(2)	2.776(3)	2.7782(5)	2.764(6)	2.777(4)	2.791(2)
<i>bpr</i> (Ag1) / %	50.9	51.6	52.6	55.4	52.5	51.7
$\tau(Ag1)$	0.52	0.52	0.50	0.47	0.51	0.51
Cl3…Cl3 ^b / Å	3.2968(14)	3.2978(14)	3.319(2)	3.331(3)	3.294(2)	3.3008(12)
C37–Cl3····Cl3 ^b / $^{\circ}$	153.48(11)	153.13(13)	154.50(16)	154.1(3)	153.84(17)	153.66(11)
V(CLF1 void) / Å ³	125	123	127	129	126	121
V(CLF2 void) / Å ³	145	145	142	155	143	146

Table S4 Comparison of the different parameters for **8a** - **8f** mentioned in the main text. Symmetry operators: a = x - 1, y, z; b = 1 - x, 1 - y, 1 - z. *bpr*: progress along the berry pseudorotation axis.



Fig. S4 Simulated and experimental powder pattern of 4.



Fig. S5 Simulated and experimental powder pattern of 5.



Fig. S6 Simulated and experimental powder pattern of 6a.



Fig. S7 Simulated and experimental powder pattern of 6b.



Fig. S8 Simulated powder pattern of 7c and simulated and experimental powder pattern of 6c. For this compound 6c is the dominant crystal structure that represents the bulk material. No experimental powder pattern matching 7c could be recorded.

For the Fe^{III} complex **7d** no matching powder pattern could be recorded.



Fig. S9 Simulated and experimental powder pattern of 7e.



Fig. S10 Simulated and experimental powder pattern of 8a.



Fig. S11 Simulated and experimental powder pattern of 8b.



Fig. S12 Simulated and experimental powder pattern of 8c.



Fig. S13 Simulated and experimental powder pattern of 8d.



Fig. S14 Simulated and experimental powder pattern of 8e.

S4 NMR spectra



Fig. S15 1 H-NMR spectra of 4 (bottom) and 4-d₃ (top). Measured in CD₂Cl₂ at 298 K and 400 MHz.







Fig. S17 HMBC spectrum of 4. Measured in CD_2CI_2 at 298 K.



Fig. S18 $^1\text{H-NMR}$ spectrum of 5. Measured in CD_2Cl_2 at 298 K and 400 MHz.



Fig. S19 HSQC spectrum of 5. Measured in CD_2Cl_2 at 298 K.





S5 Crystal data

Table S5 Crystal data for 4, 5, 6a, 7e and 8a.

Compound	8b	8c	8d	8e	8f
Empirical formula	$\mathrm{C}_{38}\mathrm{H}_{41}\mathrm{A}\mathrm{gAlCl}_{7}\mathrm{O}_{10}\mathrm{S}_{3}$	$C_{38}H_{41}AgCl_7Cr$ $O_{10}S_{20}$	C ₃₈ H ₄₁ AgFeCl ₇ O.oS ₂	$\mathrm{C}_{38}\mathrm{H}_{41}\mathrm{AgGaCl}_{7}\mathrm{O}_{10}\mathrm{S}_{3}$	$C_{38}H_{41}AgAl_{0.55}Cl_7Ga_{0.45}O_{10}S_3$
Moiety formula	$C_{36}H_{39}AgAlClO_{10}S_3, 2(CHCl_3)$	$C_{36}^{0.0-3}$ $C_{36}^{0.0}$ H ₃₉ AgCrClO ₁₀ S ₃ , 2 (CHCl ₃)	C_{36}^{1003} H $_{39}^{3}$ AgFeClO $_{10}$ S $_{3}^{3}$, 2 (CHCl $_{3}$)	$C_{36}H_{39}AgGaClO_{10}S_3, 2(CHCl_3)$	$C_{36}H_{39}AgAl_{0.55}CIGa_{0.45}O_{10}S_{3}, 2 (CHCl_{3})$
$M_{\rm r}$ / g mol ⁻¹	1136.89	1161.91	1165.76	1179.63	1156.01
T / K Curred docorintion	100(2) solourloss noodlo	100(2)	100(2) 20d 20d	100(2) colourlass rad	100(2) colourdons bloot
Crystal size / mm ³	$0.37 \times 0.09 \times 0.07$	putpic block $0.33 \times 0.12 \times 0.08$	$0.50 \times 0.09 \times 0.09$	$0.42 \times 0.10 \times 0.08$	$0.18 \times 0.17 \times 0.10$
Crystal system	monoclinic	monoclinic	monoclinic	monoclinic	monoclinic
Space group (No.)	$P2_1/c$ (14)	$P2_1/c$ (14)	$P2_1/c$ (14)	$P2_{1}/c$ (14)	$P2_{1}/c$ (14)
<i>a /</i> Å	7.6805(7)	7.6885(14)	7.674(3)	7.6796(17)	7.6934(9)
b / Å	31.867(3)	31.944(6)	32.075(13)	31.828(7)	31.866(4)
<i>c</i> / Å	19.0168(18)	19.105(4)	19.065(8)	19.013(5)	19.046(3)
β∕°	97.514(2)	97.544(3)	97.790(7)	97.932(4)	97.751(2)
V / Å ³	4614.4(7)	4651.7(15)	4649(3)	4602.8(19)	4626.6(11)
Z	4	4	4	4	4
μ / mm^{-1}	1.05	1.244	1.323	1.607	1.294
Total/unique reflections	69783/13106	64891/11607	43264/8823	58880/10207	68006/12305
Rint	0.1159	0.1426		0.0969	0.1195
$(\sin(\theta)/\lambda)_{\max}$	0.70	0.67	0.61	0.64	0.64
Completeness to θ_{\max}	0.996	0.995	0.993	0.988	0.999
$R\left[F^2 > 2\sigma(F^2) ight]$	0.0511	0.0515	0.0654	0.0536	0.0394
$WR_2(F^2)$	0.1210	0.1232	0.1486	0.1234	0.0859
GOF	1.030	0.1038	1.077	1.110	0.98
No. of parameters	550	550	551	550	551
No. of restraints		0	0	0	0
$(\Delta ho_{ m max}/\Delta ho_{ m min})$ / eÅ ⁻³ CSD mumber	1.717/-0.924 1968735	0.988/—0.925 1968736	1.118/-0.947 1968737	0.988/-0.682 1968738	0.648/-0.598 1968739
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Table S6 Crystal data for 8b, 8c, 8d, 8e and 8f.

References

- 1 Bruker, SAINT+: Program for Reduction of Data Collected on Bruker CCD Area Detector Diffractometer, 2009.
- 2 Bruker, SADABS, 2008.
- 3 G. M. Sheldrick, Acta Crystallogr., Sect. A: Found. Crystallogr., 2015, 71, 3-8.
- 4 A. L. Spek, Acta Crystallogr., Sect. C: Cryst. Struct. Commun., 2015, 71, 9–18.
- 5 D. R. Allan, S. J. Clark, R. M. Ibberson, S. Parsons, C. R. Pulham and L. Sawyer, *Chem. Commun.*, 1999, 751–752.
- 6 H. Kroll, T. Lueder, H. Schlenz, A. Kirfel and T. Vad, Eur. J. Mineral., 1997, 9, 705–733.