

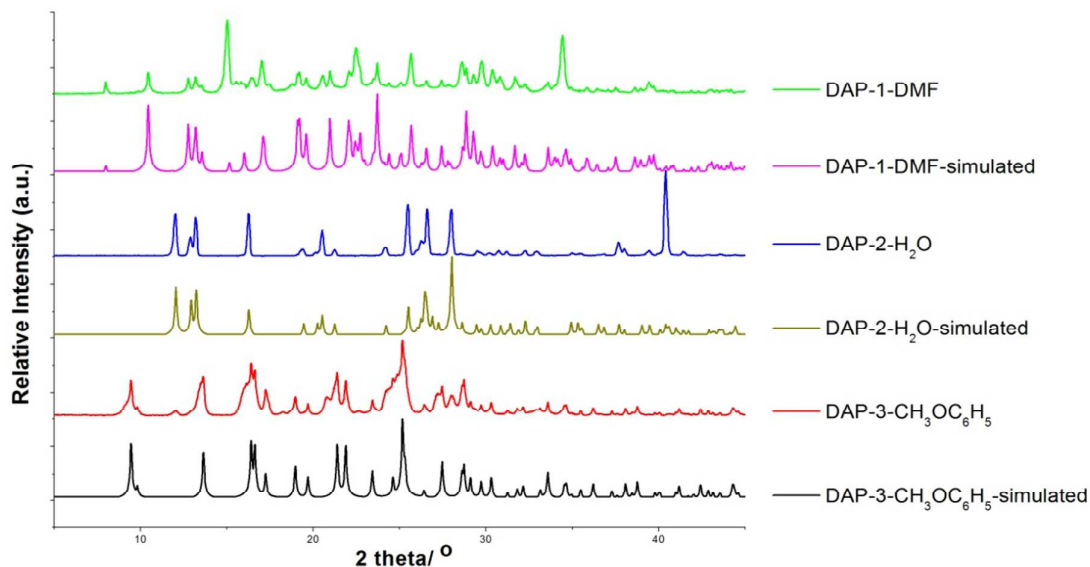
## Solvent Dependent Structures of Hydrogen-Bonded Organic Frameworks of 2,6-Diamino Purine

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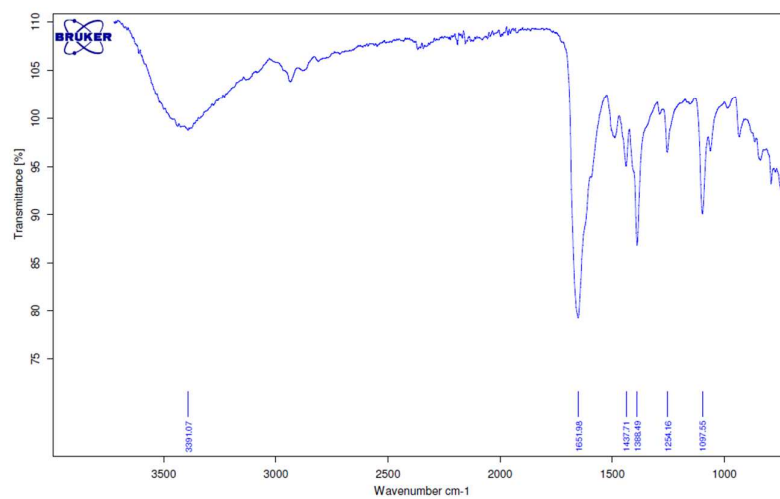
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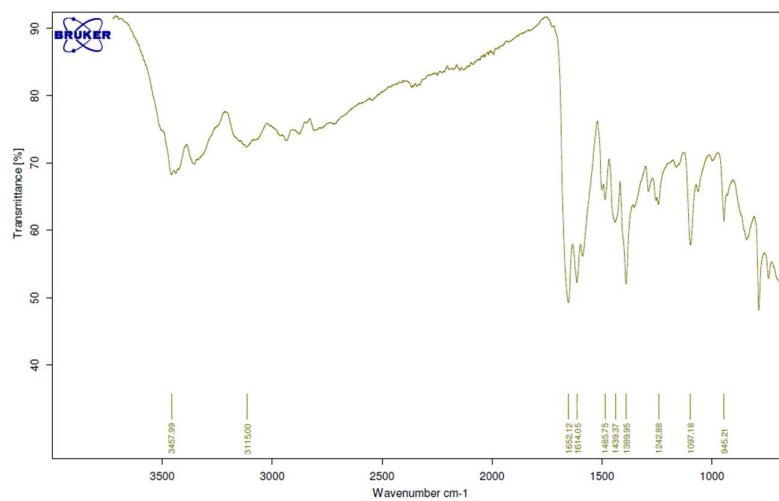
### 1. Characterizations



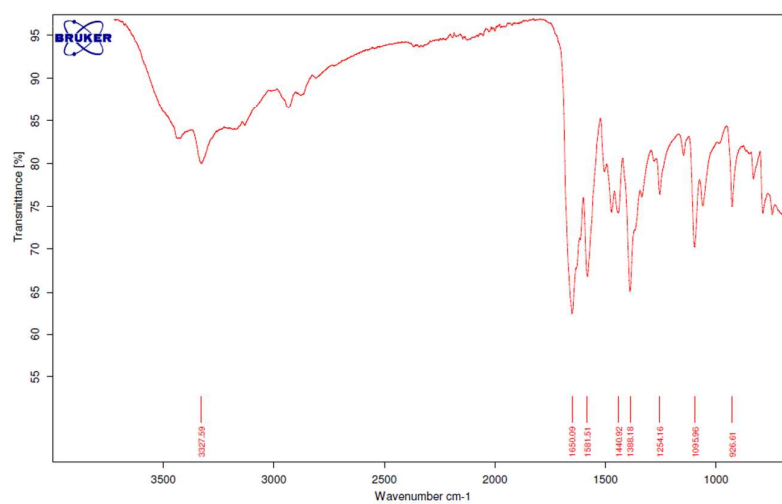
**Figure S1.** Powder X-ray diffraction patterns of simulated **DAP-1-DMF**, **DAP-2-H<sub>2</sub>O** and **DAP-3-CH<sub>3</sub>OC<sub>6</sub>H<sub>5</sub>** and as-synthesized ones.



(a)

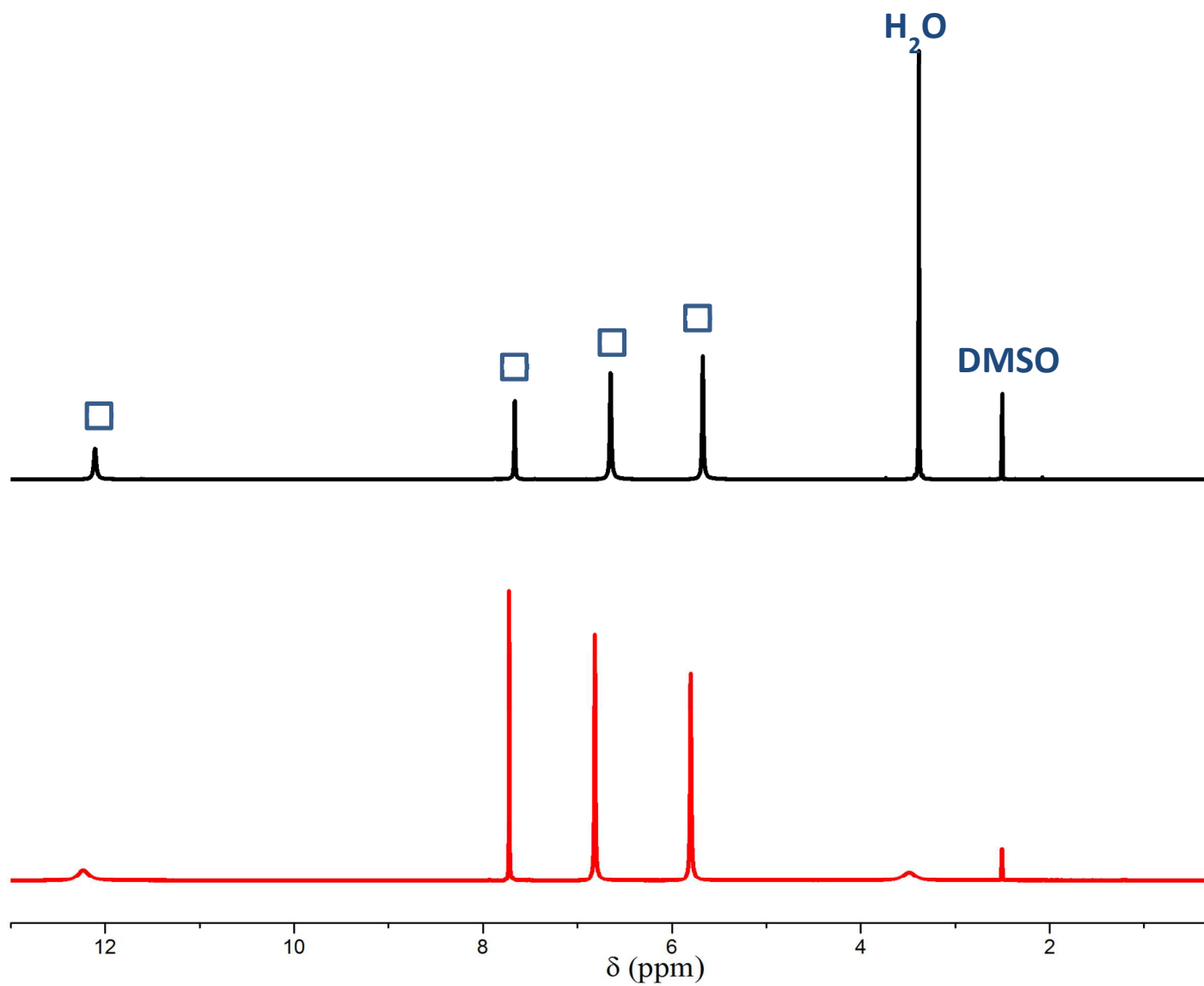


(b)



(c)

**Figure S2.** FT-IR patterns of **DAP-1-DMF**, **DAP-2-H<sub>2</sub>O** and **DAP-3-CH<sub>3</sub>OC<sub>6</sub>H<sub>5</sub>**.



**Figure S3.**  $^1\text{H}$ NMR spectra of the **DAP-3-CH<sub>3</sub>OC<sub>6</sub>H<sub>5</sub>** and DAP without solvent digested in DMSO- $d_6$ . The labeled peaks (blue squares) represent DAP molecule.