



Supporting Online Material for

Role of Solvent-Host Interactions That Lead to Very Large Swelling of Hybrid Frameworks

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Materials and Methods

Preparation of the samples. MIL-88B, MIL-88C and MIL-88D were obtained as reported before (SI). Prior to the collection of x-ray data, recrystallized carboxylic acid was first evacuated by extensive washing of the solids in dimethylformamide (MIL-88C) or pyridine (MIL-88B and D) at room temperature; each solid was then dried in air. In a second step, MIL-88B and MIL-88C samples were calcined overnight in air at 150°C to evacuate the solvent (pyridine, DMF). When returned to room temperature, a significant rehydration occurred for MIL-88B, with an approximate uptake of 15% water, whereas MIL-88C adsorbed water as only 3–4% of its weight. No treatment was applied to MIL-88D. These samples were then dispersed by stirring (i.e., 10 mg of solid for 1 ml of liquid) of MIL-88B, MIL-88C, and MIL-88D in various solvents such as water, methanol, ethanol, butanol, dimethylformamide, diethylformamide, dimethylsulfoxide, nitrobenzene, acetone, acetonitrile, pyridine, lutidine (2,6-dimethylpyridine), hexane, and toluene, for a few hours at room temperature. Glass capillaries were then filled with slurries and sealed to prevent evaporation of the solvent.

X-ray diffraction data. The *in situ* synchrotron powder diffraction experiments have been carried out at the Swiss-Norwegian Beamlines at the European Synchrotron Radiation Facility. The data were collected on the 1.0-mm quartz capillaries filled with the sample and a solvent, using MAR345 imaging plate at a sample-to-detector distance of 340 mm, $\lambda = 0.71118 \text{ \AA}$. The data were integrated using Fit2D program (Dr. A. Hammersley, ESRF) and a calibration measurement of a NIST LaB₆ standard sample.

Computer simulation. Computer simulations were used in order to yield structural models for the frameworks of the MIL-88A-B-C-D series of compounds, both in their contracted forms (narrow pore) and their solvated forms (open pores), and possibly initiate Rietveld refinement of the structures for selected compounds. For each MIL88-X solid (X = A, B, C, D), the starting structural models were obtained from the MIL-88A crystal structure, using our computational “ligand-replacement” strategy (SI). By substituting in the original MIL-88A topology the required ligand for the fumaric carboxylate, the framework models of MIL88-B (terephthalate), MIL-88C (2,6-naphthalene dicarboxylate), and MIL-88D (4,4'-biphenyl dicarboxylate) were generated that correspond to their observed as-synthesized cell parameters.

In this study, the as-synthesized forms of MIL-88X frameworks were used as initial models for modeling, MIL-88X-*as*. Starting from each of them, the dried (narrow pore) and solvated (open pores) forms were simulated using the cell parameters obtained by indexing the observed diffraction patterns. The simulations were performed on frameworks exclusively—i.e., in the absence of solvent molecules—in

order to yield starting framework models for further Rietveld refinement where solvent molecules could be located from difference Fourier maps.

The computational strategy is similar to the one we used for describing the swelling of the MIL-88A structure (S2). The main computational step consists of energy minimization for the hybrid frameworks in their original space group, *P*-62*c*, relaxing all atomic coordinates and cell parameters in a sequential fashion, *enforcing* either a contraction or an expansion of the cell, using the observed cell parameters as the targeted values. The key feature here is a use of carefully chosen external constraints along the cell axes—i.e., hydrostatic pressure—so as to enforce the elongation or the shrinkage of the crystal structure along specific cell axes during the energy minimizations. The hydrostatic pressure is therefore applied anisotropically, for example increasing it along *a* while keeping fixed along *c*, and vice versa. This pressure is chosen as positive when a shrinkage of the cell parameter is required or negative in the case of an elongation. This way, the initial models for all MIL-88X-*as* had their energy minimized sequentially, where the anisotropic hydrostatic pressure was carefully adjusted at each minimization step until the simulated cell parameters finally reached the targeted experimental values. This is the key point of these simulations, where the flexible hybrid framework of MIL-88 is enforced to adopt the targeted experimental cell parameters while keeping its network's connectivity identical. The above simulations were performed using the 'minimizer' of Cerius2 suite of software (S3) and the universal force field (S4) with periodic bound conditions. The force field is used as a simple mean to ensure that the so-minimized hybrid frameworks are realistic, i.e., kept within an acceptable range of structural features (bond distances, angles, and so on). Electrostatic interactions were not computed in this work.

Typically, in order to simulate the expected extended cell of MIL-88D-open ($a_{\text{final}} = 20.04 \text{ \AA}$), a negative hydrostatic pressure was first applied along the *a* axis when minimizing the initial model framework ($a_{\text{initial}} = 12.05 \text{ \AA}$), while keeping the *c* cell parameter fixed. A second step was needed for applying the adequate positive hydrostatic pressure along the *c* axis ($c_{\text{initial}} = 27.50 \text{ \AA}$), while keeping the *a* parameter fixed, in order to converge towards the smaller expected value ($c = 22.91 \text{ \AA}$). A similar approach was successively applied in order to simulate the crystal structures of all the extended and contracted cells of MIL-88X. All models converged relatively easily towards the targeted experimental cell parameters while maintaining the framework topology within realistic structural limits. Finally, the powder diffraction patterns of the different types of MIL-88X were simulated, and their consistency was checked against the experimental patterns.

Structure refinement. Structure refinements were done using the atomic coordinates from the computer simulation and the program Fullprof2k and its graphical interface Winplotr (S5, S6). Successive Fourier differences were performed using Shelxtl to locate the approximate position of the guest species (S7). Please note that we propose here only an approached structure of the latter solids filled with solvent. It is obvious that when using powder data with a low signal-to-background ratio, the position of the free solvent molecules, sometimes disordered at room temperature, is approximate, particularly in the case of the largest pore materials MIL-88C and MIL-88D.

Table S1 : Crystal data and structure refinement parameters for **MIL-88B_methanol** or $Cr_6O_2F_2[1,4\text{ BDC}]_6.48CH_3OH$, **MIL-88C_pyridine** or $Fe_6O_2X_2[2,6\text{ NDC}]_6.30C_5H_5N.48H_2O$ and **MIL-88D_lutidine** or $Cr_6O_2F_2[4,4'\text{ BPDC}]_6.18C_7H_9N.30H_2O$

| Formula | MIL88B_Methanol | MIL-88C_Pyridine | MIL-88D_Lutidine |
|---|------------------------------|----------------------------|-------------------------------------|
| Chemical formula | $Cr_6F_2O_{66}C_{96}H_{216}$ | $Fe_6O_{80}C_{246}H_{264}$ | $Cr_6F_2O_{64}N_{18}C_{258}H_{270}$ |
| Solvent | Methanol | Pyridine | 2,6-Lutidine |
| Molar weight ($g \cdot mol^{-1}$) | 2774 | 5250 | 4992 |
| Calculated density ($g \cdot cm^{-3}$) | 1.36 | 1.53 | 1.04 |
| Crystal system | Hexagonal | Hexagonal | Hexagonal |
| Space group | P $6_3/mmc$ (n°194) | P $-6\ 2\ c$ (n°190) | P $-6\ 2\ c$ (n°190) |
| a (Å) | 15.626 (1) | 18.753(1) | 20.040(1) |
| c (Å) | 15.960(1) | 18.743(2) | 22.911(2) |
| V (Å ³) | 3375.0(3) | 5708.2(4) | 7968.7(2) |
| Z | 1 | 1 | 1 |
| Figures of merit | $M_{15}/F_{15}=20/50$ | $M_{14}/F_{14}=25/70$ | $M_{20}/F_{20}=16/50$ |
| Radiation λ (Å) | 0.71118 | 0.71118 | 0.71118 |
| Temperature (K) | 296 | 296 | 296 |
| 2θ range (°) | 2-26.7 | 2-26.7 | 2-26.7 |
| N. reflections | 202 | 347 | 450 |
| N. independent atoms | 18 | 36 | 48 |
| N. structural parameters | 33 | 89 | 120 |
| N. profile parameters | 8 | 8 | 8 |
| N. soft distance constraints | 36 | 92 | 128 |
| R_p | 0.5 | 3.1 | 1.4 |
| R_{Bragg} | 21.7 | 19.3 | 23.7 |
| Isotropic overall atomic displacement parameter | 4.3(2) | 5.1(2) | 4.5(2) |
| Profile function | Pearson VII | Pearson VII | Pearson VII |
| Background | Experimental | Experimental | Experimental |
| N. of asymmetry parameters | 2 | 2 | 2 |
| Secondary phase | ----- | Fe_2O_3 | ----- |

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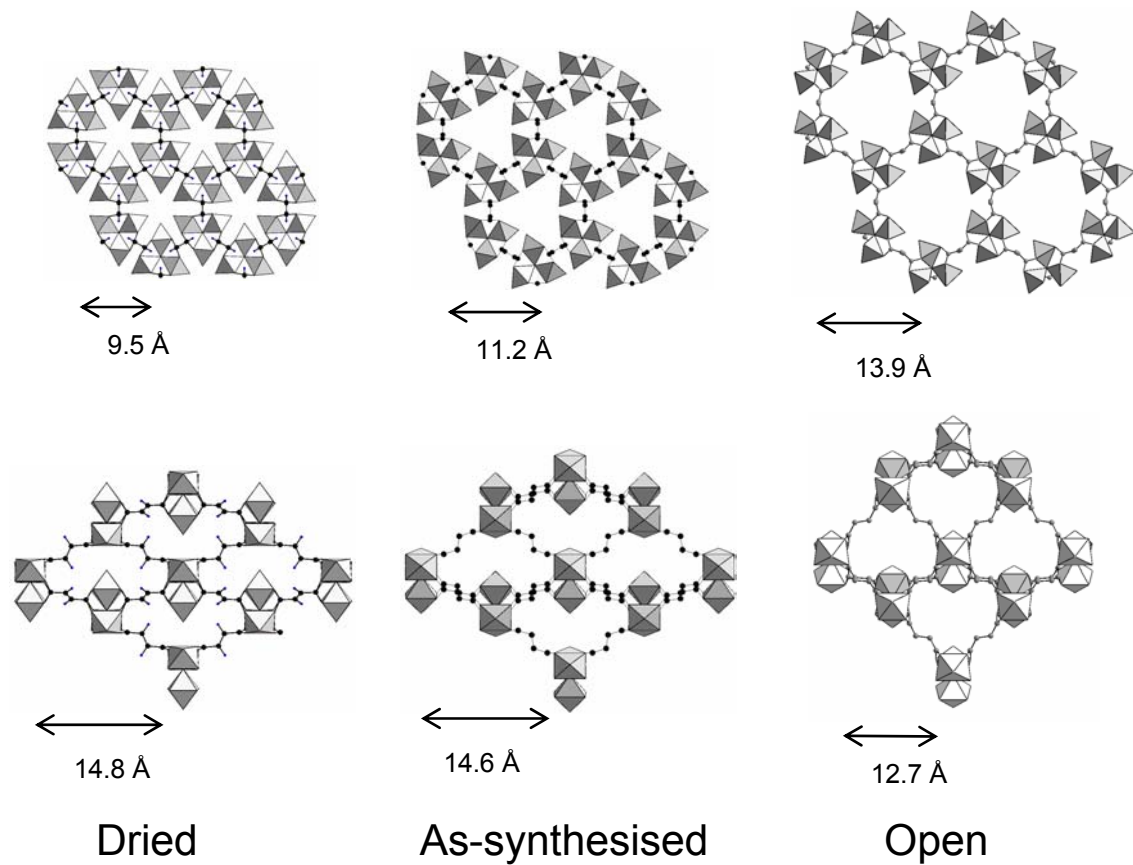


Fig. S1: view of the structure of the MIL-88A in its dry, as and open forms; top: along the c axis; bottom: along the a axis

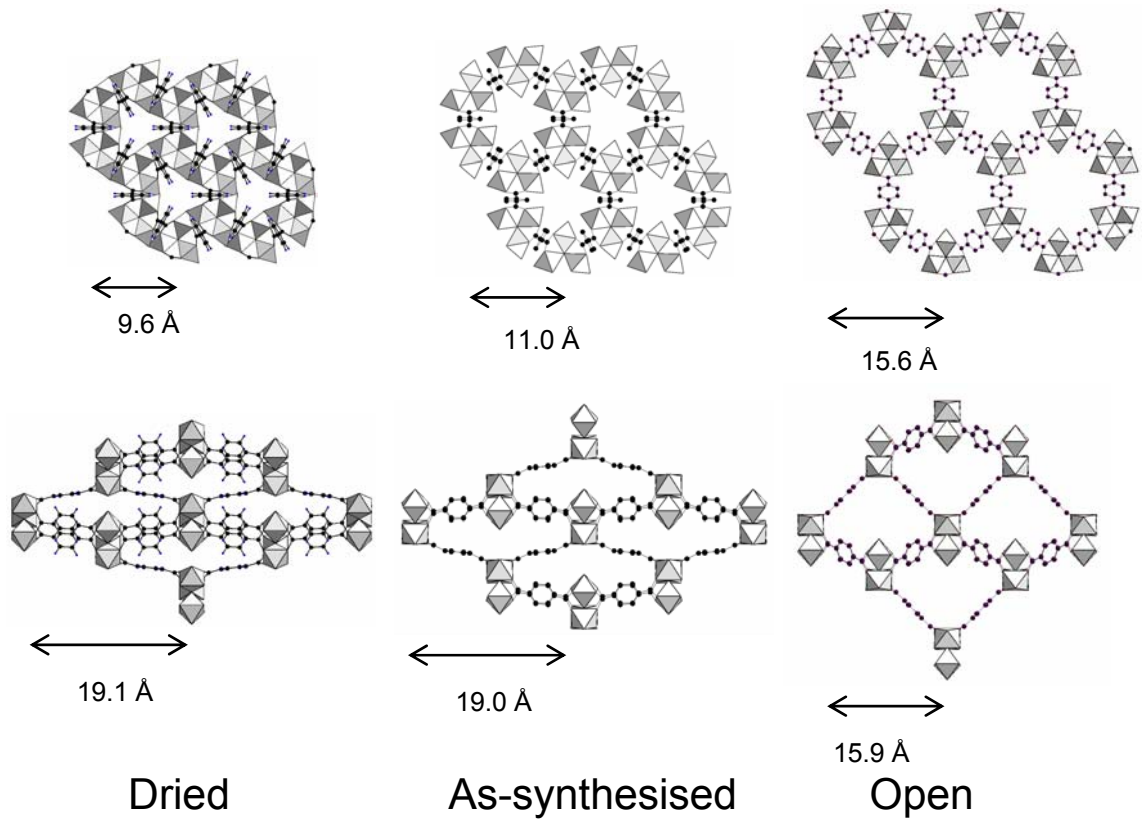


Fig. S2: view of the structure of the MIL-88B in its dry, as and open forms; top: along the c axis; bottom: along the a axis

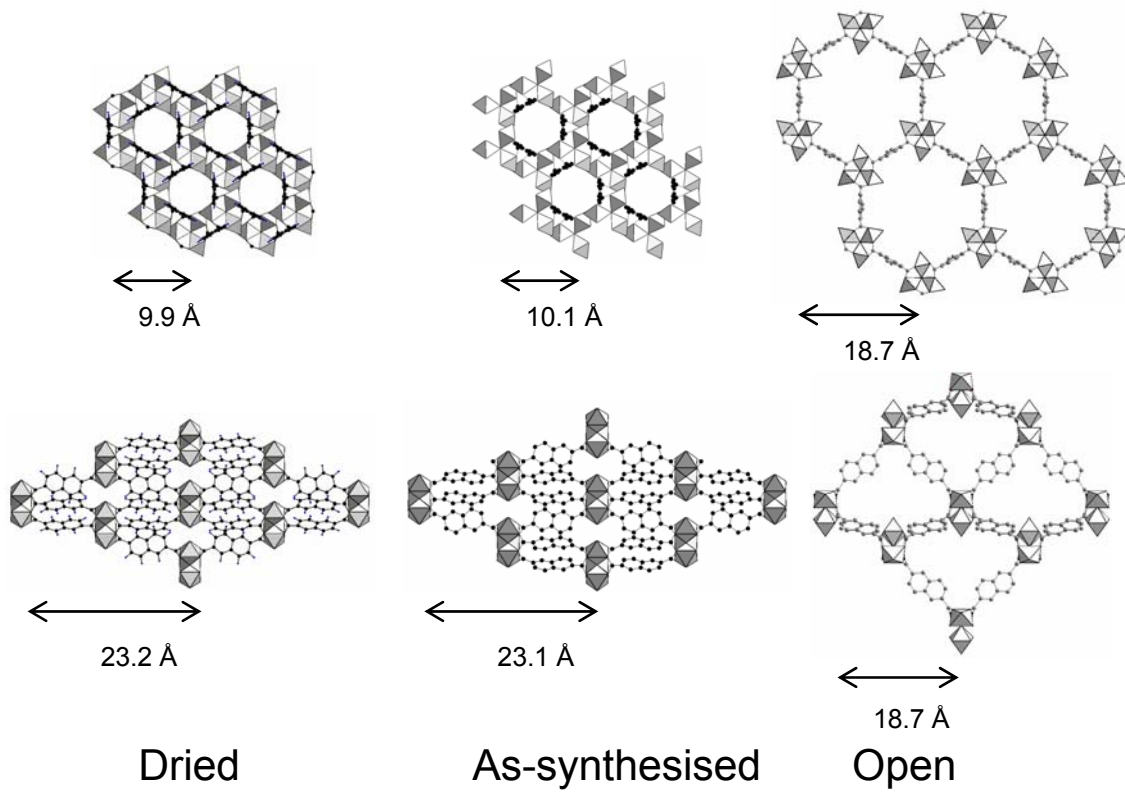


Fig. S3: view of the structure of the MIL-88C in its dry, as and open forms; top: along the c axis; bottom: along the a axis

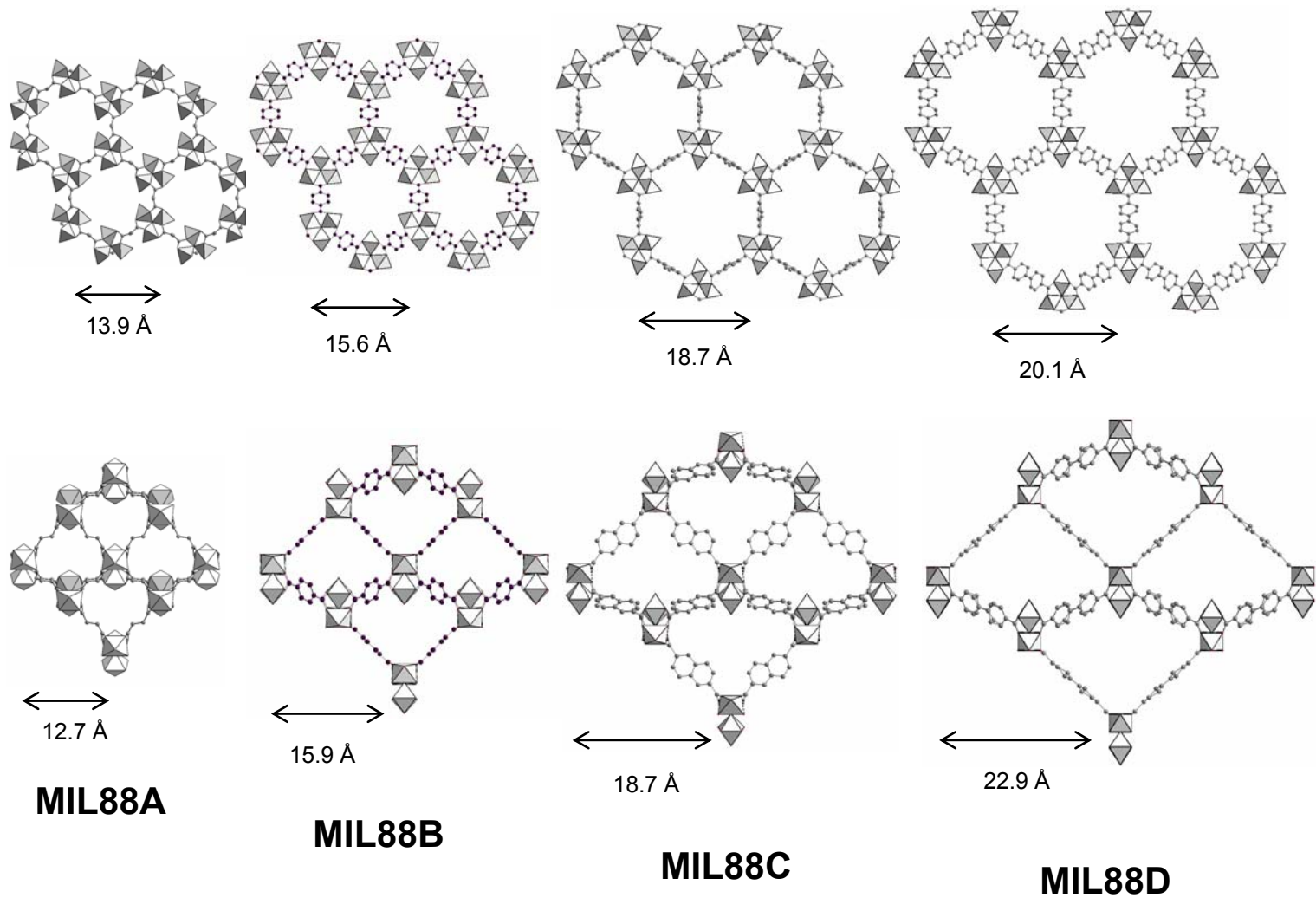


Fig. S4: view of the structure of the open forms of MIL-88A, B, C, D; top: along the c axis; bottom: along the a axis

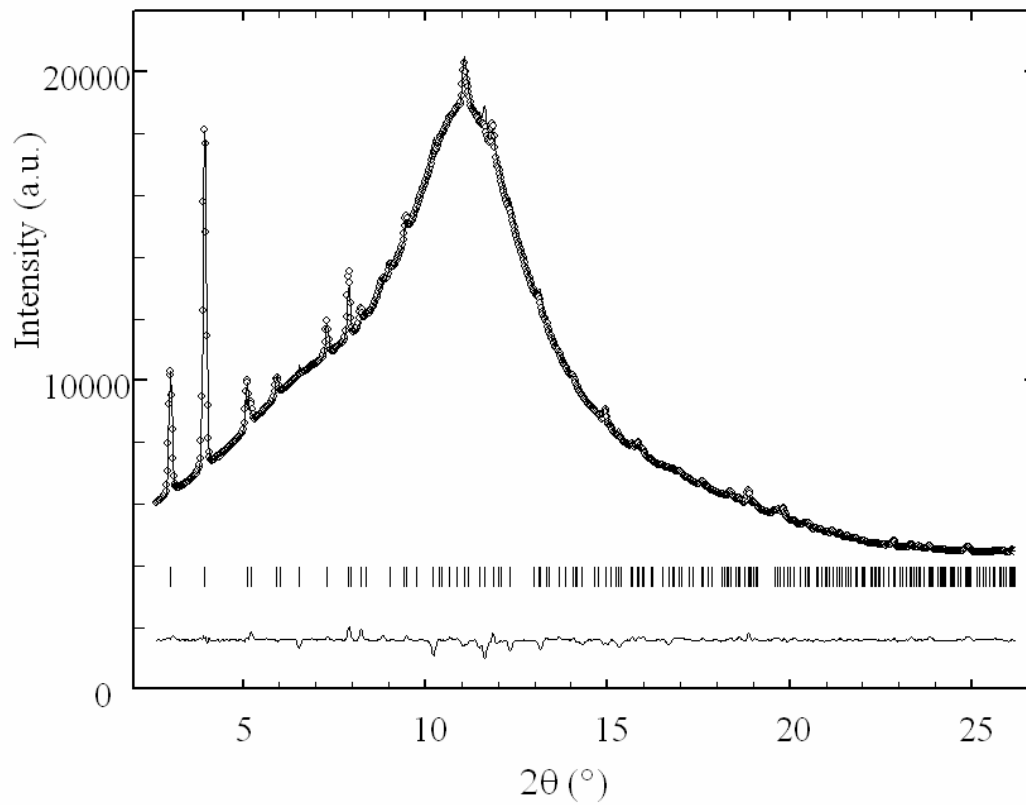


Fig. S5: Rietveld plot of MIL88B_methanol
(NB: the large hump at 2-theta=12° is due to the adsorption of methanol)

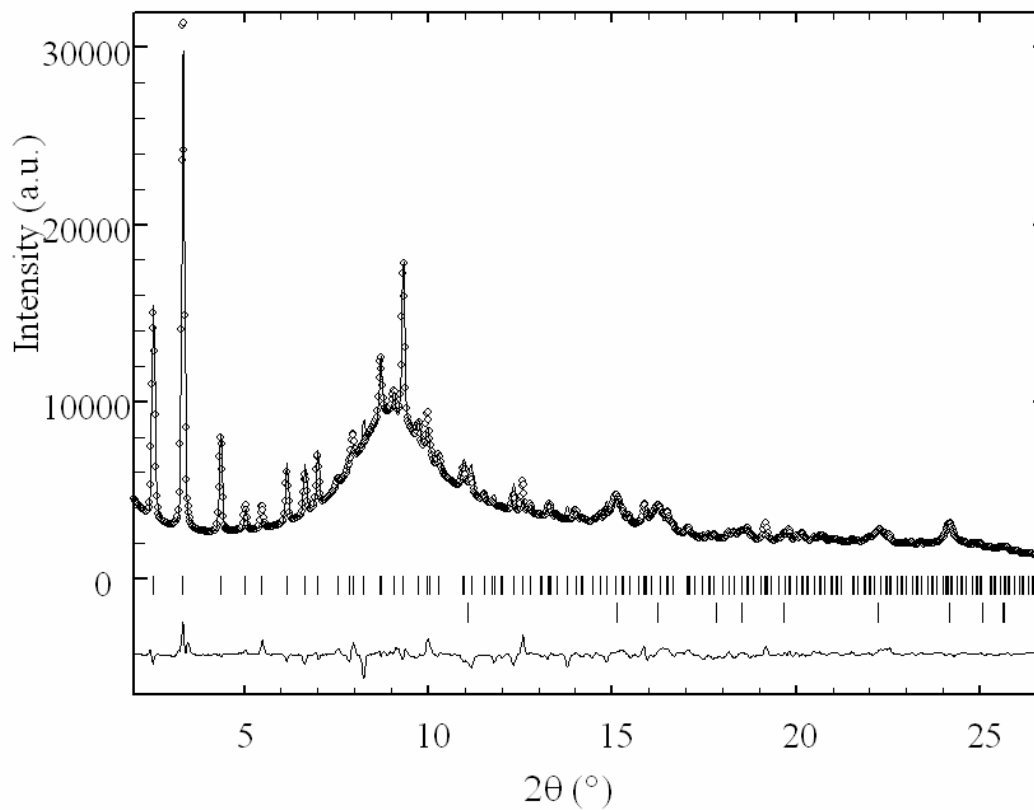


Fig. S6: Rietveld plot of MIL88C_pyridine
(NB: the large hump at 2-theta=9° is due to the adsorption of pyridine)

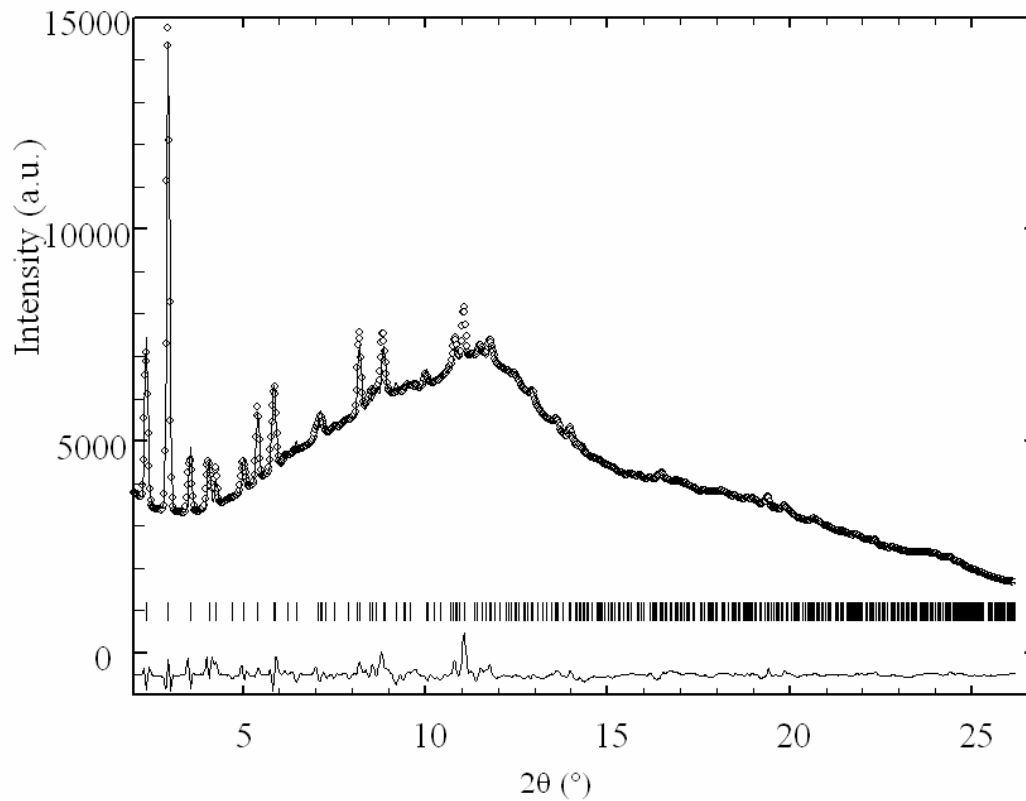


Fig. S7: Rietveld plot of MIL88D_lutidine
(NB: the large hump at 2-theta=12° is due to the adsorption of lutidine)

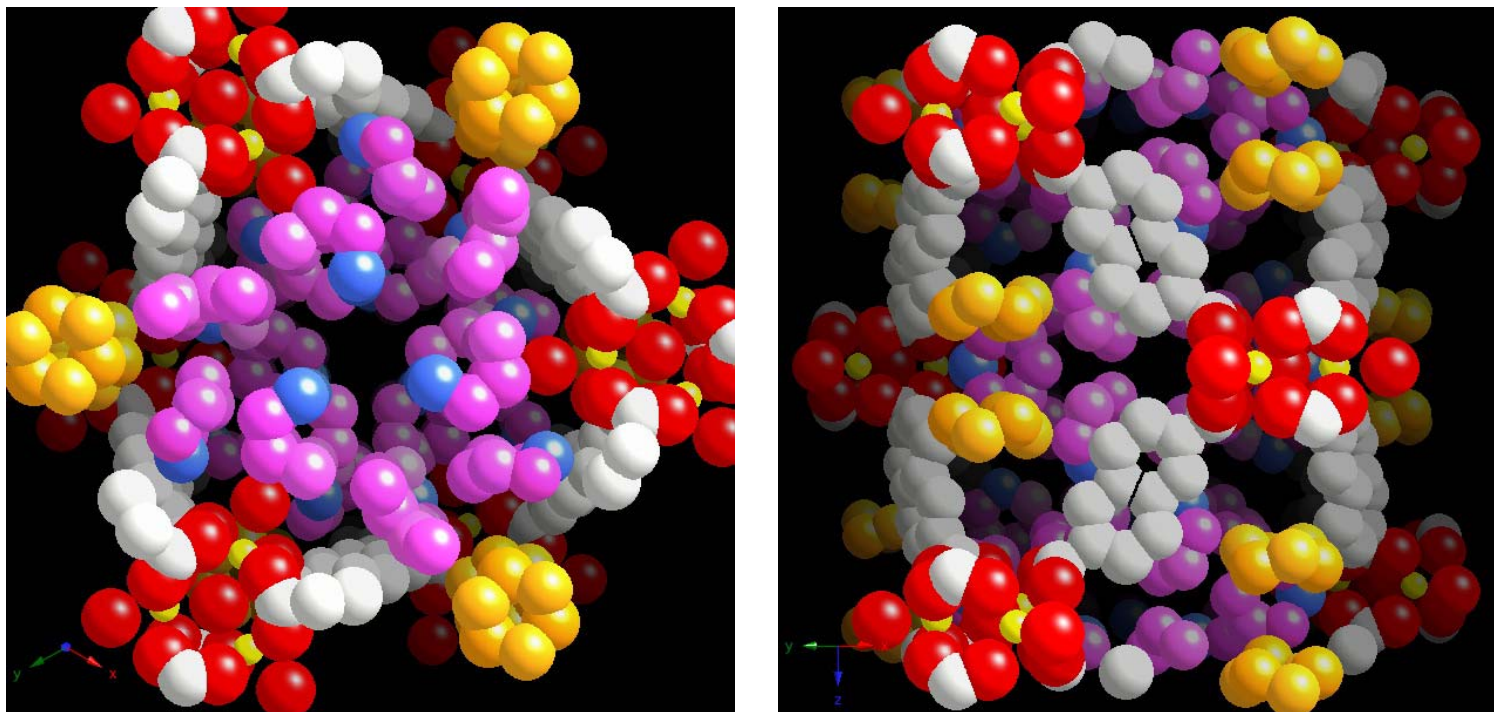


Fig. S8: Left : view of the tunnels of MIL-88C filled with pyridine; right: along the a axis Inorganic section (yellow and red), spacer (white) and pyridine (blue, purple).

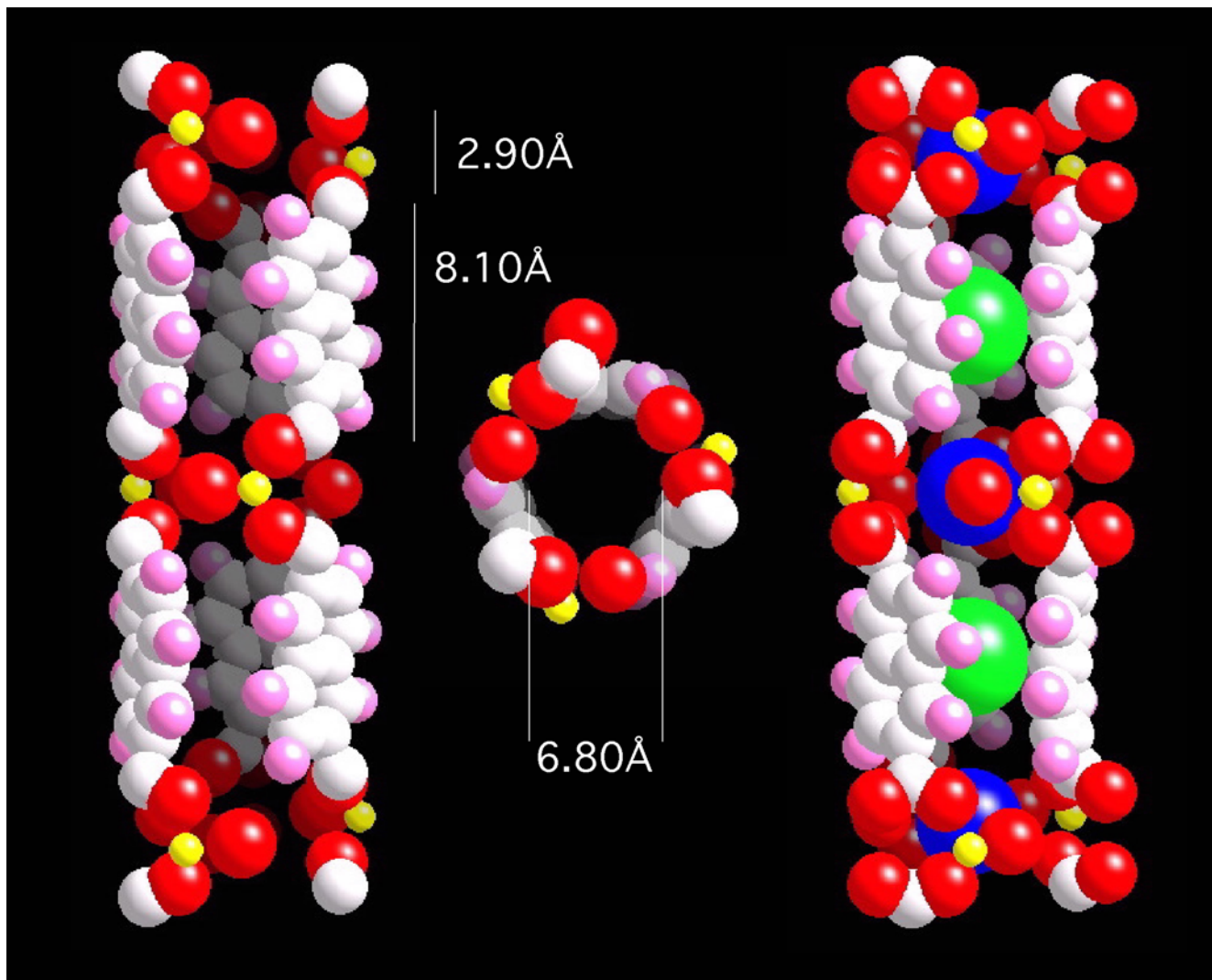
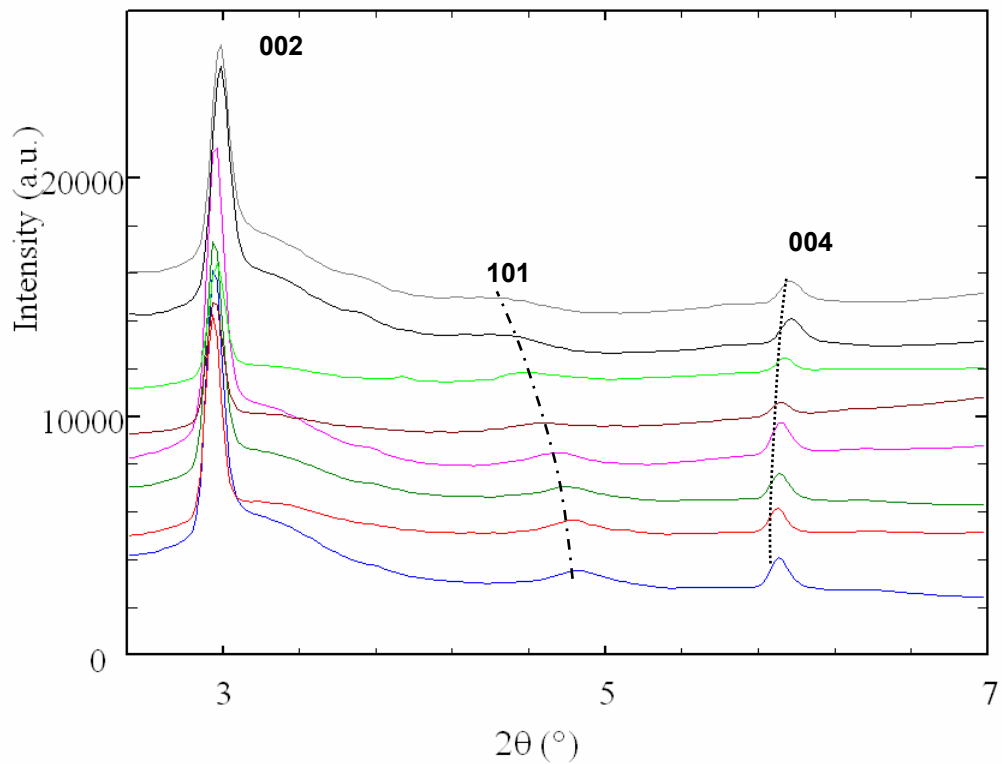


Fig. S9: Perspective view of the tunnels of MIL-88C; center: section of the tunnel; right: the tunnel with dummy atoms corresponding to the free aperture of the Inorganic section (blue) and organic one (green)



SG P-62c (n°190)

| Solvent | V(Å ³) | ΔV/Vdried |
|----------------------------|--------------------|-----------|
| Pyridine | 2990 | 20 % |
| DMF | 2965 | 19 % |
| MeOH, BuOH, DMSO | 2800 | 12-13 % |
| Hexane, acetone, DMC, EtOH | 2685 | 7-8 % |
| Lutidine, DEF | 2585 | 3-4 % |
| H ₂ O | 2535 | 2 % |
| Toluene | 2500 | <1 % |
| Dried | 2490 | --- |

Fig. S10: adsorption of liquids in MIL-88Ddry

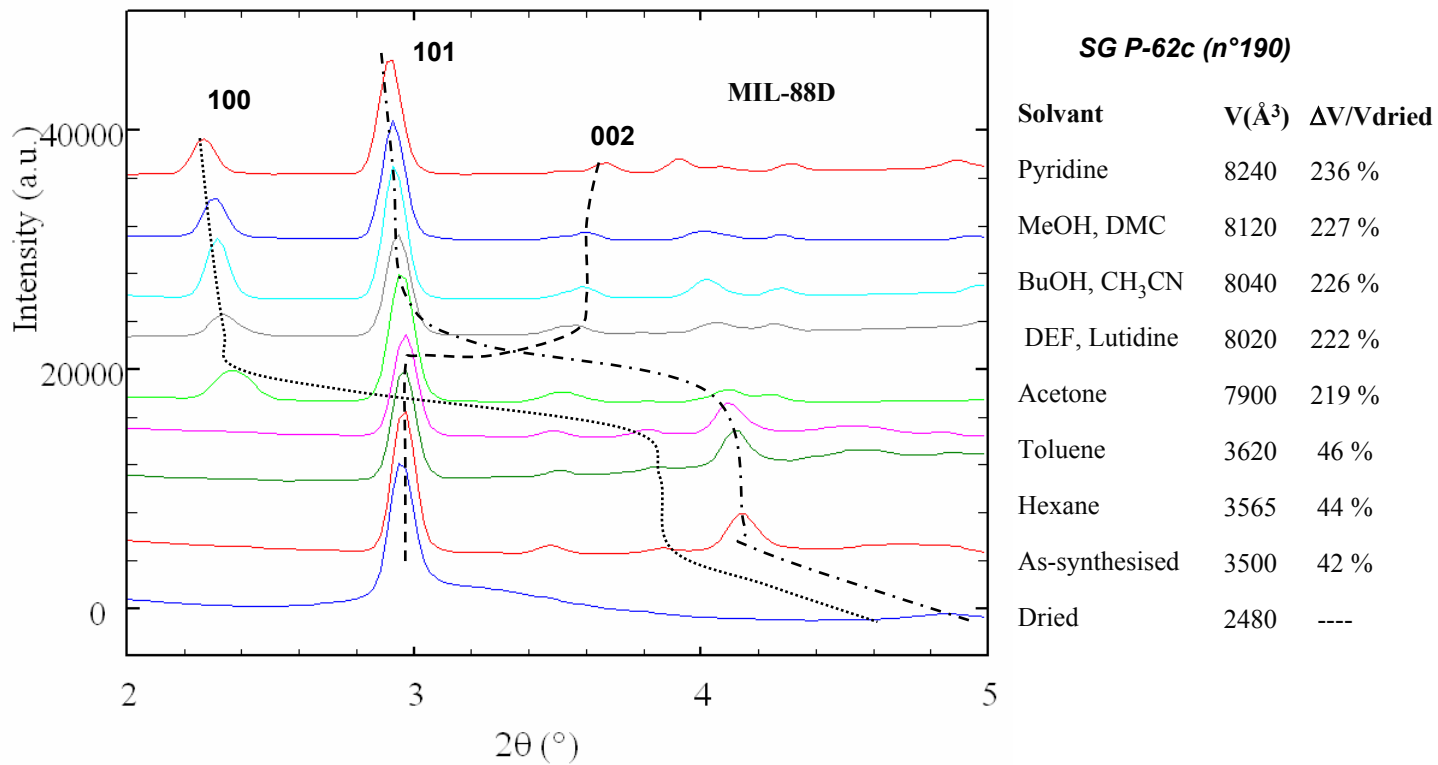


Fig. S11: adsorption of liquids in MIL-88Das

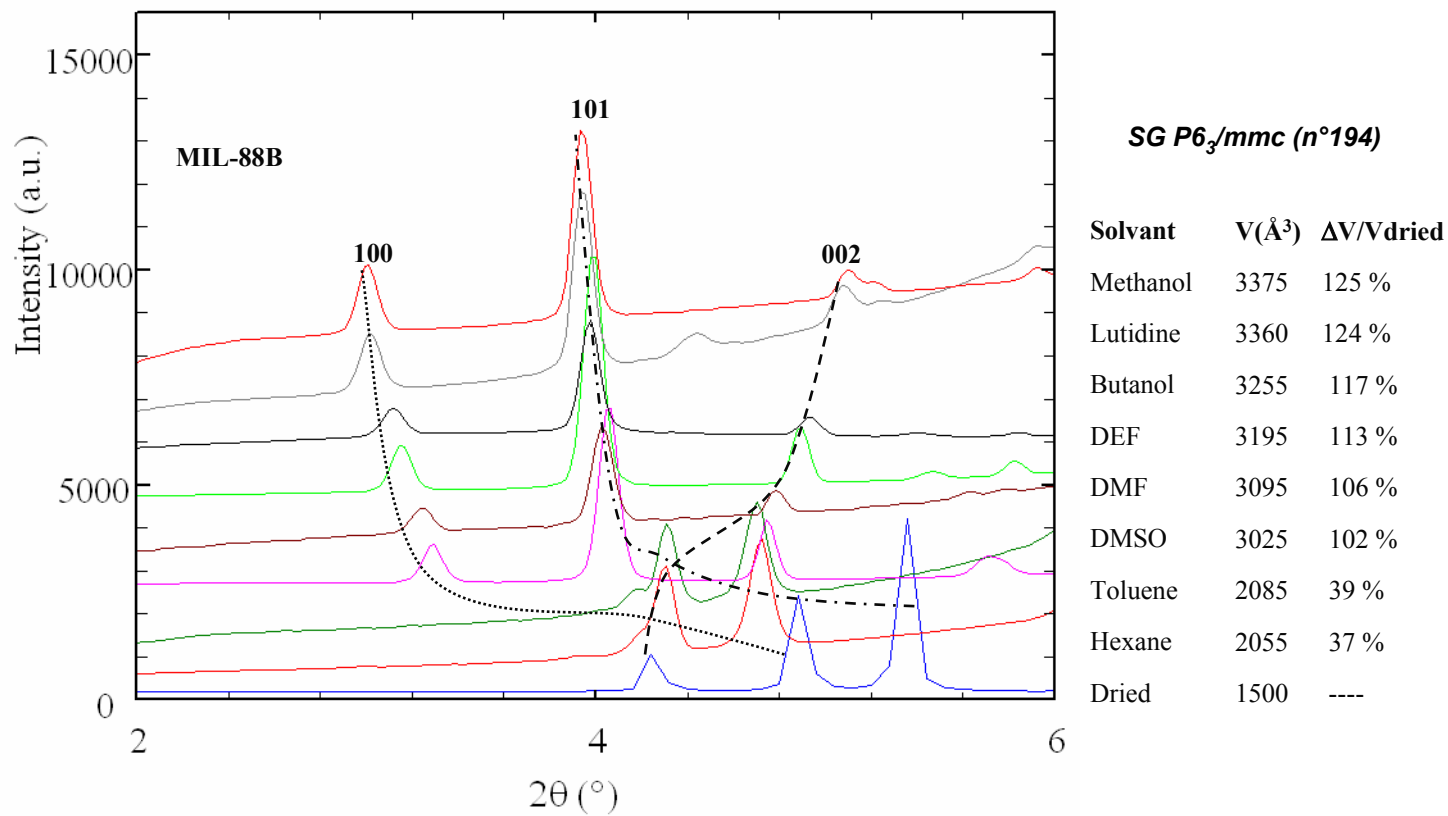


Fig. S12: adsorption of liquids in MIL-88Bas

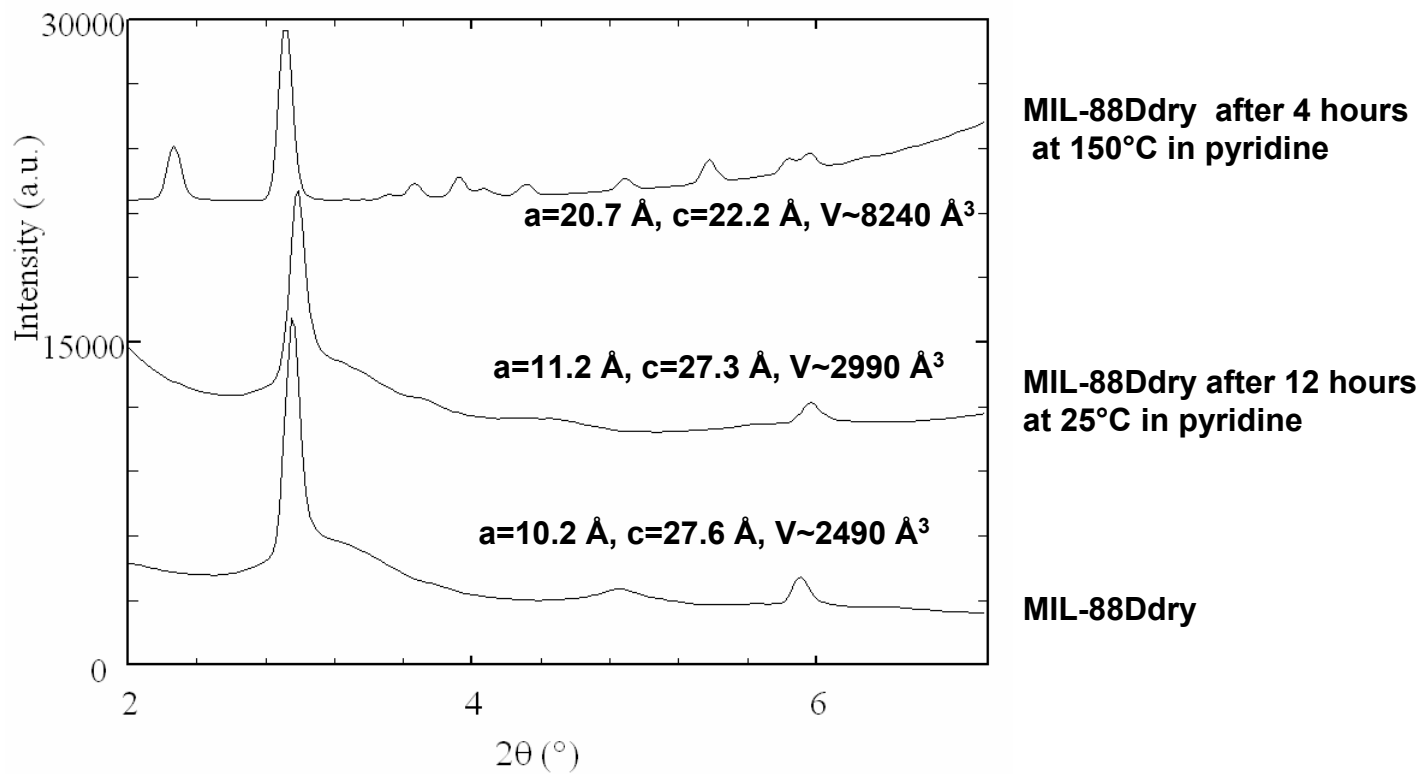


Fig. S13: effect on temperature on pyridine adsorption in MIL-88D

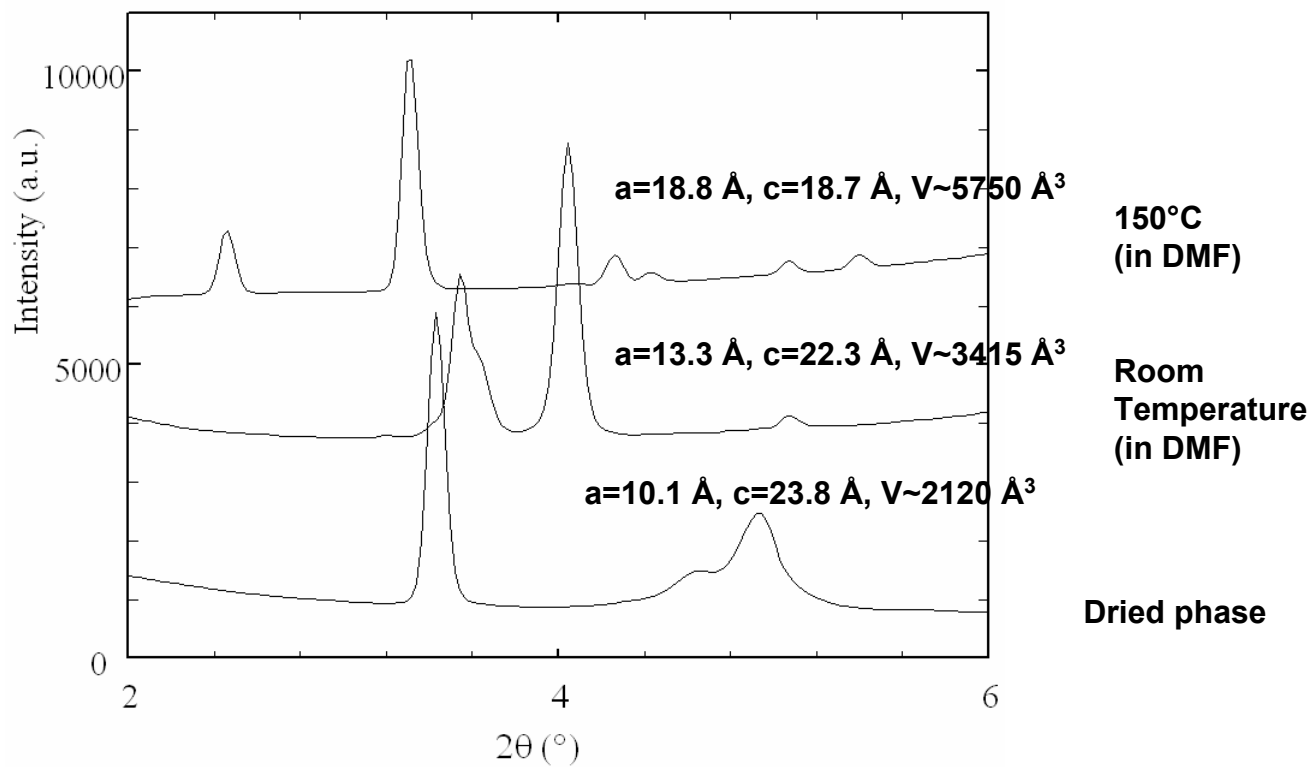


Fig. S14: effect on temperature on DMF adsorption in MIL-88C

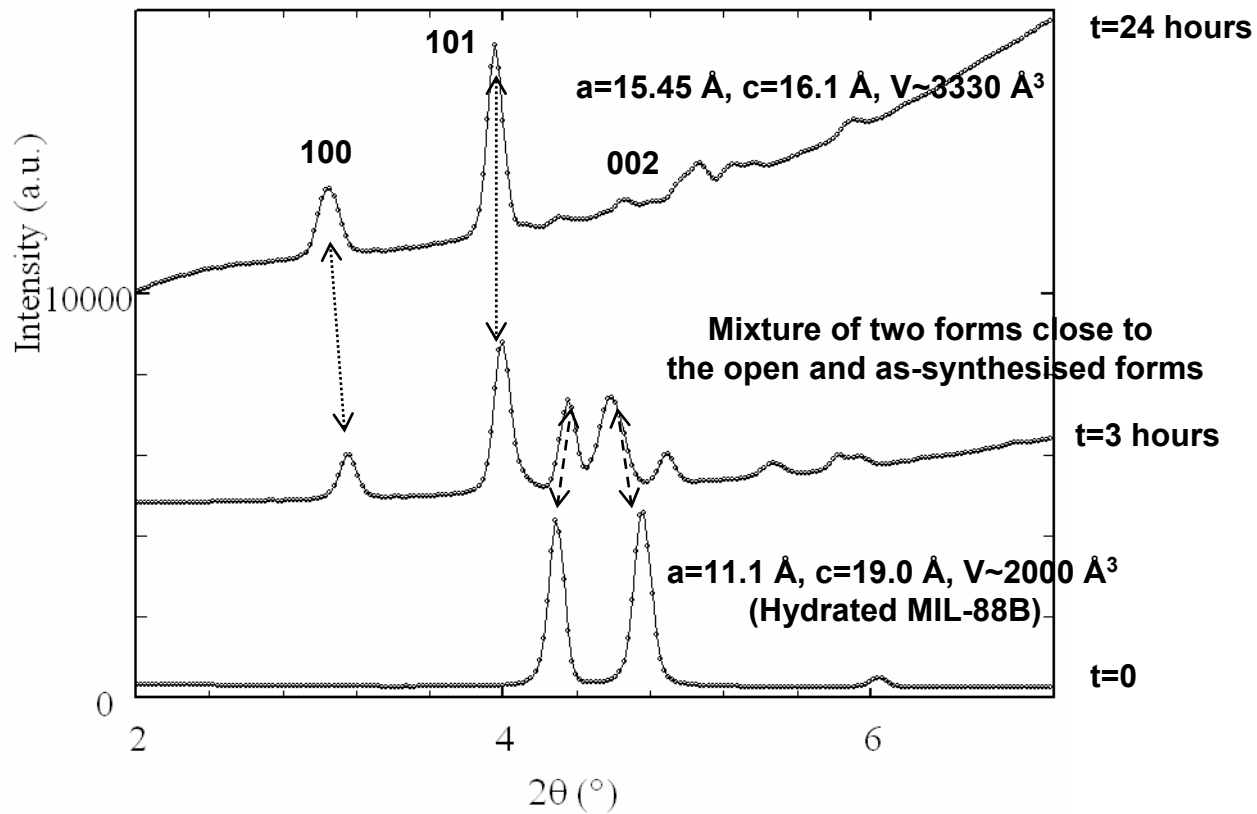


Fig.S15: X-Ray powder pattern of MIL-88B dispersed in nitrobenzene, evidencing the slow kinetics of nitrobenzene adsorption

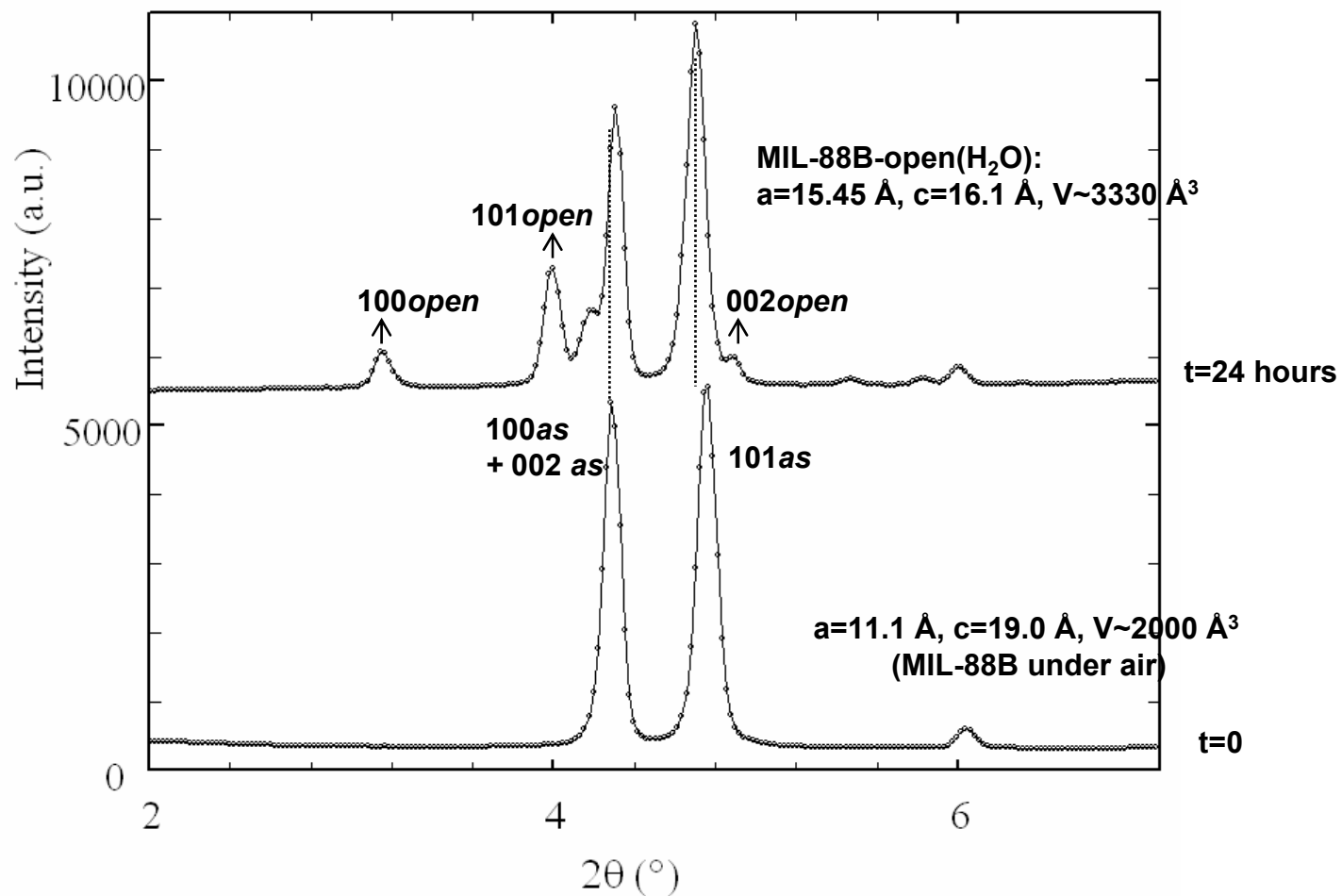


Fig.S16: X-Ray powder pattern of MIL-88B dispersed in water, evidencing the very slow kinetics of water adsorption

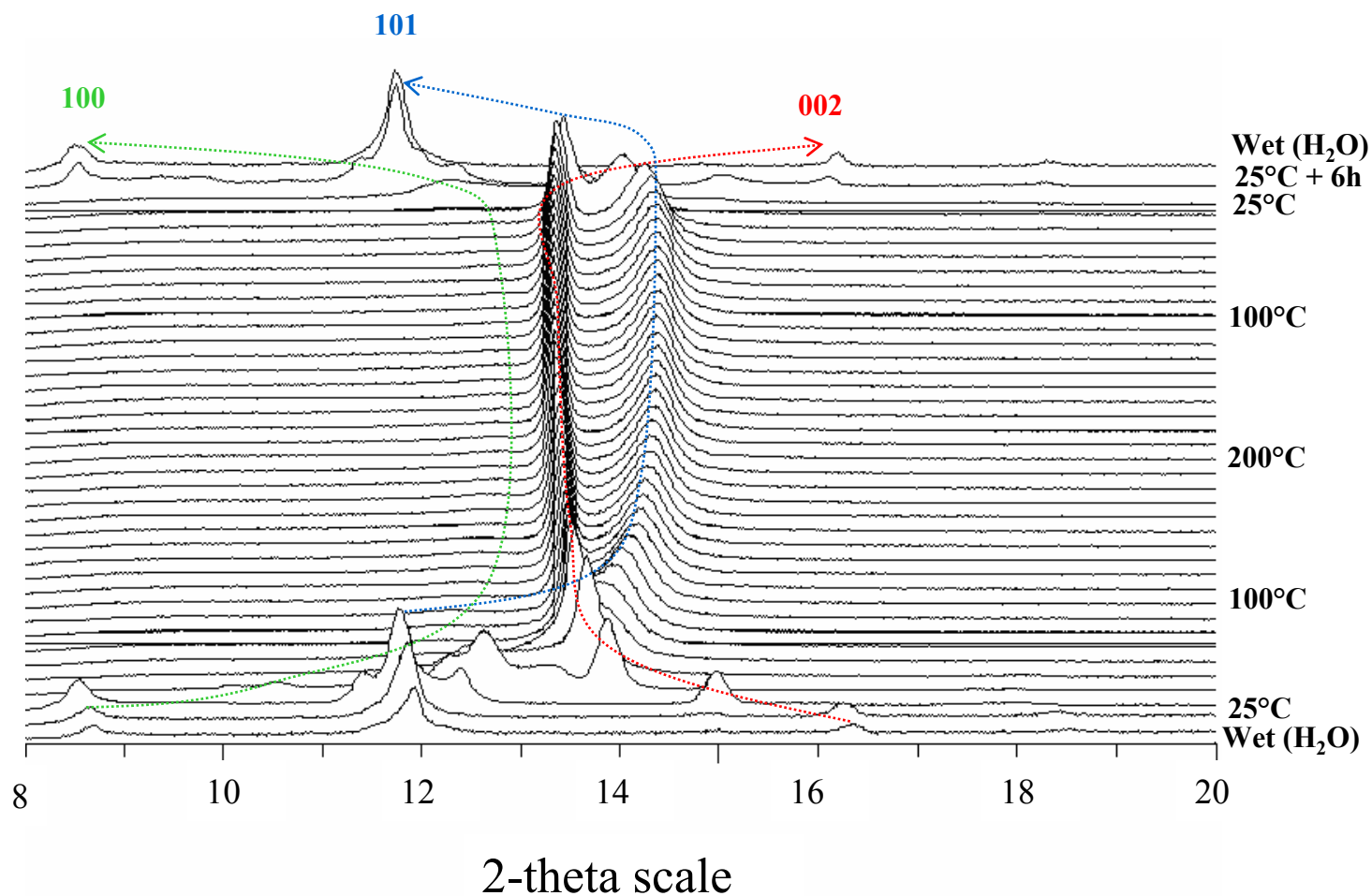


Fig.S17: X-Ray thermodiffractometry of MIL-88A under air ($\lambda \sim 1.79 \text{ \AA}$)

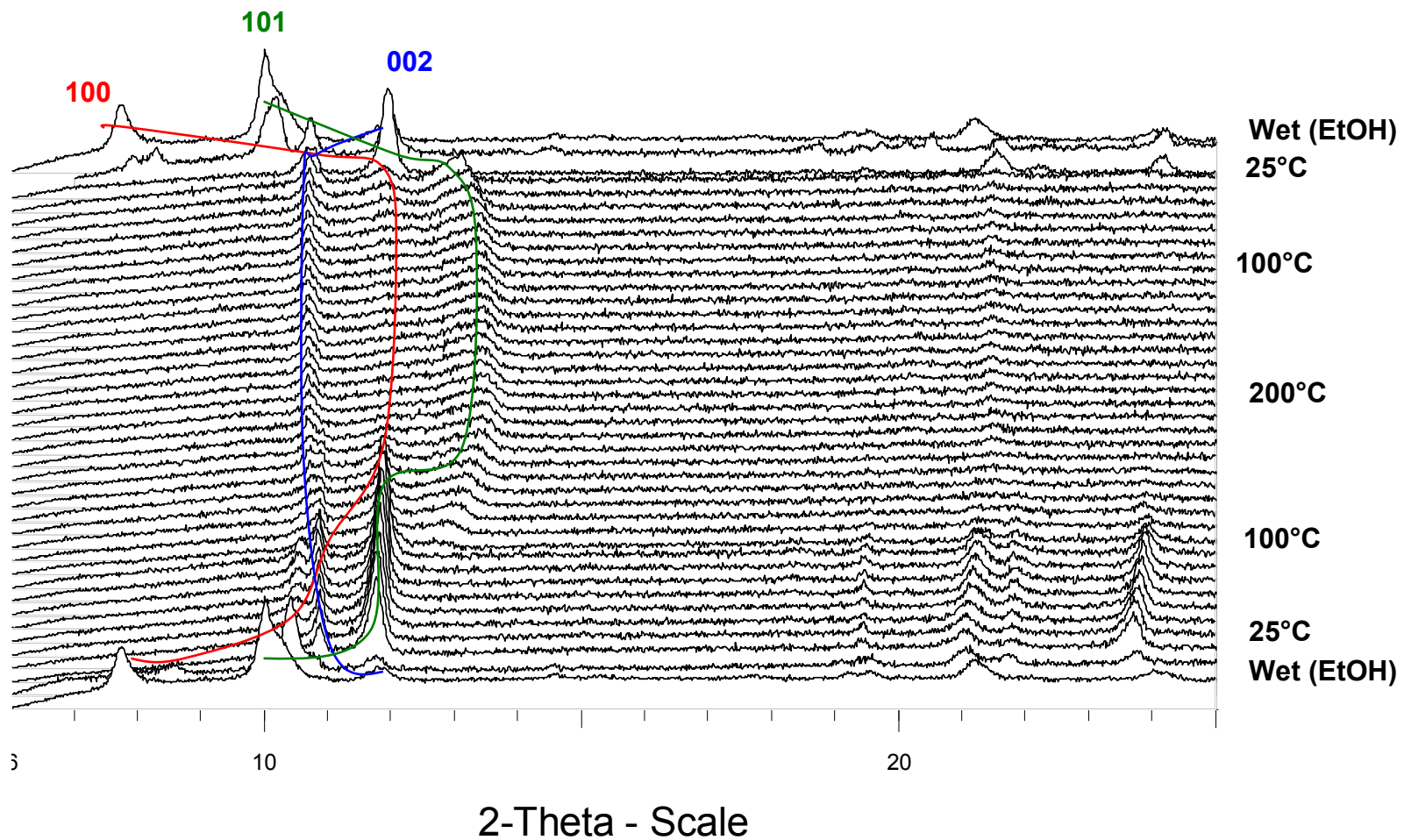
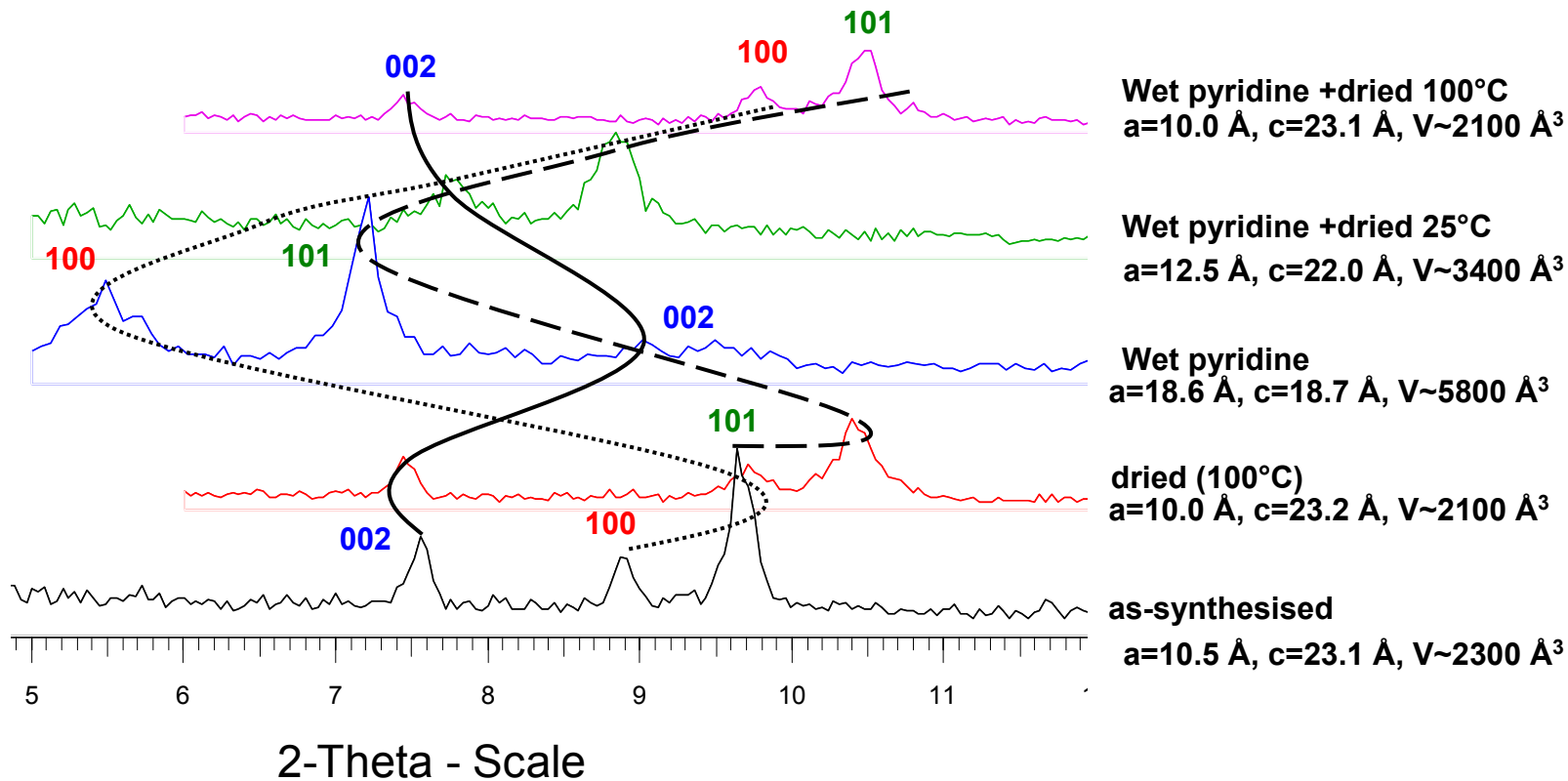
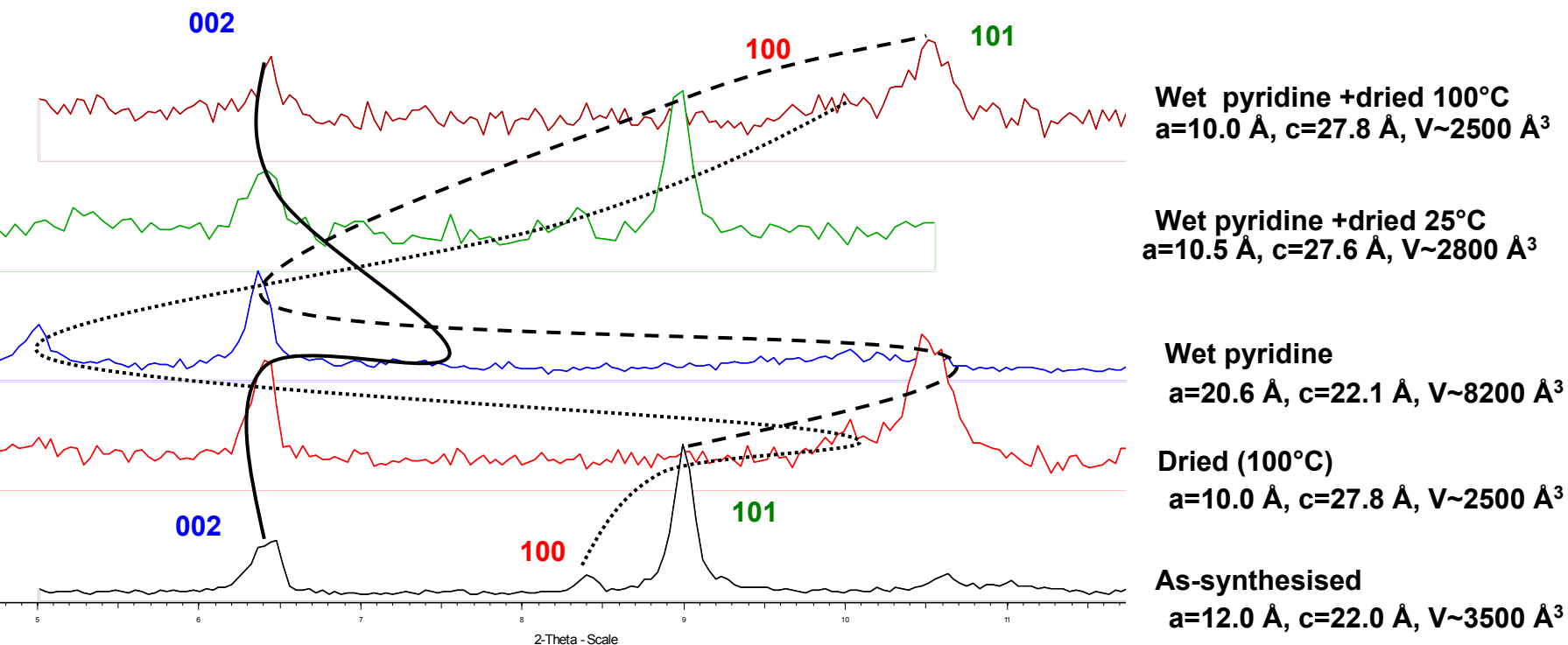


Fig.S18: X-Ray thermodiffraction of MIL-88B under air ($\lambda \sim 1.79 \text{ \AA}$)



Exp. conditions : a few mg of MIL-88C has been dispersed on a glass sample holder
 « Wet pyridine » means that one drop of pyridine has been dispersed on the solid

Fig.S19: X-Ray diffraction patterns of MIL-88C (powder)
 under various conditions ($\lambda\sim 1.54 \text{ \AA}$)



Exp. conditions : a few mg of MIL-88D has been dispersed on a glass sample holder
 « Wet pyridine » means that one drop of pyridine has been dispersed on the solid

Fig.S20: X-Ray diffraction patterns of MIL-88D (powder)
 under various conditions ($\lambda\sim 1.54 \text{ \AA}$)