

**Desolvation Process in the Flexible Metal-Organic Framework
[Cu(Me-4py-trz-ia)], Adsorption of Dihydrogen and Related
Structure Responses**

Supporting Information

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1. MOF Synthesis

Synthesis of **1a** was performed in a similar way as reported in [1]. A steel autoclave with a Teflon vessel (PARR) was loaded with 32.4 mg (0.1 mmol) of the protonated ligand, H₂(Me-4py-trz-ia), 25.0 mg (0.1 mmol) CuSO₄·5 H₂O and 5 ml of a 1:1 mixture of water/acetonitrile. After addition of 1 drop of 6 mol·l⁻¹ hydrochloric acid the autoclave was sealed, heated to 140 °C and kept at this temperature for five hours. After cooling the autoclave slowly to room temperature during a period of 60 hours blue crystals of **1a** were filtered off, washed with water/acetonitrile and stored in this solvent mixture. After drying at the air the yield was 49.3 mg blue crystals of **1a**.

Solvent exchange procedures:

For **1b**: After the synthesis of crystals of **1a** the solvent was exchanged with methanol 5 times over a period of 5 days. Then the crystals of **1b** were kept in methanol.

For **1c**: After the synthesis of crystals of **1a** the solvent was exchanged continuously by Soxhlet extraction with unhydrous methanol for 5 days. Then the crystals of **1c** were kept in methanol.

2. Single crystal structure data

Table SI-1. Single crystal structure data of **1a**, **1b**, **1c** and **1e**.

Compound	1a	1b	1c ^{*2}	1e
crystal size / mm	0.38 · 0.27 · 0.24	0.14 · 0.12 · 0.08	0.10 · 0.07 · 0.07	0.16 · 0.10 · 0.08
formula	C ₁₆ H ₁₂ N ₄ O ₅ Cu	C ₁₆ H ₁₂ N ₄ O ₅ Cu	C ₃₃ H ₂₄ N ₈ O ₉ Cu ₂	C ₁₆ H ₁₀ N ₄ O ₄ Cu
M / g·mol ⁻¹	403.84	403.84	803.68	385.82
temperature / K	213(2)	213(2)	150(2)	150(2)
diffractometer	STOE IPDS-1	STOE STADIVARI	STOE STADIVARI	STOE STADIVARI
wavelength / pm	71.073	154.186	154.186	154.186
crystal system	monoclinic	monoclinic	monoclinic	monoclinic
space group	<i>P2₁/c</i> (no. 14)	<i>P2₁/c</i> (no. 14)	<i>Pc</i> (no. 7)	<i>P2₁/c</i> (no. 14)
unit cell parameters / pm, °	a = 1425.1(2) b = 1339.9(2) c = 1463.7(2) β = 93.001(11)	a = 1386.63(4) b = 1323.50(4) c = 1502.41(4) β = 93.199(2)	a = 1371.36(4) b = 1322.70(5) c = 1517.95(5) β = 93.415(3)	a = 1355.18(4) b = 1391.61(4) c = 1482.83(4) β = 90.716(3)
volume / 10 ⁶ pm ³	2790.9(7)	2752.9(1)	2748.5(2)	2796.2(1)
Z	4	4	2	4
density / g·cm ³	0.961	0.974	0.971	0.916
absorption coefficient μ / mm ⁻¹	0.804	1.326	1.313	1.262
θ range / °	2.5 – 27.0	4.4 – 70.0	4.4 – 70.0	4.4 – 70.5
number of measured reflections	23693	29057	24796	33399
number of independent reflections	6048	5133	6716	5258
number of observed reflections	3520	3605	5142	3932
R _{int}	0.0576	0.0456	0.0493	0.0502
parameters	244	244	473	229
R ₁	0.0324 ^{*1}	0.0556 ^{*1}	0.0440 ^{*1}	0.0593
wR ₂	0.0740 ^{*1}	0.1965 ^{*1}	0.1166 ^{*1}	0.1772
max. / min. residual e ⁻ density / 10 ⁻⁶ pm ⁻³	0.35 / -0.25	0.40 / -0.64	0.47 / -0.46	1.00 / -0.71

^{*1} The PLATON/SQUEEZE-routine⁴ was applied.

^{*2} Refined as an inversion twin. Absolute structure parameter: 0.41(6).

3. Powder X-ray diffraction (PXRD)

The PXRD measurements were carried out on a STOE STADI-P diffractometer using Cu-K α_1 radiation ($\lambda = 154.060$ pm). The samples for these measurements were prepared in capillaries (outer diameter 0.5 mm). Activated samples were handled under inert conditions. The powder diffraction patterns were processed with WinXPow.⁶ Indexing of the diffraction patterns was performed using EXPO2014 via the program N-TREOR09.⁷ Subsequently, the selected unit cells were tested and refined with the Pawley method in TOPAS.⁸

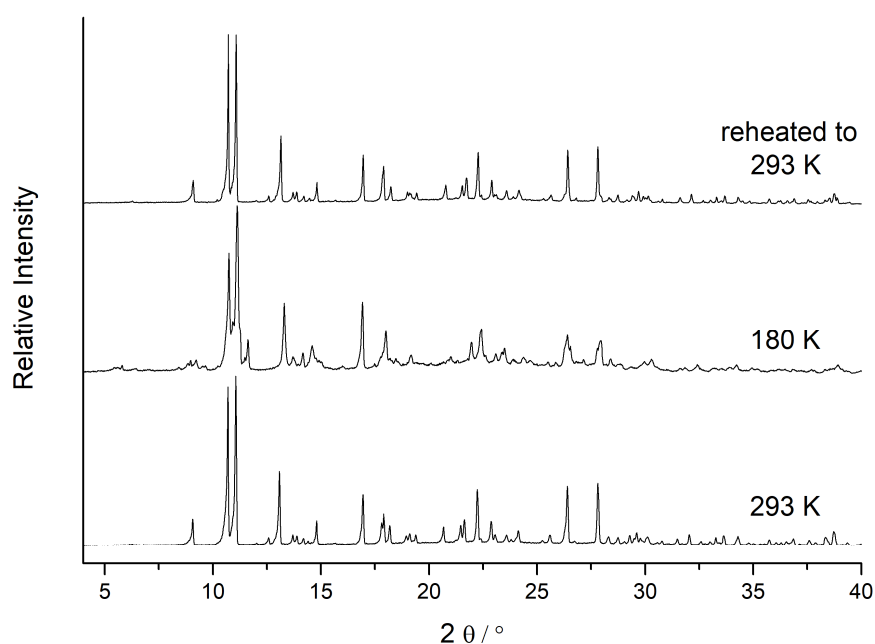


Figure SI-1. To demonstrate the influence of temperature powder diffraction patterns of an as-synthesized sample of **1a** were collected at 293 K, at 180 K and after warming up again to 293 K. Changes of the diffraction pattern at 180 K (e.g., peak splitting at $2\theta = 8^\circ$, additional peak at $2\theta = 11.5^\circ$) indicate reversible temperature-dependent structural changes of the as-synthesized material. This behaviour prevented a successful single crystal structure analysis at 180 K.

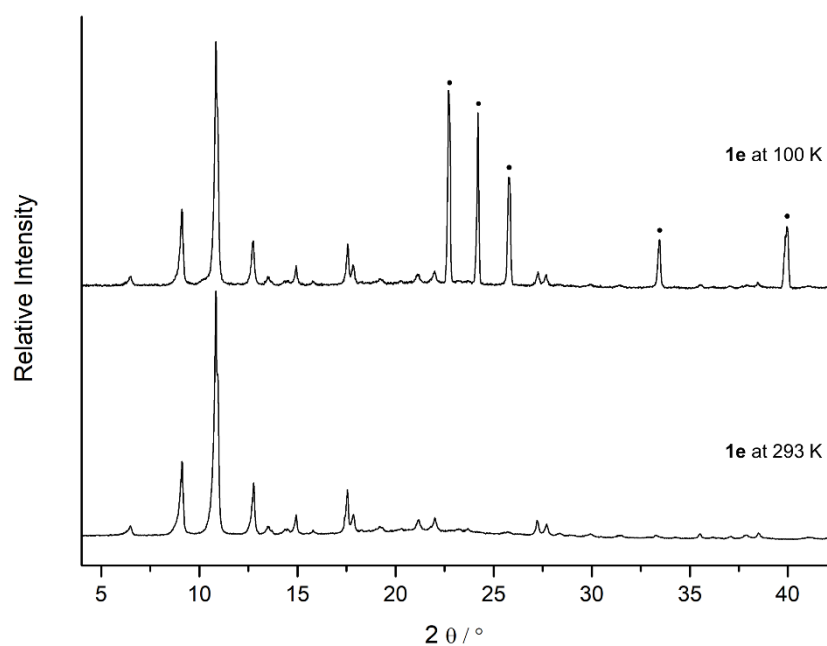


Figure SI-2. The PXRD pattern of the fully desolvated phase **1e** at 100 K is identical to that measured at room temperature, indicating no structural changes of the empty framework at lower temperatures. The additional peaks at 100 K (marked with black dots) belong to ice crystals that started to form at the outer wall of the glass capillary while cooling.

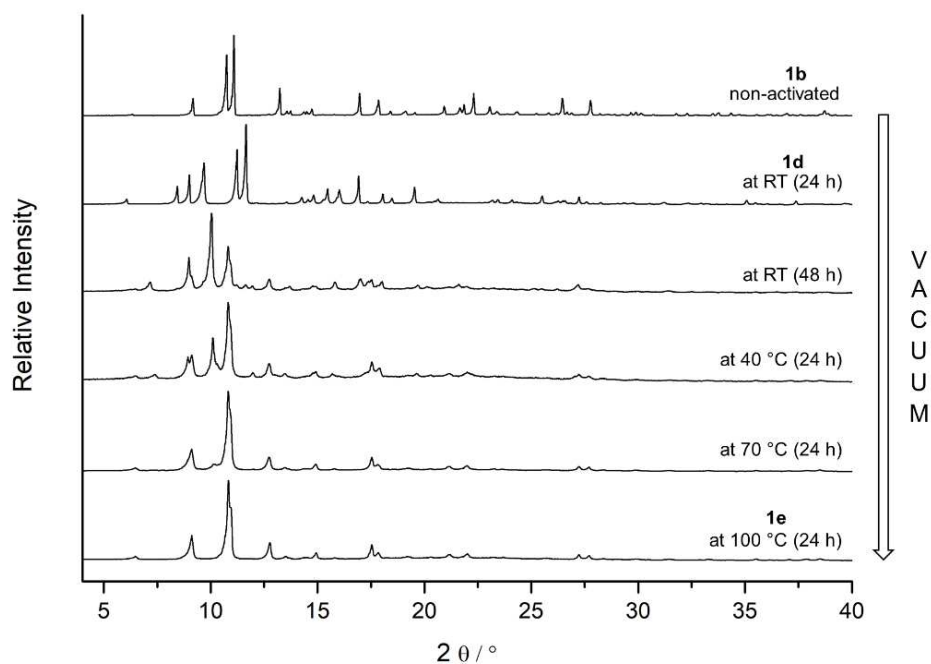


Figure SI-3. Room temperature X-ray powder diffraction patterns of sample **1b** (incomplete solvent exchange with methanol; pores are filled with methanol, two water molecules coordinating to Cu1) and after activation under vacuum at various temperatures or for different periods. The remaining water molecules at the Cu^{II} centres after removal of the non-coordinating solvent molecules from the pores cause the formation of the quite stable phase **1d** during activation in vacuum. Therefore, elevated temperatures or longer periods of activation are needed to obtain the fully desolvated phase **1e**. Since both **1a** and **1b** have to pass phase **1d**, they show similar structural transformations under vacuum. Such a behaviour is also observed for sample **1c** when exposed to humidity prior to activation.

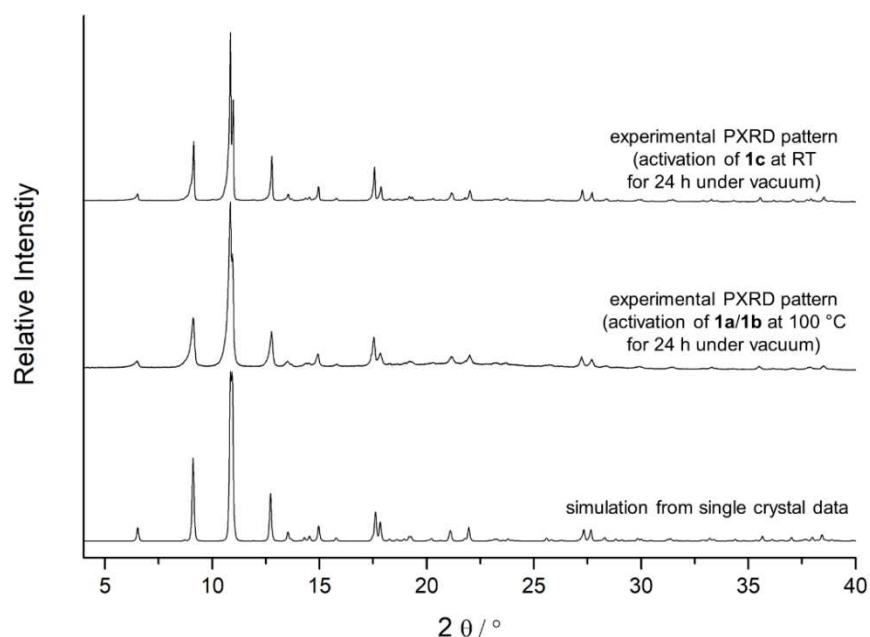


Figure SI-4. Comparison of the XRD pattern simulated from single crystal data of **1e** (a small crystal of **1e** was activated under vacuum at RT) (bottom) and experimental PXRd patterns of **1e** after activation of **1a/1b** for 24 h at 100 °C under vacuum (middle) and after activation of **1c** for 24 h at RT under vacuum (top). The peak at ca. 11° consists of two reflections, -1-11 and 111. Due to crystal fragmentation that occurs when activated along path A peaks are noticeably broader, -1-11 and 111 reflections are overlapping. When activated along path B (after Soxhlet extraction with methanol) the peaks are sharper and better resolved.

Table SI-2. Unit cell parameters and agreement factors of **1e** from indexing and subsequent Pawley refinement of PXRd pattern of an activated powder sample of **1b** (for 24 h under vacuum at 100 °C) and from single crystal structure analysis of an activated small crystal of **1e**.

Compound	1e	1e
Method	Pawley refinement	Single crystal analysis
Temperature	293 K	150 K
Space group	$P2_1/c$ (no. 14)	$P2_1/c$ (no. 14)
a / pm	1354.8(1)	1355.18(4)
b / pm	1383.9(1)	1391.61(4)
c / pm	1485.2(1)	1482.83(4)
$\beta / ^\circ$	90.928(3)	90.716(3)
V / 10^6 pm^3	2784.2(4)	2796.2(1)
Z	-	4
R_{int}	-	0.0502
R_1	-	0.0593
wR_2	-	0.1710
R_p	0.0322	-
R_{wp}	0.0422	-

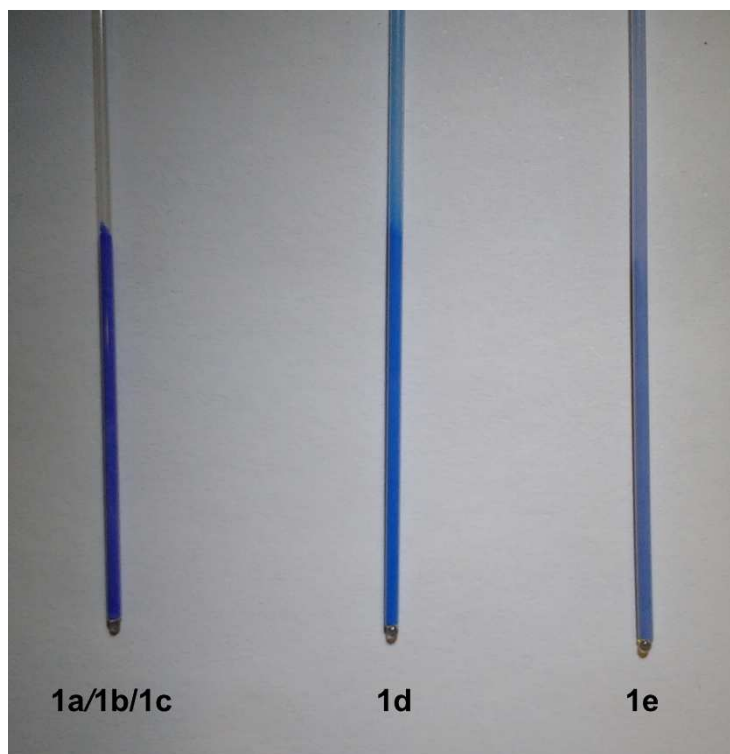


Figure SI-5. Comparison of the colours of the different samples of **1**. Air-dried samples **1a**, **1b** and **1c** showing the same shade of blue (left), partially activated **1d** (middle) and fully activated sample **1e** (right).

4. Structure determination of 1d from powder diffraction data

Space group determination and structure solution were performed with EXPO2014.⁷ A reasonable structure model could be obtained by the simulated annealing approach in the space group $P2_1$ using the derived lattice parameters from the indexing procedure. Assuming symmetry reduction, for the starting structure model two linker molecules and two copper ions were given as fragments. Since the activation is obviously not completed at this stage two water molecules were added as fragments to the starting model. Torsion angles of the linker molecules were not restrained during this process. The obtained structure model was refined by the Rietveld method using TOPAS.⁸ In the course of refinement the linker molecules were treated as rigid bodies using the Z-matrix notation allowing translation of the molecules and rotation of the phenyl and pyridine rings at the triazole group as well as the carboxylate groups at the phenyl ring (torsion angles to be refined freely). Atomic positions of the copper ions and the water molecules were not constrained/restrained during the refinement.

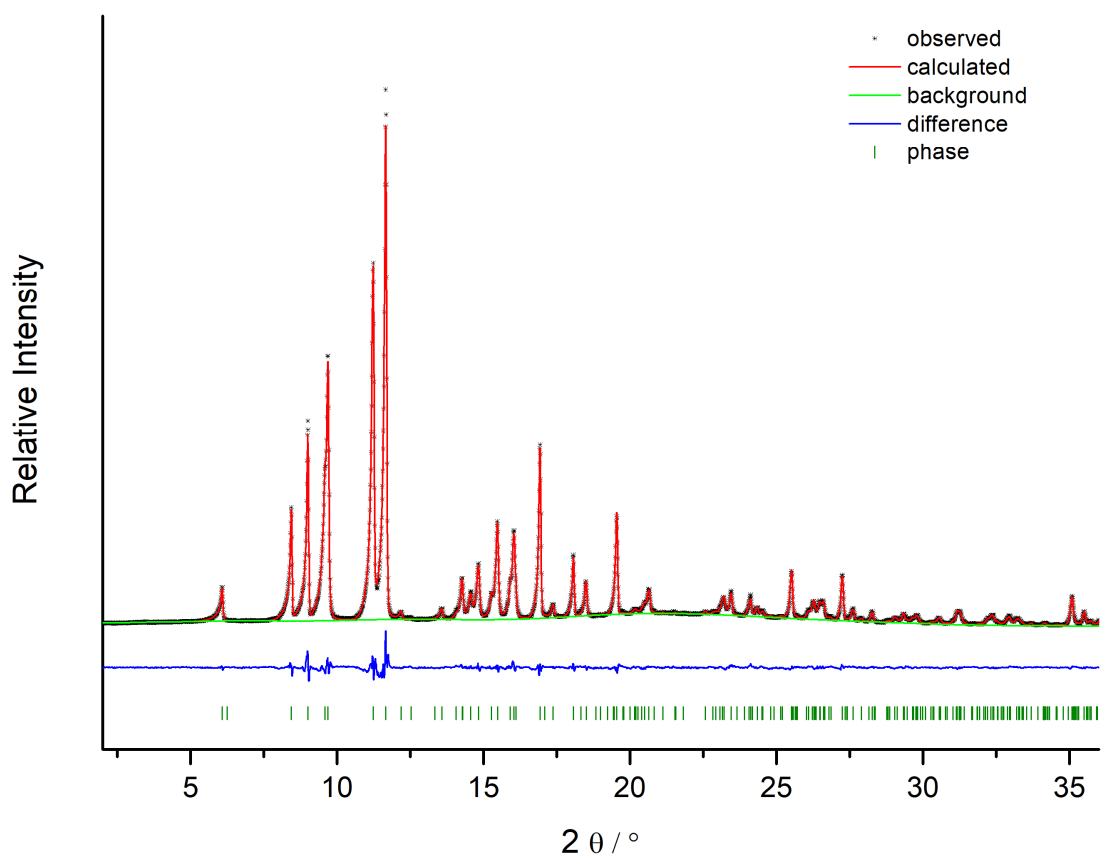
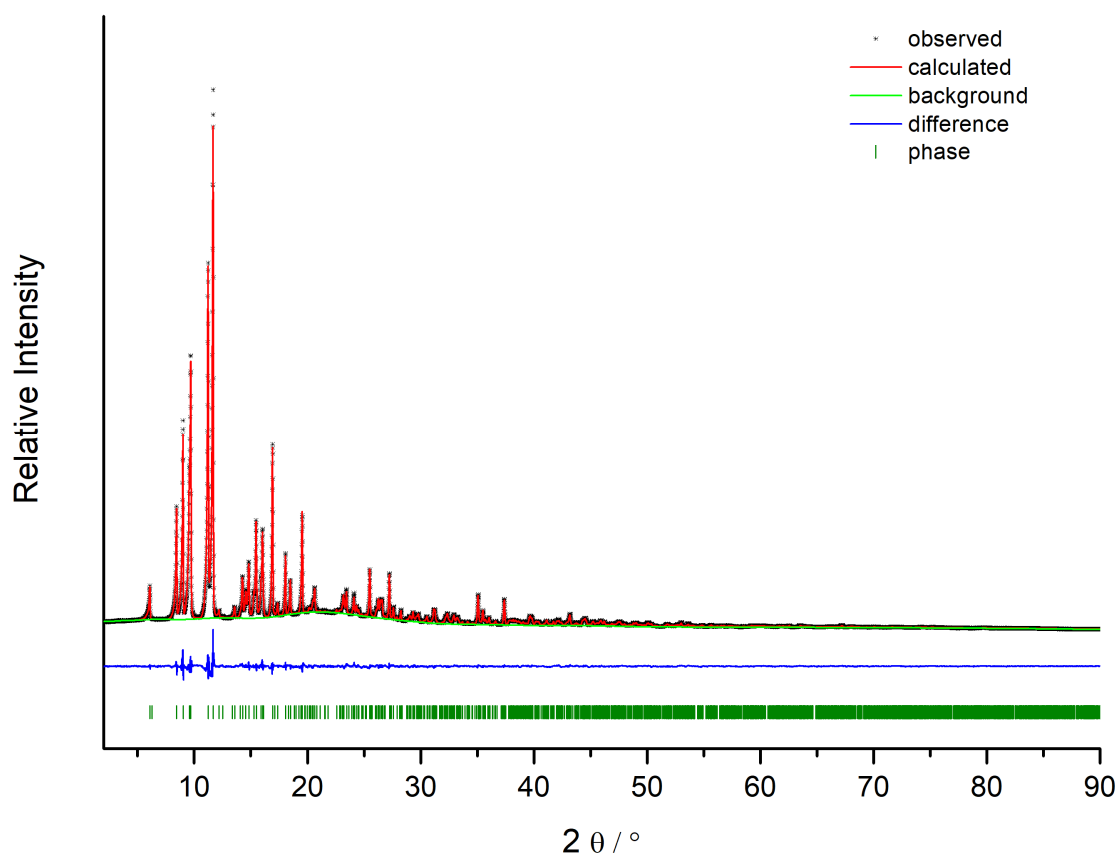
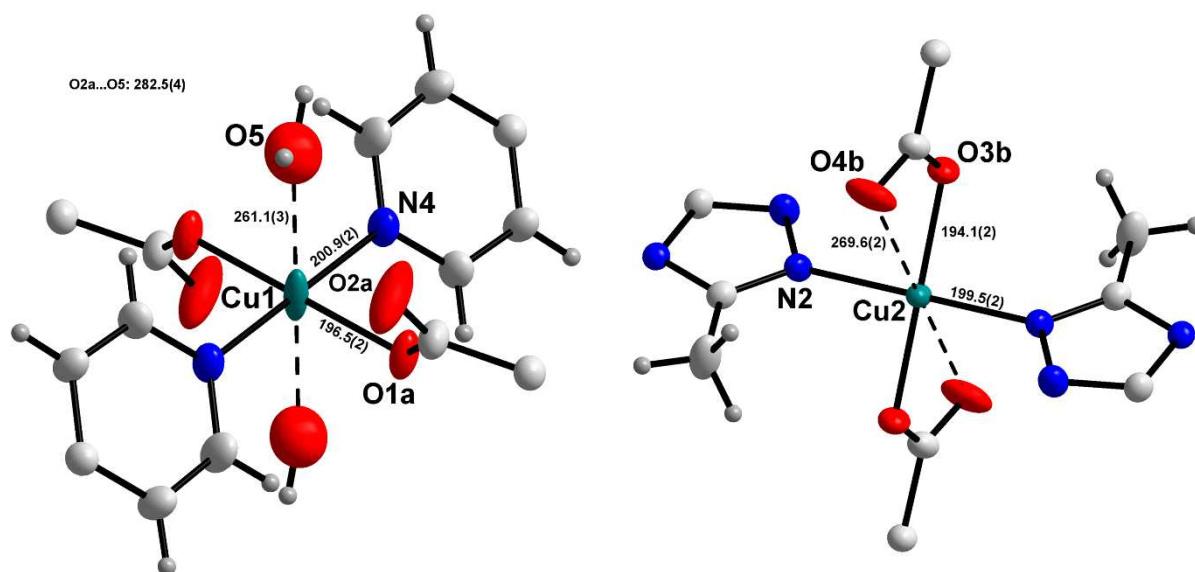
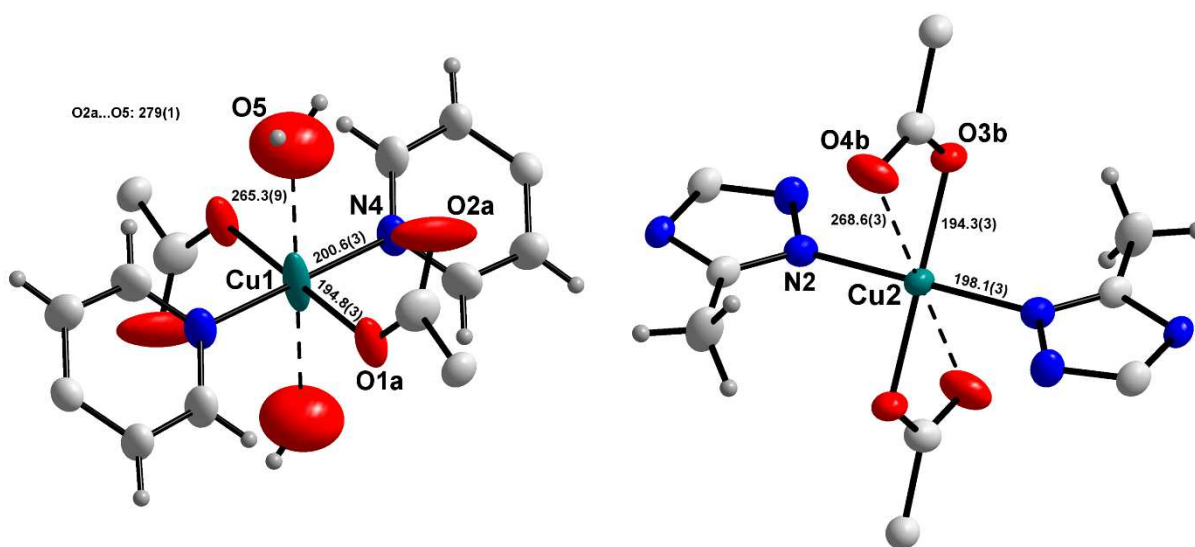


Figure SI-6. Rietveld refinement of phase **1b**. Space group $P2_1$ (no. 4); $a = 1454.93(3)$ pm, $b = 1193.95(1)$ pm, $c = 1415.17(3)$ pm, $\beta = 93.6939(5)^\circ$; $Z = 2$; $V = 2453.20(7) \cdot 10^6$ pm³; $R_p = 0.0286$, $R_{wp} = 0.0379$).

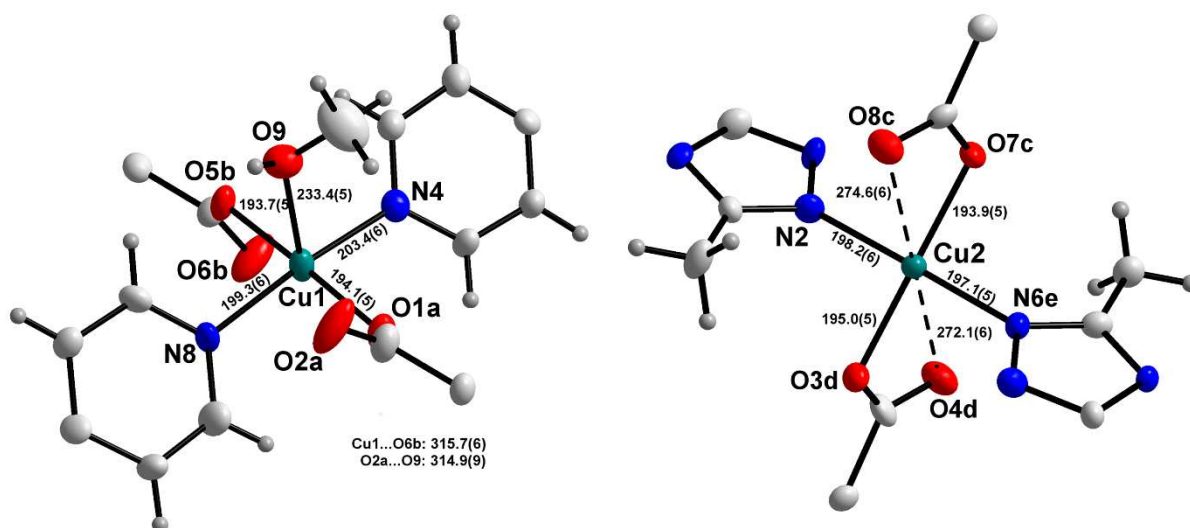
5. Coordination modes of the Cu²⁺ centres Cu1 and Cu2



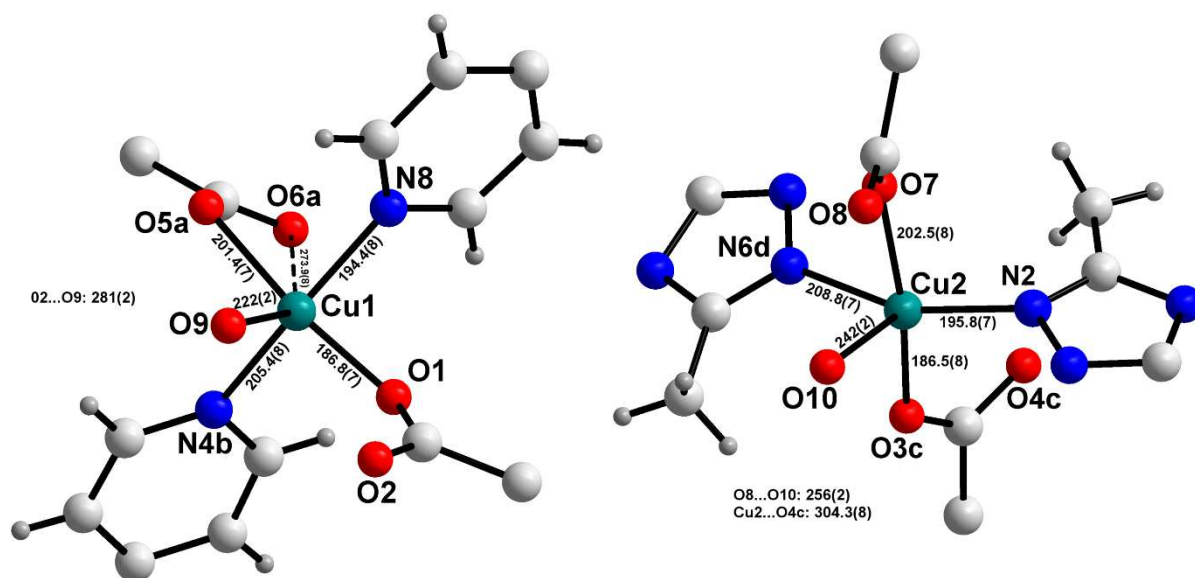
(a) **1a** (single crystal data). Symmetry codes: a: 1-x, -0.5+y, 0.5-z; b: -x, -0.5+y, 0.5-z.



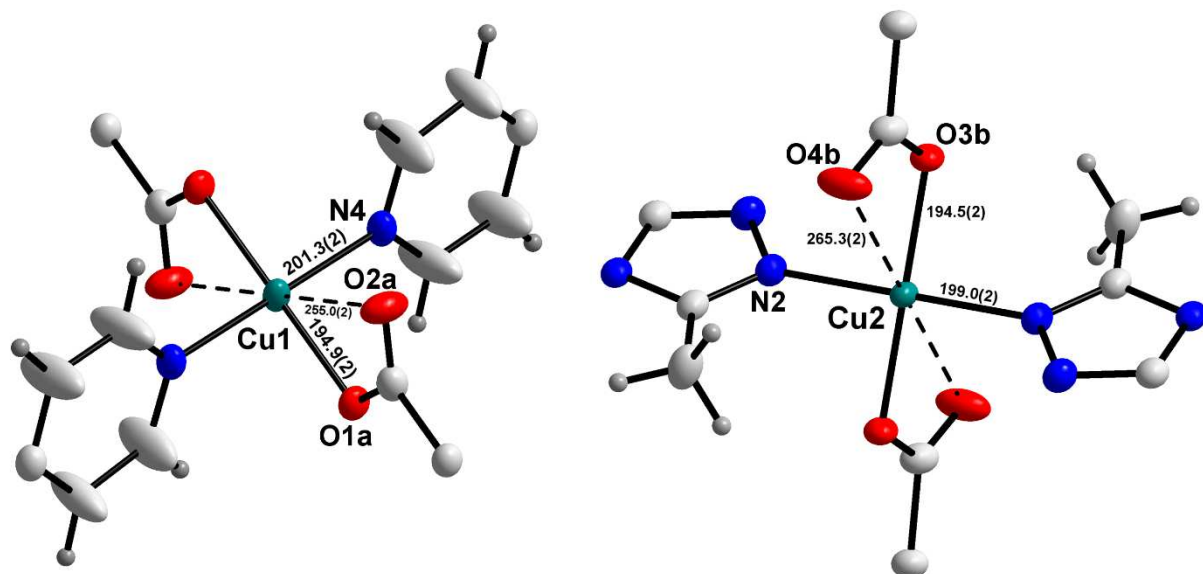
(b) **1b** (single crystal data). Symmetry codes: a: 1-x, -0.5+y, 0.5-z; b: -x, -0.5+y, 0.5-z.



(c) **1c** (single crystal data). Symmetry codes: a: $x, 1-y, 0.5+z$; b: $x, -y, -0.5+z$; c: $-1+x, -y, -0.5+z$; d: $x, 1-y, -0.5+z$; e: $-1+x, y, -1+z$.



(d) **1d** (powder data). Symmetry codes: a: $-x, -0.5+y, 1-z$; b: $-x, -0.5+y, -z$; c: $1-x, 0.5+y, -z$; d: $1-x, 0.5+y, 1-z$.



(e) **1e** (single crystal data). Symmetry codes: a: $1-x, -0.5+y, 0.5-z$; b: $-x, -0.5+y, 0.5-z$.

Figure SI-7. Fragments of the crystals structures of **1a** -**1e** showing the coordination environment of the copper atoms. The ellipsoids drawn for the structures of **1a**, **1b**, **1c** and **1e** represent 50% probability.

6. Temperature dependent PXRD patterns

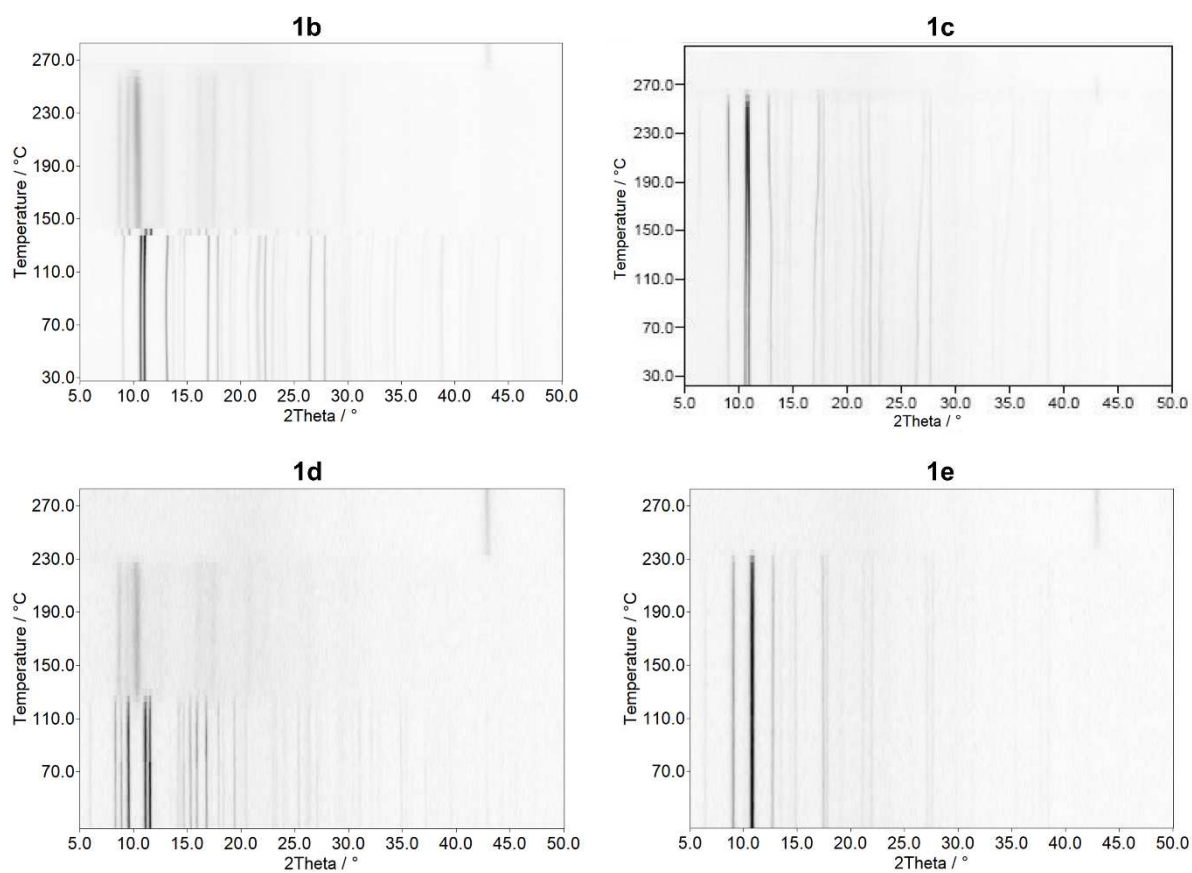


Figure SI-8. Temperature-dependent PXRD patterns of samples **1b**, **1c**, **1d** and **1e**. The diagram for **1b** indicates the formation of phase **1d** at ca. 140 °C, which quickly transforms to a high-temperature phase, while the diagram for **1c** shows a continuous cell transformation from **1c** to the fully desolvated material **1e**. This is in agreement with the observed behaviour under vacuum. [The reason for the higher thermal stabilities of samples **1b** and **1c** in comparison to that of **1d** and **1e** is not fully understood yet.]

7. DFT calculations

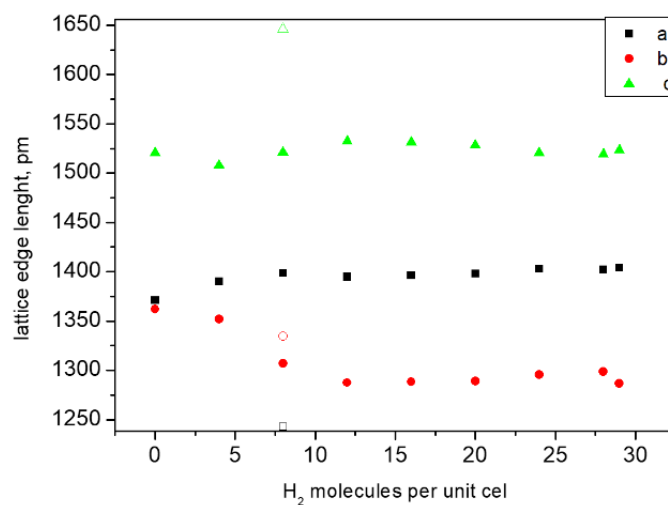


Figure SI-9. Unit cell parameter response (DFT simulations) upon H₂ loading (filled symbols) and H₂O chemisorbed at Cu1 and Cu2 sites (open symbols).

Table SI-3. Comparison of the unit cell parameters of the simulated structures.

	Simulated empty framework	Simulated structure with 2 H ₂ O at Cu1	Simulated structure with 2 H ₂ O each at Cu1 and Cu2	Simulated structure with 1 H ₂ O each at Cu1 and Cu2	Simulated structure with 1 H ₂ O each at Cu1 and Cu2	Simulated structure with 29 H ₂ in unit cell
Space group	<i>P2₁/c</i>	<i>P2₁/c</i>	<i>P2₁/c</i>	<i>P2₁</i>	<i>P1</i>	<i>P1</i>
a / pm	1371.2	1389.6	1242.6	1484.2	1485.8	1404.0
b / pm	1361.9	1306.6	1334.3	1197.0	1196.5	1286.6
c / pm	1520.3	1541.4	1646.0	1396.1	1394.7	1523.2
α / °	-	-	-	-	90.00	90.00
β / °	92.31	94.00	93.39	94.26	94.27	93.29
γ / °	-	-	-	-	90.00	90.00
V / 10 ⁶ pm ³	2836.9	2791.7	2724.5	2473.5	2472.5	2747.0

CIFs of simulated structures:

Simulated empty framework:

```
data_simulation_empty_framework
_symmetry_space_group_name_H-M 'P21/c'
_symmetry_Int_Tables_number 14
_symmetry_cell_setting
monoclinic
loop_
_symmetry_equiv_pos_as_xyz
  x,y,z
  -x,y+1/2,-z+1/2
  -x,-y,-z
  x,-y+1/2,z+1/2
_cell_length_a 13.7121
_cell_length_b 13.6194
_cell_length_c 15.2034
_cell_angle_alpha 90.0000
_cell_angle_beta 92.3109
_cell_angle_gamma 90.0000
loop_
_atom_site_label
_atom_site_type_symbol
_atom_site_fract_x
_atom_site_fract_y
_atom_site_fract_z
Cu1 Cu 0.5 0 0.5
Cu2 Cu 0 0 0
O1 O 0.42225 0.41761 0.07864
O2 O 0.36956 0.56788 0.11997
O3 O 0.05991 0.42636 0.40359
O4 O 0.12695 0.57254 0.36656
N1 N 0.15422 0.18561 0.16648
N2 N 0.0882 0.06842 0.08655
N3 N 0.15305 0.02483 0.14308
N4 N 0.40289 0.02963 0.4004
C1 C 0.3649 0.47656 0.12069
C2 C 0.2863 0.42486 0.17087
C3 C 0.2598 0.32776 0.1495
C4 C 0.18281 0.28355 0.19252
C5 C 0.13336 0.33249 0.25833
C6 C 0.16176 0.4288 0.2814
C7 C 0.2366 0.47518 0.23632
C8 C 0.11307 0.48283 0.35555
C9 C 0.02236 0.23702 0.05292
C10 C 0.08734 0.16515 0.09973
C11 C 0.19426 0.09568 0.19204
C12 C 0.26762 0.07689 0.26286
C13 C 0.2983 0.14771 0.32585
C14 C 0.36515 0.12072 0.39323
C15 C 0.37574 0.96161 0.33952
C16 C 0.30845 0.98175 0.27064
H1 H 0.29859 0.28798 0.09925
H2 H 0.0741 0.29689 0.29214
H3 H 0.25616 0.55103 0.25244
H4 H 0.04609 0.31272 0.06696
H5 H 0.94733 0.22591 0.0748
H6 H 0.02246 0.22239 0.98169
H7 H 0.27053 0.22263 0.32425
H8 H 0.38915 0.1727 0.44431
H9 H 0.41022 0.88964 0.34717
H10 H 0.28701 0.92466 0.22331
```

Simulated structure with 2 H₂O at Cu1 in P2₁/c:

```
data_simulation_2H2O_at_Cu1_inP21/c
_symmetry_space_group_name_H-M 'P21/C'
_symmetry_Int_Tables_number 14
_symmetry_cell_setting
monoclinic
loop_
_symmetry_equiv_pos_as_xyz
  x,y,z
  -x,y+1/2,-z+1/2
  -x,-y,-z
  x,-y+1/2,z+1/2
_cell_length_a 13.8955
_cell_length_b 13.0655
_cell_length_c 15.4143
_cell_angle_alpha 90.0000
_cell_angle_beta 93.9945
_cell_angle_gamma 90.0000
loop_
_atom_site_label
_atom_site_type_symbol
_atom_site_fract_x
_atom_site_fract_y
_atom_site_fract_z
Cu1 Cu 0.5 0 0.5
Cu2 Cu 0 0 0
O1 O 0.40386 0.44855 0.07927
O2 O 0.36655 0.5994 0.14375
O3 O 0.06057 0.4301 0.4038
O4 O 0.12645 0.58522 0.37726
N1 N 0.15641 0.19411 0.16481
N2 N 0.09038 0.07157 0.08524
N3 N 0.15446 0.02634 0.14294
N4 N 0.40728 0.03423 0.39857
C1 C 0.35676 0.50502 0.13069
C2 C 0.28077 0.44714 0.1776
C3 C 0.25629 0.34639 0.15255
C4 C 0.1827 0.29605 0.19303
C5 C 0.13406 0.34254 0.25913
C6 C 0.16063 0.44214 0.28604
C7 C 0.23271 0.49459 0.24414
C8 C 0.11291 0.49259 0.36098
C9 C 0.0275 0.24759 0.04791
C10 C 0.09031 0.17255 0.09728
C11 C 0.19559 0.10044 0.19172
C12 C 0.26942 0.08156 0.26265
C13 C 0.29512 0.15328 0.32866
C14 C 0.36381 0.12636 0.39528
C15 C 0.38452 0.96518 0.33528
C16 C 0.31618 0.98533 0.2668
H1 H 0.29429 0.30887 0.10164
H2 H 0.07704 0.30219 0.29051
H3 H 0.25139 0.57287 0.26361
H4 H 0.02998 0.23242 0.9778
H5 H 0.05067 0.32643 0.063
H6 H 0.95255 0.23613 0.06538
H7 H 0.26238 0.22895 0.32988
H8 H 0.38642 0.17922 0.44767
H9 H 0.42327 0.89243 0.34108
H10 H 0.29871 0.9276 0.21699
Ow O 0.55584 0.19207 0.48934
H1w H 0.607 0.22916 0.52173
H2w H 0.58574 0.16497 0.43662
```

Simulated structure with 2 H₂O at Cu1 and Cu2 in *P*₂/c:

data_simulation_2H2O_at_Cu1andCu2_inP21/c		C1	C	0.35713	0.49245	0.12927
_symmetry_space_group_name_H-M	'P21/C'	C2	C	0.28158	0.43276	0.17972
_symmetry_Int_Tables_number	14	C3	C	0.24877	0.33628	0.15527
_symmetry_cell_setting	monoclinic	C4	C	0.17638	0.28374	0.20141
loop_		C5	C	0.136	0.32431	0.27199
_symmetry_equiv_pos_as_xyz		C6	C	0.17087	0.42057	0.29709
x,y,z		C7	C	0.24272	0.47442	0.25062
-x,y+1/2,-z+1/2		C8	C	0.13206	0.46914	0.37355
-x,-y,-z		C9	C	0.02155	0.24492	0.05611
x,-y+1/2,z+1/2		C10	C	0.0825	0.16705	0.10414
_cell_length_a	12.4263	C11	C	0.18816	0.09253	0.19807
_cell_length_b	13.3432	C12	C	0.26168	0.07278	0.26942
_cell_length_c	16.4603	C13	C	0.32667	0.14756	0.30804
_cell_angle_alpha	90.0000	C14	C	0.39342	0.12226	0.37614
_cell_angle_beta	93.3873	C15	C	0.33779	0.9548	0.36917
_cell_angle_gamma	90.0000	C16	C	0.26854	0.97439	0.30107
loop_		H1	H	0.28054	0.30266	0.10111
_atom_site_label		H2	H	0.0796	0.28282	0.3078
_atom_site_type_symbol		H3	H	0.26831	0.54944	0.26992
_atom_site_fract_x		H4	H	0.95932	0.20985	0.01507
_atom_site_fract_y		H5	H	0.07655	0.28856	0.01953
_atom_site_fract_z		H6	H	0.98287	0.29709	0.09734
Cu1 Cu 0.5 -2.83186E-33 0.5		H7	H	0.3278	0.22438	0.28588
Cu2 Cu 0 0 0		H8	H	0.44551	0.17772	0.40777
O1 O 0.39669 0.44117 0.07092		H9	H	0.34542	0.88059	0.39765
O2 O 0.37422 0.58263 0.14594		H10	H	0.21988	0.91426	0.27283
O3 O 0.07028 0.41491 0.41619		Ow	O	0.61705	0.20975	0.48477
O4 O 0.16088 0.55767 0.38955		H1w	H	0.67214	0.26215	0.47939
N1 N 0.14699 0.18487 0.17304		H2w	H	0.62417	0.16726	0.43628
N2 N 0.08483 0.06905 0.08946		Ow	O	0.79322	0.15148	0.96549
N3 N 0.14937 0.02225 0.14702		H3w	H	0.79889	0.10426	0.92055
N4 N 0.39808 0.0281 0.4056		H4w	H	0.19371	0.61166	0.48397

Simulated structure with 1 H₂O at Cu1 and Cu2 in *P*₂:

data_simulation_1H2O_at_Cu1andCu2_inP21		N61	N	0.8136	0.89852	0.10216
_symmetry_space_group_name_H-M	'P21'	N62	N	0.85052	0.05497	0.17484
_symmetry_Int_Tables_number	4	N63	N	0.78659	0.08047	0.10446
_symmetry_cell_setting	monoclinic	N64	N	0.53893	0.97159	0.84928
loop_		C1	C	0.43357	0.52268	0.37135
_symmetry_equiv_pos_as_xyz		C2	C	0.3482	0.49915	0.42142
x,y,z		C3	C	0.2957	0.40485	0.39461
-x,y+1/2,-z		C5	C	0.21279	0.38962	0.4345
_cell_length_a	14.8415	C6	C	0.1824	0.46398	0.50229
_cell_length_b	11.9701	C8	C	0.23636	0.55638	0.53141
_cell_length_c	13.9614	C9	C	0.31843	0.57468	0.48961
_cell_angle_alpha	90.0000	C11	C	0.2038	0.6346	0.60724
_cell_angle_beta	94.2569	C12	C	0.05747	0.40628	0.28096
_cell_angle_gamma	90.0000	C16	C	0.08366	0.30472	0.33773
loop_		C17	C	0.1567	0.18709	0.43709
_atom_site_label		C18	C	0.22201	0.1362	0.5082
_atom_site_type_symbol		C19	C	0.227	0.01854	0.51445
_atom_site_fract_x		C21	C	0.289	0.96905	0.58184
_atom_site_fract_y		C23	C	0.3409	0.14254	0.63642
_atom_site_fract_z		C25	C	0.28081	0.19877	0.57165
Cu53 Cu 0.43499 0.97105 0.74624		C27	C	0.7054	0.50634	0.11705
Cu54 Cu 0.93339 0.1622 0.24734		C28	C	0.76791	0.59272	0.07813
O55 O 0.39085 0.7943 0.78046		C29	C	0.76245	0.70434	0.10888
O56 O 0.8514 0.32232 0.24178		C31	C	0.81975	0.7839	0.07266
O65 O 0.47564 0.43305 0.34606		C32	C	0.88272	0.75567	0.00663
O66 O 0.45499 0.62121 0.3551		C34	C	0.88744	0.64443	0.97511
O67 O 0.1235 0.61488 0.63071		C35	C	0.83047	0.5632	0.01162
O68 O 0.25591 0.71142 0.64062		C37	C	0.95101	0.61092	0.8989
O69 O 0.63984 0.5483 0.16074		C38	C	0.92953	0.88224	0.24362
O70 O 0.72094 0.40375 0.10516		C42	C	0.86716	0.94541	0.17521
O71 O 0.99706 0.69344 0.86661		C43	C	0.76302	0.98556	0.05971
O72 O 0.9528 0.51146 0.87271		C44	C	0.68906	0.97754	0.98427
N57 N 0.15572 0.29692 0.40477		C45	C	0.66688	0.87987	0.93066
N58 N 0.0436 0.20517 0.3317		C47	C	0.59155	0.88084	0.86421
N59 N 0.08809 0.13206 0.39219		C51	C	0.63401	0.07286	0.96625
N60 N 0.34487 0.03017 0.64176		C49	C	0.56007	0.06603	0.89888
		H4	H	0.3196	0.34544	0.34278

H7	H	0.1176	0.45134	0.533	H39	H	0.89235	0.81576	0.27881
H10	H	0.3592	0.64823	0.50955	H40	H	0.98411	0.84429	0.2054
H13	H	0.07371	0.48221	0.32327	H41	H	0.95983	0.94036	0.2978
H14	H	0.09457	0.40924	0.21533	H46	H	0.70624	0.80334	0.94003
H15	H	0.98462	0.40235	0.25974	H48	H	0.57086	0.80718	0.82179
H20	H	0.18294	0.96676	0.46671	H50	H	0.51483	0.13689	0.88406
H22	H	0.29403	0.87822	0.58867	H52	H	0.64867	0.15014	0.00599
H24	H	0.38891	0.18796	0.68492	H73	H	0.35347	0.75483	0.73042
H26	H	0.28113	0.28974	0.57145	H74	H	0.34811	0.81929	0.82785
H30	H	0.71306	0.72786	0.15945	H75	H	0.80603	0.29268	0.28433
H33	H	0.92733	0.81889	0.97928	H76	H	0.81653	0.34263	0.18163
H36	H	0.8347	0.47689	0.9869					

Simulated structure with 1 H₂O at Cu1 and Cu2 in P1

data_simulation_1H2O_at_Cu1andCu2_inP1	C35	C	0.83013	0.56286	0.01147				
_symmetry_space_group_name_H-M	C37	C	0.9509	0.61049	0.89902				
_symmetry_Int_Tables_number	C38	C	0.92959	0.88189	0.24384				
_symmetry_cell_setting	C42	C	0.86727	0.94515	0.17539				
loop_	C43	C	0.76298	0.98535	0.06002				
_symmetry_equiv_pos_as_xyz	C44	C	0.68895	0.97735	0.98465				
x,y,z	C45	C	0.66714	0.87993	0.93044				
_cell_length_a	C47	C	0.59184	0.88096	0.86397				
_cell_length_b	C51	C	0.63354	0.07246	0.96727				
_cell_length_c	C49	C	0.55966	0.06569	0.89984				
_cell_angle_alpha	H4	H	0.3195	0.34571	0.34257				
_cell_angle_beta	H7	H	0.11747	0.45171	0.53264				
_cell_angle_gamma	H10	H	0.35886	0.6487	0.50948				
loop_	H13	H	0.07264	0.48233	0.32356				
_atom_site_label	H14	H	0.0947	0.41016	0.2155				
_atom_site_type_symbol	H15	H	0.98452	0.40188	0.25905				
_atom_site_fract_x	H20	H	0.18326	0.96684	0.46657				
_atom_site_fract_y	H22	H	0.29431	0.87841	0.58862				
_atom_site_fract_z	H24	H	0.3897	0.18836	0.68413				
Cu53	Cu	0.4352	0.97123	0.74617	H26	H	0.28192	0.29008	0.57055
Cu54	Cu	0.93358	0.16209	0.2473	H30	H	0.71285	0.72769	0.15939
O55	O	0.39107	0.79434	0.78032	H33	H	0.92738	0.81848	0.97963
O56	O	0.85132	0.32204	0.24095	H36	H	0.83427	0.47653	0.98664
O65	O	0.47533	0.43336	0.34602	H39	H	0.89237	0.81564	0.27929
O66	O	0.45471	0.62162	0.35504	H40	H	0.98392	0.8436	0.20551
O67	O	0.12325	0.61534	0.63045	H41	H	0.96013	0.94006	0.2979
O68	O	0.25562	0.71171	0.64076	H46	H	0.70679	0.80358	0.93934
O69	O	0.63977	0.54822	0.16086	H48	H	0.57147	0.80753	0.82104
O70	O	0.72031	0.40346	0.10477	H50	H	0.51414	0.13637	0.88549
O71	O	0.99705	0.69304	0.8669	H52	H	0.64789	0.14952	0.00753
O72	O	0.95268	0.511	0.87273	H73	H	0.35358	0.75493	0.7303
N57	N	0.1558	0.29712	0.4044	H74	H	0.34852	0.81931	0.82792
N58	N	0.04377	0.20513	0.33151	H75	H	0.8061	0.29255	0.28374
N59	N	0.08849	0.13204	0.3919	H76	H	0.81634	0.34208	0.18075
N60	N	0.34538	0.03047	0.64139	Cu53	Cu	0.5648	0.47121	0.25382
N61	N	0.81361	0.89827	0.10237	Cu54	Cu	0.06642	0.6621	0.75271
N62	N	0.85067	0.05476	0.17506	O55	O	0.60894	0.29432	0.21967
N63	N	0.78663	0.0803	0.10477	O56	O	0.14868	0.82204	0.75906
N64	N	0.5389	0.97152	0.8496	O65	O	0.52468	0.93338	0.65398
C1	C	0.43329	0.52305	0.37127	O66	O	0.5453	0.12164	0.64497
C2	C	0.34796	0.49954	0.42128	O67	O	0.87675	0.11535	0.36956
C3	C	0.29555	0.40519	0.39437	O68	O	0.74438	0.2117	0.35924
C5	C	0.21268	0.38997	0.43416	O69	O	0.36023	0.04823	0.83914
C6	C	0.18223	0.46437	0.50196	O70	O	0.27971	0.90347	0.89524
C8	C	0.23611	0.55681	0.5312	O71	O	0.00295	0.19303	0.13311
C9	C	0.31814	0.57511	0.48948	O72	O	0.04731	0.01099	0.12726
C11	C	0.20354	0.63499	0.60713	N57	N	0.84422	0.79714	0.5956
C12	C	0.05716	0.40644	0.28083	N58	N	0.95624	0.70516	0.66849
C16	C	0.08363	0.30481	0.33751	N59	N	0.91153	0.63207	0.60809
C17	C	0.15704	0.18721	0.43669	N60	N	0.65463	0.53047	0.3586
C18	C	0.2225	0.13641	0.50773	N61	N	0.18637	0.39825	0.89763
C19	C	0.2274	0.01871	0.51417	N62	N	0.14932	0.55475	0.82494
C21	C	0.28941	0.96927	0.58161	N63	N	0.21337	0.58028	0.89522
C23	C	0.34156	0.14289	0.6358	N64	N	0.46109	0.47149	0.15039
C25	C	0.28147	0.19907	0.57096	C1	C	0.56672	0.02307	0.62873
C27	C	0.70504	0.50612	0.11689	C2	C	0.65205	0.99956	0.57873
C28	C	0.76758	0.59244	0.07797	C3	C	0.70446	0.90522	0.60564
C29	C	0.76222	0.70409	0.10883	C5	C	0.78733	0.89	0.56584
C31	C	0.81965	0.78361	0.07277	C6	C	0.81778	0.96438	0.49804
C32	C	0.88265	0.75531	0.00681	C8	C	0.7639	0.05682	0.46881
C34	C	0.88725	0.64404	0.97515	C9	C	0.68187	0.07513	0.51053

C11	C	0.79647	0.135	0.39287	H4	H	0.68052	0.84574	0.65744
C12	C	0.94284	0.90646	0.71919	H7	H	0.88254	0.95173	0.46736
C16	C	0.91637	0.80484	0.6625	H10	H	0.64115	0.14871	0.49052
C17	C	0.84297	0.68724	0.5633	H13	H	0.92734	0.98235	0.67647
C18	C	0.77751	0.63643	0.49226	H14	H	0.9053	0.91018	0.78452
C19	C	0.77261	0.51873	0.48582	H15	H	0.01548	0.90193	0.74097
C21	C	0.71061	0.46928	0.41838	H20	H	0.81676	0.46688	0.53342
C23	C	0.65845	0.64289	0.36419	H22	H	0.70571	0.37842	0.41138
C25	C	0.71854	0.69909	0.42902	H24	H	0.61031	0.68836	0.31586
C27	C	0.29496	0.00613	0.88312	H26	H	0.7181	0.79009	0.42943
C28	C	0.23242	0.09244	0.92204	H30	H	0.28713	0.22769	0.84061
C29	C	0.23776	0.20409	0.89117	H33	H	0.07259	0.31846	0.02035
C31	C	0.18033	0.28359	0.92722	H36	H	0.16576	0.97653	0.01339
C32	C	0.11733	0.25529	0.99318	H39	H	0.10757	0.31562	0.72072
C34	C	0.11275	0.14403	0.02486	H40	H	0.01604	0.3436	0.7945
C35	C	0.16988	0.06286	0.98855	H41	H	0.03983	0.44004	0.70211
C37	C	0.04909	0.11047	0.10098	H46	H	0.29317	0.30357	0.06067
C38	C	0.07036	0.38188	0.75618	H48	H	0.42849	0.30751	0.17896
C42	C	0.1327	0.44514	0.82461	H50	H	0.48589	0.63632	0.11446
C43	C	0.23701	0.48533	0.93997	H52	H	0.35214	0.64947	0.99243
C44	C	0.31105	0.47732	0.01534	H73	H	0.64643	0.25492	0.26969
C45	C	0.33284	0.37992	0.06956	H74	H	0.6515	0.3193	0.17207
C47	C	0.40814	0.38094	0.13602	H75	H	0.1939	0.79256	0.71627
C51	C	0.36648	0.57242	0.0327	H76	H	0.18367	0.84208	0.81926
C49	C	0.44036	0.56565	0.10013					

Simulated structure with 29 H₂ in unit cell in P1

data_simulation_29H2_in_unit_cell_inP1					H3	H	0.25944	0.57547	0.25875
_symmetry_space_group_name_H-M				'P1'	H4	H	0.06611	0.32702	0.05357
_symmetry_Int_Tables_number				1	H5	H	0.96112	0.2524	0.07454
_symmetry_cell_setting				triclinic	H6	H	0.02488	0.22568	0.9789
loop_					H7	H	0.25424	0.22326	0.33531
_symmetry_equiv_pos_as_xyz					H8	H	0.37516	0.17326	0.45328
x,y,z					H9	H	0.42003	0.88539	0.34049
_cell_length_a				14.0400	H10	H	0.29588	0.91958	0.21788
_cell_length_b				12.8660	H11	H	0.56096	0.19724	0.47871
_cell_length_c				15.2321	H12	H	0.53727	0.21782	0.51841
_cell_angle_alpha				89.99996	H13	H	0.89532	0.19831	0.77257
_cell_angle_beta				93.29066	H4	H	0.89038	0.24688	0.74498
_cell_angle_gamma				89.99978	H15	H	0.52527	0.30473	0.74505
loop_					H16	H	0.49188	0.2737	0.71472
_atom_site_label					H17	H	0.71403	0.64464	0.50917
_atom_site_type_symbol					H18	H	0.76554	0.635	0.52308
_atom_site_fract_x					H19	H	0.80452	0.45298	0.04138
_atom_site_fract_y					H20	H	0.77739	0.4272	0.00342
_atom_site_fract_z					H21	H	0.26783	0.69957	0.37299
Cu1	Cu	0.50011	-1.62175E-4	0.49981	H22	H	0.31676	0.72362	0.37608
Cu2	Cu	8.3861E-5	3.5056E-5	-2.57229E-4	H23	H	0.55387	0.65617	0.64942
O1	O	0.4164	0.43358	0.08231	H24	H	0.58929	0.6124	0.65171
O2	O	0.38057	0.59196	0.13822	Cu1	Cu	0.5001	0.50008	-1.42322E-4
O3	O	0.07021	0.43633	0.40523	Cu2	Cu	9.82329E-5	0.49998	0.4998
O4	O	0.12029	0.59696	0.36538	O1	O	0.58365	0.93338	0.41724
N1	N	0.15693	0.19125	0.16618	O2	O	0.61947	0.09186	0.36153
N2	N	0.08815	0.06873	0.08713	O3	O	0.92995	0.93652	0.09443
N3	N	0.14951	0.02079	0.14544	O4	O	0.87998	0.09714	0.13443
N4	N	0.40146	0.02803	0.40117	N1	N	0.84331	0.69137	0.33352
C1	C	0.36804	0.49656	0.13028	N2	N	0.91211	0.56875	0.41241
C2	C	0.28872	0.44284	0.17683	N3	N	0.8506	0.5209	0.3542
C3	C	0.26234	0.34065	0.15346	N4	N	0.59849	0.52843	0.09867
C4	C	0.18598	0.29407	0.19342	C1	C	0.632	0.99645	0.36933
C5	C	0.13652	0.34542	0.25762	C2	C	0.71135	0.94282	0.32275
C6	C	0.16457	0.44665	0.28266	C3	C	0.7378	0.84066	0.34613
C7	C	0.23927	0.49584	0.24099	C4	C	0.81428	0.79419	0.30628
C8	C	0.11495	0.5009	0.35594	C5	C	0.86379	0.8456	0.24214
C9	C	0.03275	0.24975	0.0483	C6	C	0.83566	0.94679	0.21707
C10	C	0.09138	0.17131	0.09847	C7	C	0.76085	0.99589	0.25865
C11	C	0.19234	0.09511	0.19389	C8	C	0.88528	1.00107	0.1438
C12	C	0.26453	0.07501	0.2653	C9	C	0.96773	0.74969	0.45127
C13	C	0.288	0.14721	0.3328	C10	C	0.90894	0.67134	0.40113
C14	C	0.35603	0.12021	0.39921	C11	C	0.80775	0.59529	0.30584
C15	C	0.38025	0.95836	0.33666	C12	C	0.73545	0.57531	0.23452
C16	C	0.3121	0.97816	0.26839	C13	C	0.71209	0.64749	0.16697
H1	H	0.30062	0.299	0.10397	C14	C	0.64403	0.62053	0.10058
H2	H	0.07731	0.30777	0.28854	C15	C	0.61957	0.45881	0.16324

C16 C 0.68772 0.47856 0.23152
 H1 H 0.69949 0.79892 0.39555
 H2 H 0.92308 0.80803 0.21127
 H3 H 0.74066 0.07552 0.24091
 H4 H 0.93461 0.82708 0.44581
 H5 H 0.03942 0.75201 0.42516
 H6 H 0.97538 0.72577 0.52072
 H7 H 0.74598 0.72346 0.16437
 H8 H 0.62498 0.67355 0.04647
 H9 H 0.57969 0.3859 0.15943
 H10 H 0.70385 0.42 0.28208
 H11 H 0.4389 0.69674 0.02061
 H12 H 0.46269 0.71662 0.98071
 H13 H 0.10498 0.69802 0.72734
 H4 H 0.10986 0.74663 0.75488
 H15 H 0.47451 0.80675 0.7553
 H16 H 0.50778 0.77614 0.78605
 H17 H 0.283 0.15099 0.98676
 H18 H 0.23167 0.13891 0.97385
 H19 H 0.19598 0.95107 0.45889
 H20 H 0.22369 0.92582 0.49678
 H21 H 0.73378 0.2014 0.12773
 H22 H 0.68527 0.22644 0.12466
 H23 H 0.44909 0.15298 0.85743
 H24 H 0.41264 0.11025 0.85402
 O1 O 0.58384 0.56644 0.91739
 O2 O 0.61952 0.408 0.86149
 O3 O 0.92985 0.56346 0.59434
 O4 O 0.88002 0.4028 0.63438
 N1 N 0.84338 0.80854 0.83345
 N2 N 0.91211 0.93118 0.91238
 N3 N 0.85064 0.97901 0.85412
 N4 N 0.59867 0.97146 0.59848
 C1 C 0.63215 0.50338 0.86942
 C2 C 0.71151 0.55701 0.82286
 C3 C 0.73796 0.65918 0.84622
 C4 C 0.81434 0.70571 0.80626
 C5 C 0.86377 0.65433 0.74205
 C6 C 0.83566 0.55313 0.71702
 C7 C 0.76093 0.50398 0.75869
 C8 C 0.88525 0.49887 0.64373
 C9 C 0.96771 0.75025 0.95129
 C10 C 0.90897 0.82859 0.9011
 C11 C 0.80784 0.90461 0.80573
 C12 C 0.73562 0.92461 0.73434
 C13 C 0.71225 0.85239 0.66682
 C14 C 0.64419 0.87933 0.60042
 C15 C 0.61981 0.04113 0.66299
 C16 C 0.68796 0.0214 0.73127
 H1 H 0.69972 0.70087 0.89572
 H2 H 0.92298 0.69194 0.7111
 H3 H 0.74073 0.42436 0.74093
 H4 H 0.93455 0.67289 0.94589
 H5 H 0.0394 0.74786 0.92518
 H6 H 0.97537 0.77424 0.02072
 H7 H 0.74609 0.7764 0.6643
 H8 H 0.62511 0.82629 0.54634
 H9 H 0.57996 0.11406 0.65916
 H10 H 0.70412 0.07998 0.7818
 H11 H 0.43936 0.80257 0.52111
 H12 H 0.46306 0.78196 0.48143
 H13 H 0.10511 0.80196 0.22743

H4 H 0.11 0.75335 0.25497
 H15 H 0.47506 0.69528 0.25437
 H16 H 0.50856 0.72635 0.28456
 H17 H 0.28628 0.35523 0.49065
 H18 H 0.23477 0.36484 0.47674
 H19 H 0.19615 0.54905 0.9588
 H20 H 0.22381 0.57431 0.99672
 H21 H 0.73309 0.29949 0.62721
 H22 H 0.68433 0.27503 0.62412
 H23 H 0.4461 0.34343 0.34993
 H24 H 0.41065 0.38717 0.34785
 O1 O 0.41664 0.06646 0.58238
 O2 O 0.38074 0.90809 0.63825
 O3 O 0.07001 0.06369 0.90497
 O4 O 0.12032 0.90308 0.86531
 N1 N 0.15709 0.30878 0.66609
 N2 N 0.08817 0.43133 0.58719
 N3 N 0.14962 0.47924 0.64543
 N4 N 0.40164 0.47189 0.90108
 C1 C 0.36827 0.0035 0.63037
 C2 C 0.28898 0.05725 0.67694
 C3 C 0.26261 0.15944 0.65355
 C4 C 0.18612 0.20596 0.69334
 C5 C 0.13652 0.15457 0.75743
 C6 C 0.16459 0.05337 0.78254
 C7 C 0.23941 0.00423 0.74101
 C8 C 0.11489 -8.68627E-4 0.85578
 C9 C 0.03274 0.25031 0.54833
 C10 C 0.09145 0.32874 0.59846
 C11 C 0.19254 0.40489 0.69378
 C12 C 0.2648 0.42493 0.76514
 C13 C 0.2882 0.35273 0.83266
 C14 C 0.35619 0.37972 0.8991
 C15 C 0.38053 0.54152 0.83653
 C16 C 0.31243 0.52175 0.76821
 H1 H 0.30098 0.20114 0.60414
 H2 H 0.07721 0.19218 0.78824
 H3 H 0.25956 0.92459 0.75877
 H4 H 0.06603 0.17301 0.55366
 H5 H 0.96111 0.24776 0.57458
 H6 H 0.02489 0.27432 0.47892
 H7 H 0.25438 0.27672 0.8352
 H8 H 0.37525 0.32669 0.9532
 H9 H 0.42034 0.61447 0.84038
 H10 H 0.29627 0.58032 0.71768
 H11 H 0.56147 0.3034 0.97919
 H12 H 0.53769 0.28351 0.01909
 H13 H 0.89554 0.30156 0.27265
 H4 H 0.89062 0.25299 0.24506
 H15 H 0.52685 0.19402 0.24449
 H16 H 0.4937 0.22476 0.21373
 H17 H 0.71754 0.84899 0.01258
 H18 H 0.76881 0.86118 0.02562
 H19 H 0.80459 0.04713 0.54125
 H20 H 0.77738 0.07284 0.50329
 H21 H 0.2667 0.79919 0.8723
 H22 H 0.31521 0.77415 0.87547
 H23 H 0.55216 0.84771 0.14202
 H24 H 0.58889 0.89016 0.14556
 H25 H 0.49824 0.48401 0.47865
 H26 H 0.50392 0.51853 0.51835

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