Bisulfate anion clusters were introduced as soft joints to construct porous molecular crystals, one of which is converted to a single-crystalline hydrogen-bonded cross-linked organic framework H$_2$OF-6 through photo-irradiated thiol-ene reaction. Upon the uptake and the removal of phenol, the electrostatically repulsive bisulfate anions in the anion cluster of H$_2$OF-6 were disrupted and reformed, allowing the material to expand and contract rapidly.
An ultra-dynamic anion-cluster-based organic framework

Jayanta Samanta,1 Rick W. Dorn,2,3 Wenlin Zhang,1,* Xuanfeng Jiang,1,4 Mingshi Zhang,1 Richard J. Staples,5 Aaron J. Rossini,2,3,* and Chenfeng Ke1,6,*

SUMMARY
Porous organic frameworks that can dynamically expand their voids to guest uptake remain formidable to design. The challenge comes from finding the balance between rigidity and flexibility. Rigidity is required to retain the framework, and the flexibility is needed for reversible expansion/contraction. Herein, we introduced bisulfate anion clusters as soft joints to construct a single-crystalline hydrogen-bonded cross-linked organic framework HCOF-6. Upon the uptake of guest molecules, the anion clusters of HCOF-6 were disrupted, allowing the crystal to expand rapidly (<40 min) to more than twice its original length. Removing these guests restored the anion clusters and the crystallinity of HCOF-6 quickly (<10 min). These guest-induced material size expansions and contractions are highly reversible, and the dynamic anion cluster dissociation-reassociation is confirmed by molecular dynamics simulations and solid-state NMR experiments. The guest-induced size change of HCOF-6 highlights a new approach to synthesize dynamic framework materials by introducing anion clusters.

INTRODUCTION
Dynamic porous crystals that possess responsive framework architectures offer stimuli-responsive sorption and separation capabilities beyond just size-sieving.1 Featurong adjustable metal-ligand coordination geometries, a group of metal-organic framework (MOF) materials reversibly change their framework geometries upon the sorption of substrates, demonstrating gated sorption1,2 and breathing behaviors.3,4 Unlike these soft MOFs, it remains nontrivial to trigger guest-gated framework dynamics in crystalline organic framework materials, such as covalent organic frameworks (COFs) or hydrogen-bonded organic frameworks (HOFs). Compared with metal-ligand coordinations, the covalent bond in COFs and hydrogen bonds in HOFs have limited bond rotation capabilities. Therefore, to activate the framework dynamics, the adsorbed guests must possess strong interactions with the framework to compete against the multivalent solid-state packing forces and overcome the energetically demanding covalent bond rotation or hydrogen-bond disruption processes. Most reported dynamic COFs14–17 and HOFs18–20 are responsive to solvent molecules to maximize the substrate-framework interactions, with one exception is the demetallated amorphous woven COFs,21 which enabled large-sized substrate uptake.22 In comparison, hydrogen-bonded cross-linked organic frameworks23–25 (H2OFs) that are doubly cross-linked by rigid hydrogen-bonding networks and flexible covalent cross-linkers demonstrated crystalline-to-amorphous framework dynamics upon guest uptake.25 Upon the disruption of the hydrogen-bonded network, the flexible cross-linking moieties in H2OFs...
allow the framework to expand, thus activating dormant voids to accommodate more guests. However, to achieve the delicate balance between the framework's rigidity and flexibility, tightly bound substrates are still required to disrupt the strong hydrogen-bonded networks in H$_2$OFs. Hence, the development of dynamic porous organic frameworks calls for new synthetic methods to construct robust network architectures with easily accessible framework dynamics.

Here, we present a versatile method to construct H$_2$OF materials with high structural integrity and easily triggered framework dynamics by introducing anion clusters as responsive joints (Figure 1A). A linear monomer (LM, Figure 1C) and a three-arm monomer (TM) consisting of piperazine and diallylmelamine moieties were synthesized to reduce the π–π interactions between stacked layers in the solid state, owing to piperazine’s semi-rigid chair conformation. Protonation of the diallylmelamine moieties generated charge-reinforced hydrogen-bonded motifs, and the donor-acceptor-donor (DAD) diallylmelamine was converted to donor-donor-donor (DDD) diallylmelaminium (Figure 1B). The assembly of LM with NO$_3$- or HSO$_4$- afforded supramolecular polymers with linearly connected anion dimers, whereas TM formed porous single crystals possessing unique tetrakis-anion (HSO$_4$)$_4$ and tris-anion (HSO$_4$)$_3$ clusters at different temperatures. Covalently cross-linking the allyl groups in the preorganized molecular crystal through photo-irradiated thiol-ene single-crystal to single-crystal (SCSC) transformation afforded porous H$_2$OF-6 crystals with (HSO$_4$)$_3$ joints (Figure 1D). Compared with the disruption of cation-anion networks, such as guanidinium–, ammonium–, or amidinium–organo-sulfonate HOFs, a considerably smaller energy penalty is required for the disruption of anion dimers or clusters due to the electrostatic repulsions between anions. Hydrogen-bond competing substrates, such as phenol and its analogs, demonstrated fast and effective disruption and fragmentation of the anion clusters in H$_2$OF-6, activating the dormant voids to allow for a fast crystal expansion. Removal of the adsorbed phenol in H$_2$OF-6 rapidly contracted the framework and restored the anion cluster joint. Although anion clusters have been investigated in the solid-state supramolecular systems rarely, if ever, has an anion cluster been introduced to construct a porous network to enable network dynamics upon its association and dissociation. The reversible switching of H$_2$OF-6’s network between the crystalline contracted state and the amorphous expanded state bridges the gap between traditional rigid frameworks and swellable cross-linked polymers, providing new opportunities to design porous organic materials that allow the adsorption and separation of substrates beyond just size-sieving.

RESULTS

Of the more than 200 melaminium derivatives reported in the Cambridge Crystal Database, only a few examples demonstrated complementary hydrogen-bonding arrays of melaminium/anion in the solid state. To validate the feasibility of anion (dimer/cluster)-directed assembly of monomers in the solid state, LM was synthesized with two diallylmelamine moieties. A semi-rigid piperazine moiety was introduced to prevent strong π–π stacking between LMs in the solid state. We obtained single crystals of LM (Figure 2A, left) through solvent evaporation. The diallylmelamine moieties of LM were hydrogen-bonded in DA-AD fashion, and the piperazine moieties adopted the chair conformation in the solid state (Figures S26 and S27), affording one-dimensional (1D) supramolecular polymers (LM)$_n$. When LM was protonated by HNO$_3$, the single-crystal structure of [LM•2H$_2$]$_{14}$•2NO$_3$- (Figure 2A, middle) showed that pairs of diallylmelaminium were bridged by anion-dimers (NO$_3$)$_2$, forming complementary (DDD-AAA)$_2$ hydrogen-bonded arrays with...
strong hydrogen bonds (N–H...O–N = 2.09, 2.10, and 2.31 Å, Figure S28). Mixing LM and H_2SO_4 afforded single crystals of [LM•2H]^2+•2HSO_4^-. In the solid state (Figure 2A, right), hydrogen-bonded anion-dimers (HSO_4^-) were found to bridge neighboring pairs of diallyl-melaminiums of [LM•2H]^2+, forming (DDD-AAA)_2 hydrogen-bonding arrays with even shorter hydrogen bonds (N–H...O–S = 1.99, 2.08, and 2.14 Å, Figure S30). These LM-based assemblies were closely packed.
with no appreciable void space, and the chair-formed piperazine moieties prevented any strong π–π stacking (Figures S26–S31). The assembly of [LM•2H]^{2+} \cdot 2NO_3^- and [LM•2H]^{2+} \cdot 2HSO_4^- with complementary (DDD-AAA)_2 hydrogen-bonding arrays motivated us to synthesize a TM for H$_2$OF construction. TM was synthesized in four steps (Scheme S2): the mono-Boc-protection of piperazine, reaction with intermediate 2,4-diallylamino-6-chloro-1,3,5-triazine, followed by Boc-deprotection, and reaction with cyanuric chloride. Slow evaporation of a mixture of TM and fuming HNO_3 in dioxane-DMSO (v/v = 3:1) afforded single crystals of [TM•3H]^{3+} \cdot 3NO_3^- \cdot 3DMSO in the monoclinic P2_1/n space group (Figure 2B).
[TM●3H]^{4+}●3NO_3^-●3DMSO formed a distorted hexagonal framework with solvent-occupied voids, featuring a pore size of 24.0 × 16.5 Å. The diallylmelaminium moieties of three [TM●3H]^{3+} were connected by a cluster of three NO_3^- anions and three DMSO (Figures 2B, top and S33). Unfortunately, due to the participation of DMSO in the cluster, the hydrogen-bonded network of [TM●3H]^{3+}●3NO_3^-●3DMSO is highly unstable, and crystallinity was lost upon solvent removal or exposure to thiol cross-linkers.

When H_2SO_4 was mixed with TM, slow evaporation of the reaction mixture in dioxane-DMSO (v/v = 2:1) afforded single crystals of [TM●4H]^{4+}●4HSO_4^- (Figure 2C) in the monoclinic P2_1/n space group with unit cell parameters of a = 13.6804(2), b = 22.9165(4), c = 20.7502(3) Å, α = γ = 90°, β = 102.646(2)°. Different from [TM●3H]^{3+}●3NO_3^-●3DMSO, all three triazines of [TM●4H]^{4+}●4HSO_4^- were protonated. A unique rhombic tetrakis-anion cluster (HSO_4^-)_{4} was discovered to bridge four diallylmelaminium moieties (Figure 2C), forming a 4-fold (DDD-AAA) hydrogen-bonding array. In this tetrakis-anion cluster, individual HSO_4^- anions were stabilized by S-O-H●●●O-S hydrogen bonds, two of which also formed hydrogen bonds with the protonated central triazine of the [TM●4H]^{4+} (O●●●H-N^+ = 1.84 Å. Figure S37) at the upper and lower layers. Surprisingly, only two out of the three diallylmelaminium arms of the [TM●4H]^{4+} participated in anion cluster formation, leaving one diallylmelaminium/HSO_4^- ion-pair freely pointing toward the framework’s void space (Figure 2C (middle)). Although the allyl groups of [TM●4H]^{4+} surmounted the voids, only 7% of discontinuous voids were measured in this hydrogen-bonded network (Figure 2C, bottom).

The existence of one non-participating diallylmelaminium in [TM●4H]^{4+}●4HSO_4^- implied that there are other possible superstructures in the solid state. Therefore, we increased the crystallization temperature from room temperature to 80°C searching for a more stable network. After testing multiple solvent combinations for crystallization at 75°C-80°C range (Table S1), single crystals of [TM●3H]^{3+}●3HSO_4^- (Figure 2D) were successfully obtained at 80°C in a DMSO-nitrobenzene (v/v = 1:3) mixture. [TM●3H]^{3+}●3HSO_4^- was crystallized in the monoclinic C2 space group with unit cell parameters of a = 43.1933(15) Å, b = 23.2259(8) Å, c = 22.9907(5), α = γ = 90°, β = 91.430(2)°. In this hexagonal hydrogen-bonded network (Figure 2D), a tris-anion cluster (HSO_4^-)_{3} was discovered to bridge three diallylmelaminium units, forming a complementary 3-fold (DDD-AAA) hydrogen-bonding arrays with hydrogen bond distances ranging between 1.81–2.14 Å (Figure S43A). [TM●3H]^{3+} and (HSO_4^-)_{3} clusters were packed alternatively along the c-axis, stabilizing the hydrogen-bonded 2D hexagonal networks through weak π–π, anion–π, and CH–π interactions (Figure S43B). Linear hexagonal channels of void in [TM●3H]^{3+}●3HSO_4^- were measured to have a pore size diameter of 15.3 × 13.8 Å, and allyl groups decorated the pore surfaces (Figures 2D and S44).

To probe the reactivity and accessibility of the allyl groups in [TM●4H]^{4+}●4HSO_4^- and [TM●3H]^{3+}●3HSO_4^-, ethanethiol (ET) was diffused into these single crystals followed by photo-irradiation (Scheme S3). The formed thioether products were dissolved in DMSO-d_6 for ^1H NMR analysis. The measured thiol-to-thioether conversions (Table S2) showed that only 55% of the allyl groups were consumed in [TM●4H]^{4+}●4HSO_4^- due to the small discontinuous voids, whereas a complete thioether conversion was achieved in [TM●3H]^{3+}●3HSO_4^- (Figure S12). When the ethanedithiol (EDT) cross-linker was employed, an incomplete thiol-to-thioether conversion was again observed in the cross-linked crystalline H_2OF-5 (Figures S14–S18). 2D ^1H→^13C cross-polarization heteronuclear correlation (CP-HETCOR) magic-angle spinning (MAS) solid-state NMR spectra of H_2OF-5 confirmed the presence of...
unreacted allyl groups (Figure S17). Direct excitation $^{13}$C NMR spectra revealed that only ca. 70% of the allyl groups were cross-linked (Table S3 and Figure S16).

Photo-cross-linking EDT-diffused single crystals of [TM$d_3H]^3\cdot3\text{HSO}_4^-$ afforded single-crystalline H$_2$OF-6. When [TM$d_3H]^3\cdot3\text{HSO}_4^-$ crystals were soaked in neat EDT, elemental analysis revealed that an excess of EDT (EDT: TM = 4:1) was attached to the allyl groups in the product (Table S4). Hence, the synthesis of H$_2$OF-6 was optimized by gradually reducing the amount of EDT until a nearly stoichiometric thiol-to-thioether conversion was reached in the elemental analysis (EDT: TM = 3.05–3.10: 1, Table S4). Comparison of 2D $^1$H→$^{13}$C CP-HETCOR solid-state NMR spectra of H$_2$OF-6 recorded with short (0.1 ms) or long (2.0 ms) CP contact times (Figures 3D and S22) enabled the complete assignment of all $^1$H and $^{13}$C NMR signals (Figures 3B–3D). No appreciable olefin $^1$H or $^{13}$C NMR signals were observed in the 1D direct excitation $^1$H, $^1$H→$^{13}$C cross-polarization MAS (CPMAS) and direct excitation $^{13}$C solid-state NMR spectra (Figures S20–S23) of H$_2$OF-6, confirming the complete consumption of all allyl groups upon photo-cross-linking. New $^{13}$C and $^1$H NMR signals emerged in the solid-state NMR spectra of H$_2$OF-6 at 20–40 and 1–3 ppm, respectively, confirming the formation of thiether cross-linking moieties. Compared with the solution phase $^1$H NMR spectrum of [TM$d_3H]^3\cdot3\text{HSO}_4^-$ (Figure 3C), the proton resonances of triazinium (−N–$^1$H) and amine (N−$^1$H) of H$_2$OF-6 were shifted downfield (Figure 3D), indicating the deshielding effect of the tris-anion cluster in the solid state. After cross-linking, the olefin stretching and bending bands of H$_2$OF-6 at 1,500 and 900 cm$^{-1}$ disappeared, and no free thiol vibrational band was observed in the IR spectrum (Figure 3E), further supporting a complete allyl-to-thioether conversion.

Powder X-ray diffraction (PXRD, Figure 3H) analysis showed that H$_2$OF-6 maintained high crystallinity, which is consistent with the $^{13}$C solid-state NMR spectra that show relatively narrow $^{13}$C NMR signals for the triazinium carbon atoms. The single-crystalline morphology of H$_2$OF-6 was observed in SEM and TEM analysis (Figures 3F and 3G). In the vapor sorption analysis (Figure 3I), supercritical CO$_2$-activated H$_2$OF-6 adsorbed 103 cm$^3$/g STP of water, 105 cm$^3$/g STP of methanol, and 46 cm$^3$/g STP of toluene at 296 K, respectively, demonstrating its permanent porosity after solvent removal. Observation with an optical microscope revealed that large crystals of [TM$d_3H]^3\cdot3\text{HSO}_4^-$ were shattered into layers of transparent smaller crystals of H$_2$OF-6 (Figure 3A) after photo-cross-linking. Single-crystal X-ray diffraction (SXRD) analysis showed that the monoclinic [TM$d_3H]^3\cdot3\text{HSO}_4^-$ crystals were reconfigured to hexagonal H$_2$OF-6 crystals in the P$3_1$21 space group with the cell unit parameters of $a = b = 24.0599(11)$ Å, $c = 22.9674(9)$ Å, $\alpha = \beta = 90^\circ$, $\gamma = 120^\circ$. Anisotropic refinement of the backbone of H$_2$OF-6, which consists of [TM$d_3H]^3\cdot$ and (HSO$_4^-$)$_3$ cluster (Figure 3J), showed that TM moieties in H$_2$OF-6 possessed similar conformations to its crystalline precursor [TM$d_3H]^3\cdot3\text{HSO}_4^-. H_2$OF-6 retained distorted 1D porous channels after cross-linking (Figure 3K). The (HSO$_4^-$)$_3$ cluster was stabilized by the shallow bowl-shaped cavity of the [TM$d_3H]^3\cdot$ moiety at the lower layer through multivalent anion–π interactions (Figure 3L). However, the flexible thioether moieties in H$_2$OF-6 were heavily disordered in the crystal lattice, making unambiguous modeling of the thioether cross-linking motifs difficult (ball-stick in Figure 3J). To reveal the structural information of the flexible cross-linking moieties of H$_2$OF-6, we surveyed the nearby olefin group distances in the crystal precursor [TM$d_3H]^3\cdot3\text{HSO}_4^-$. H$_2$OF-6 distances between the nearby olefin groups suggested that the EDT cross-linking could either take place in an intralayer fashion parallel to the a/b plane (Figure 3J) to yield a stacked 2D polymer, or in an interlayer manner between two...
**Figure 3. Synthesis and characterizations of single-crystalline H$_2$OF-6**

(A) Synthesis of H$_2$OF-6 through thiol-ene SCSC transformation.
(B) Comparison of a (upper) 1D $^1$H→$^13$C CP MAS solid-state NMR spectrum of H$_2$OF-6 recorded with 8.928 kHz MAS and (lower) a direct excitation $^1$H solution NMR spectrum of [TM•3H]$^3+$•3HSO$_4^-$ in DMSO-d$_6$.
(C) Comparison of a (upper) 1D $^1$H spin-echo solid-state NMR spectrum of H$_2$OF-6 recorded with 50 kHz MAS and a (lower) $^1$H solution NMR spectrum of [TM•3H]$^3+$•3HSO$_4^-$ in DMSO-d$_6$.
(D) 2D $^1$H homonuclear dipolar decoupling, $^1$H homonuclear dipolar decoupling, a 2 ms CP contact time and $^1$H homonuclear dipolar decoupling, a 2 ms CP contact time and 8.928 kHz MAS. (B–D) All NMR spectra were recorded at room temperature (~300 K).
(E) IR spectra of [TM•3H]$^3+$•3HSO$_4^-$ and H$_2$OF-6.
(F and G) SEM and TEM images of H$_2$OF-6, respectively. Inset: electron diffraction pattern of the TEM image.
(H) Experimental and simulated PXRD profiles of [TM•3H]$^3+$•3HSO$_4^-$ (blue) and H$_2$OF-6 (red).
(I) Vapor sorption isotherms (water, methanol, and toluene) of supercritical CO$_2$ activated H$_2$OF-6 recorded at 296 K.
(J) Overlaid structures of [TM•3H]$^3+$•3HSO$_4^-$ (orange color, wireframe) and H$_2$OF-6 (capped sticks). The dithiolether moieties (ball-stick) were modeled to show the proposed intralayer cross-linking network. Hydrogen atoms were omitted for clarity.
(K) Packed structure and voids of H$_2$OF-6.
(L) The anion cluster and cross-linked DAM-H$^+$ moieties in H$_2$OF-6.
(M) Images of a H$_2$OF-6 crystal (upper) before and (lower) after expansion induced by soaking in DMSO at 80°C for 24 h and then cooling to room temperature.
neighboring layers to yield a 3D cross-linked network (Figures S49–S58), or a mixture of both. Since our previous investigation demonstrated that the macroscopic crystal expansion of HCOFs could reflect their cross-linking topology, 25 crystals of HCOF-6 were soaked in DMSO at 80°C to dismantle its hydrogen-bonded network for crystal expansion (Figure 3M). The HCOF-6•DMSO turned amorphous (Figure S60) and anisotropically expanded along the c-axis direction to more than twice its original size while shrinking slightly along the a/b plane. The anisotropic size expansion of HCOF-6 suggested that (1) the intralayer cross-linking is the predominant connection in HCOF-6, which allowed the crystal to greatly expand along the c-axis and mildly contract along a/b plane, and (2) a small amount of interlayer cross-linking also existed as defects 42 in HCOF-6, which prevented the exfoliation of HCOF-6 as 2D polymers. 43–45

Since the disruption of (HSO₄)₃ clusters in HCOF-6 could give rise to dynamic framework expansion, the stability of HCOF-6 was investigated against aqueous NaOH and BaCl₂/HCl (Figure 4A; Scheme S5). After soaking in these aqueous solutions, the HSO₄⁻ anions were extracted from the framework to afford crystal-like but amorphous polymer HCOF-6am (Figure S63). Re-protonation of the HCOF-6am using

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**Figure 4. Dynamic guest sorption in HCOF-6**

(A) Illustration of HCOF-6’s dynamic expansion upon adsorption of various guests and HCOF-6’s guest uptake capacities. (B) Time-dependent PXRD profiles of HCOF-6 upon aniline addition. Crystalline aniline sulfate nanocrystals were formed in the expanded framework. (C) PXRD investigations of the crystallinity recovery of HCOF-6am•(PhNH₃)⁺₂SO₄²⁻ upon MeOH or HCl/MeOH washing. (D) PXRD profiles of HCOF-6 upon three cycles of phenol uptake and contraction upon MeOH washing. (E) Images of a HCOF-6 single crystal upon three cycles of phenol uptake and MeOH washing.
H₂SO₄ or HCl/NaHSO₄ solutions did not regenerate crystalline H₂COF-6. Interestingly, when the organic base aniline was employed, the H₂COF-6 crystal expanded along the c-xis direction rapidly to more than twice its original length in 15–20 min at room temperature (Figure S65; Video S1). The dynamic size expansion was much faster and more extensive than H₂COF-6 in a NaOH methanolic solution (~130%, 96 h, Figure S61). Time-dependent PXRD analysis (Figure 4B) showed that H₂COF-6 gradually turned amorphous upon aniline adsorption. Meanwhile, a new set of diffraction peaks emerged, which closely matched the PXRD profile of aniline sulfate.¹⁶ This result suggested that the adsorbed aniline in H₂COF-6 was protonated by HSO₄⁻ to form nano-crystalline aniline sulfate.¹⁶ ¹H NMR desorption analysis revealed that 0.32 g/g (3.4 mmol/g) aniline was adsorbed (Figure 4A) and H₂COF-6 was converted into H₂COF-6am(PhNH₃)⁺₂SO₄²⁻. During the adsorption process, the crystallinity exchange took place between H₂COF-6 and the in situ formed aniline sulfate, and the process was also partially reversible. When H₂COF-6am(PhNH₃)⁺₂SO₄²⁻ was soaked in an HCl methanolic solution, the amorphous H₂COF-6am(PhNH₃)⁺₂SO₄²⁻ was partially converted to crystalline H₂COF-6 (Figure 4C). The regeneration of the H₂COF-6 crystalline framework was attributed to the in situ reverse ion exchange from the locally crystallized aniline sulfate, which seeded the ordered (HSO₄⁻)₃ reformation. In contrast, when the aniline sulfate nanocrystals in H₂COF-6am(PhNH₃)⁺₂SO₄²⁻ were removed by methanol washing, and the sample was then re-subjected to either H₂SO₄ or HCl/NaHSO₄ solutions, the crystallinity of H₂COF-6 was not recovered (Figure S67). These observations indicated that the transfer of entropy between aniline and highly ordered H₂COF-6 enabled the reversible crystallinity exchange, which also explains some of the unsuccessful attempts⁴⁷ to regenerate crystalline MOFs after demetallation and re-metallation cycles.

In the solutions of aniline analogs, such as cyclohexylamine, p-toluidine, and N-methylaniline (Figure 4A), H₂COF-6 also dynamically expanded its dormant void space and adsorbed 2.9, 3.0, and 2.1 mmol/g at room temperature, respectively (Table S8). In contrast, when N,N-dimethylaniline (DMA) was used, H₂COF-6 retained its crystallinity (Figure S68) and only adsorbed 0.7 mmol/g of DMA, despite its similar pKₐ value to others.⁴⁸ Since DMA lacks hydrogen-bond donating groups, we suspect that the disruption of the tris-anion clusters in H₂COF-6 using aniline and its analogs might be attributed to their hydrogen-bonding N–H groups more than their basicity. To confirm this hypothesis, H₂COF-6’s adsorptions capabilities on an acidic substrate phenol and its analogs cyclohexanol, p-cresol, anisole, 1- and 2-naphthol were investigated (Figure 4A). H₂COF-6 was immersed in concentrated dioxane solutions of these substrates, and the uptake capacities were measured by desorption ¹H NMR analysis (Table S9). Among them, only the adsorption of phenol and p-cresol in H₂COF-6 resulted in significant crystalline-to-amorphous phase transformations, suggesting that both the substrates’ chemical (hydrogen bonding) and physical (size) properties affect the framework dynamics.

In a concentrated phenol solution, H₂COF-6 crystals rapidly expanded their sizes along the c-axis direction to more than twice the original length in about 40 min at room temperature (Figure 4E; Video S2). The phase change from crystalline H₂COF-6 to amorphous H₂COF-6am•PhOH was clearly visualized (Video S2; Figures S70 and S71). The phenol adsorption-induced crystalline-to-amorphous phase transformation is highly reversible between the crystalline H₂COF-6 and amorphous H₂COF-6am•PhOH (Figures 4D and 4E). When H₂COF-6am•PhOH was washed by MeOH, the sample size quickly shrunk via phenol desorption (Figure 4E). In a parallel PXRD experiment (Figure 4D), the amorphous sample recovered its crystallinity and
converted back to H$_2$OF-6. Two more cycles of the phenol adsorption and desorption were repeated, and the crystal sample demonstrated reversible size expansion and contraction (Figures 4D and 4E; Video S2). Although H$_2$OF-6$_{am}$ ●PhOH appeared to be an amorphous material under the TEM (Figure S76), the hexagonal macroscale shape of the original crystal remained as revealed in SEM analysis (Figure S75) with a few cracks along the c-axis direction, supporting our suggested structure of H$_2$OF-6, which had small amounts of interlayer cross-links.

To reveal the origin of the phenol-induced anisotropic expansion, all-atom molecular dynamics (MD) simulations were performed on H$_2$OF-6 using the GROMACS package. We constructed a supercell with a slab of 4,000 phenols placed on top of H$_2$OF-6 as the initial state, which consisted of ●PhOH$^-$ anions (red) in H$_2$OF-6 (blue) and phenol (green) were color-coded, respectively.

(B) Supercell dimensions of interlayer and intralayer cross-linked H$_2$OF-6 measured in thermal-assisted annealing cycles (final cell size for intralayer H$_2$OF-6: 5.4 × 5.4 × 17.3 nm$^3$).

(C) A snapshot of a layer of the expanded H$_2$OF-6$_{am}$●PhOH Hydrogen atoms were omitted for clarity.

(D) 1D $^2$H solid echo ($\pi/2$–t–$\pi$–t–$\pi$) solid-state NMR spectrum of H$_2$OF-6$_{am}$●PhOH-d$_{6}$. The experimental spectrum (black) was deconvoluted by fitting to three Gaussian-Lorentzian peaks.

(E) 2D $^2$H→$^1$H CP-HETCOR NMR spectrum of H$_2$OF-6$_{am}$●PhOH-d$_{6}$ recorded with a 0.5 ms CP contact time. All NMR spectra were recorded at $B_0$ = 9.4 T with 50 kHz MAS.

(F) 2D $^1$H dipolar DQ-SQ homonuclear correlation NMR spectra of H$_2$OF-6 (red) with and (blue) without phenol-d$_{6}$.
c-axis expansions (<150%). In comparison, the 100% intralayer cross-linked virtual sample exhibited significant expansions (>250%) along the c-axis and mild contraction along the a/b plane, consistent with our suggested model, which had a predominantly intralayer cross-linked H$_2$OF-6. In the phenol expanded H$_2$OF-6 that are fully intralayer cross-linked, (HSO$_4^-$)$_3$ clusters were dissociated into (HSO$_4^-$)$_2$ anion dimers and HSO$_4^-$ anions in which the melaminium/HSO$_4^-$ ion pairs were further stabilized via hydrogen bonding interactions with phenol (Figure 5C). The disruption of the tris-anion clusters enabled phenol insertion between the layers of H$_2$OF-6 to expand the network, activating dormant voids to host more phenols in this expanded network (Figure 5A).

To confirm these simulated results experimentally, H$_2$OF-6$_{am}$●PhOH-d$_6$ (98% deuterated) was prepared for $^2$H solid-state NMR spectroscopy experiments. A direct excitation MAS ($v_r = 50$ kHz) $^2$H NMR spectrum (Figure 5D) of H$_2$OF-6$_{am}$●PhOH-d$_6$ revealed multiple $^2$H NMR signals with similar isotropic chemical shifts ($\delta_{iso}$) at ca. 5, 7, and 11 ppm but significantly different linewidths; narrow $^2$H NMR signals correspond to phenol-d$_6$, with fast motion, whereas broad $^2$H NMR signals likely correspond to phenol-d$_6$ with reduced motion. We note that all adsorbed phenol-d$_6$ contains some degree of motion (i.e., they are not static) as no spinning sidebands, which are indicative of a significant first-order quadrupolar coupling interaction were observed (Figure S74).$^{31-33}$ A 2D $^2$H→$^1$H CP-HETCOR NMR spectrum (Figure 5E) of H$_2$OF-6$_{am}$●PhOH-d$_6$ revealed correlations between the broad $^2$H NMR signal centered at ca. 11 ppm and the CH$_3$ $^1$H NMR signals of H$_2$OF-6 at ca. 2–3 ppm, which confirmed that some adsorbed phenols were in close spatial proximity to the H$_2$OF-6 framework. Comparison of 2D $^1$H dipolar double-quantum-single-quantum (DQ-SQ) homonuclear correlation solid-state NMR spectra$^{54,55}$ (Figure 5F) of H$_2$OF-6 (blue) and H$_2$OF-6$_{am}$●PhOH-d$_6$ (red) revealed similar N–H···H–N* off-diagonal correlations between the melaminium and amine $^1$H signals, suggesting that the local environment of melaminium/HSO$_4^-$ was only changed slightly after the formation of hydrogen bonds with the adsorbed phenols as indicated in the MD simulations.

**DISCUSSION**

In summary, we have successfully introduced deformable anion clusters to construct hydrogen-bonded networks and subsequently synthesized a dynamic hydrogen-bonded cross-linked organic framework H$_2$OF-6 through a thiol-ene SCSC transformation. We discovered a rare rhombic tetrakis-anion cluster (HSO$_4^-$)$_4$ and a triangle tris-anion cluster (HSO$_4^-$)$_3$ in the assembled molecular crystals, the latter of which directed the formation of a hexagonally packed hydrogen-bonded network. Photo-cross-linking the crystal precursor [TM●3H]$^{+3}$$\cdot$3HSO$_4^-$ afforded single-crystalline H$_2$OF-6 with large voids. H$_2$OF-6 exhibited rapid and reversible size expansion and contraction (>200%) upon guest uptake and switching between crystalline and amorphous phases. Molecular dynamics simulations and solid-state NMR experiments revealed that the fast size expansion of H$_2$OF-6 is attributed to the guest-induced disruption of the anion cluster, which is highly reversible for substrates, such as phenol, and partially reversible for aniline. Our strategy of introducing anion clusters to direct the synthesis of H$_2$OFs will not only expand the substrate scope for H$_2$OFs and promote their dynamic responses, but this synthetic method will also be directly transferable to the synthesis of HOFs and other supramolecular assemblies. Furthermore, the guest-induced dynamic network expansion and contraction features of H$_2$OF-6 demonstrated the potential to integrate the features of traditional elastic polymeric materials with rigid framework materials, which will provide unique sorption capabilities for separation and catalysis for the development of porous materials operated with low energy consumption.
EXPERIMENTAL PROCEDURES

Resource availability

Lead contact
Further information and requests for resources should be directed to and will be fulfilled by the lead contact, Chenfeng Ke (chenfeng.ke@dartmouth.edu).

Materials availability
All materials generated in this study are available from the lead contact without restriction.

Data and code availability
The X-ray crystallographic coordinates for structures reported in this study have been deposited at the Cambridge Crystallographic Data Centre (CCDC), under deposition numbers CCDC: 2075758–2075763 and 2078152. These data can be obtained free of charge from the CCDC via www.ccdc.cam.ac.uk/data_request/cif.

Crystallization of [TM₃H³⁺·3HSO₄]⁻
Single crystals of [TM₃H³⁺·3HSO₄]⁻ were obtained from a mixture of DMSO-nitrobenzene (2.5 mL, v/v = 1:3) solution containing TM (25 mg) and two drops of conc. H₂SO₄ (98%) were placed in a 20-mL scintillation vial and allowed to evaporate slowly at 80°C. Rhombic-shaped, colorless crystals appeared within 2 weeks. The surfaces of the crystals were washed by a mixture of 1:1 MeCN-MeOH solution three times to remove any by-products and were activated using supercritical CO₂ for the elemental analysis. IR (ν, cm⁻¹): 1,609 (S=O). Elemental analysis: for [C₄₂H₆₀N₂₄][H₂SO₄]₃.[CO₂]₄₂, calculated C: 40.21% H: 4.82% N: 24.36% S: 6.97%, found C: 39.17% H: 4.94% N: 25.44% S: 5.89%.

Synthesis of H₂OF-6
Single crystals of [TM₃H³⁺·3HSO₄]⁻ were washed by a mixture of 3:1 MeCN-MeOH solution three times to remove any by-products. Single crystals samples (~30 mg) were taken in a 20-mL glass vial to which 1 mL MeCN and 0.02 mL EDT were added and kept overnight in the dark. The mixture was replaced with a fresh portion of 0.02 mL EDT and 1 mL MeCN after 24 h. The reaction vial was irradiated under the UV light (medium-pressure 175-watt Hg lamp) for 72 h with forced air cooling. The crystal samples were collected and washed by an excess of MeCN to remove the unreacted EDT. MeCN solvent-containing samples were used for single-crystal X-ray analysis. The crystal samples were activated using supercritical CO₂ three times and used for different analyses. Elemental analysis: formula [C₄₂H₆₀N₂₄][H₂SO₄]₃.[EDT]₃.[CO₂]₄, calc: C 37.69%, H 5.13%, N 20.21%, S 17.73%, found: C 37.38%, H 5.45%, N 20.23%, S 17.89%. Solid-state ¹³C NMR (100 MHz): 161.4, 155.1, 44.3, 41.5, 33.1, 30.7, 25.8.

Regeneration of the crystallinity of H₂OF-6
Regeneration of the (HSO₄)₃⁻ anion clusters is critical for the reformation of crystalline H₂OF-6. Experimentally, H₂OF-6 sample (~5 mg) was placed on a non-diffracting PXRD plate, and its PXRD diffraction data were collected. Next, 30 µL of phenol was carefully added, immersing these crystals under liquid without spreading them out. The sample was kept still for approximately 15 min to allow extensive phenol adsorption. During this period, the plate was occasionally mounted on the instrument to check for crystallinity loss. Once the PXRD profile was almost lost, as shown further on, the excess of phenol was carefully removed by filter paper without disturbing the crystals. Thereafter, a MeOH solution (~100 µL) was gently added to
the crystals. The mixture was allowed to sit for 10 min, and PXRD data were collected. This procedure was repeated three times with the same sample.

Detailed procedures for the synthesis of monomers are described in the supplemental information, and they are fully characterized by NMR spectroscopy and high-resolution electrospray ionization mass spectrometry.

SUPPLEMENTAL INFORMATION

Supplemental information can be found online at https://doi.org/10.1016/j.chempr.2021.11.014.

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AUTHOR CONTRIBUTIONS

J.S. performed the synthesis and materials characterization under the supervision of C.K.; R.W.D. and A.J.R. performed the solid-state NMR spectroscopy experiments. W.Z. performed the MD simulations. X.J. initiated the project, and M.Z. performed some of the materials characterizations. R.S. collected the single-crystal data. The manuscript was drafted by C.K. with inputs from all co-authors.

DECLARATION OF INTERESTS

The authors declare no competing interests.

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