

1 **Nested Non-covalent Interactions Expand the Functions of Supramolecular Polymer**
2 **Networks**

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13 **Abstract**

14 Supramolecular polymer networks contain non-covalent cross-links that enable access to broadly
15 tunable mechanical properties and stimuli-responsive behaviors; the incorporation of multiple
16 unique non-covalent cross-links within such materials further expands their mechanical responses
17 and functionality. To date, however, the design of such materials has been accomplished through
18 discrete combinations of distinct interaction types in series, limiting materials design logic. Here
19 we introduce the concept of leveraging “nested” supramolecular crosslinks, wherein two distinct
20 types of non-covalent interactions exist in parallel, to control bulk material functions. To
21 demonstrate this concept, we use polymer-linked Pd₂L₄ metal-organic cage gels (polyMOCs) that
22 form hollow metal-organic cage junctions through metal-ligand coordination and can exhibit
23 well-defined host-guest binding within their cavity. In these “nested” supramolecular network
24 junctions, the thermodynamics of host-guest interactions within cage junctions affect the metal-
25 ligand interactions that form those junctions, ultimately translating to substantial guest-dependent
26 changes in bulk material properties that could not be achieved in traditional supramolecular
27 networks with multiple interactions in series.

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33 **Introduction**

34 Supramolecular polymer networks are formed through non-covalent bonds (e.g., metal–ligand
35 coordination, hydrogen bonding pairs, host-guest binding) between (macro)molecular strands and
36 junctions, leading to a wide range of network structures and properties from elastic materials to
37 highly ordered framework materials. Compared to conventional, covalently cross-linked polymer
38 networks, the reversibility of the non-covalent interactions used to construct supramolecular
39 polymer networks enables distinct functionalities (e.g., self-healing¹, stress dissipation²,
40 responsiveness to chemical stimuli³) and enhanced mechanical properties (e.g., extensibility and
41 toughness⁴),⁵ which facilitate a variety of applications, such as adhesives,⁶ biomaterials,⁷ and
42 stimuli responsive materials.³ In some cases, non-covalent bonds form discrete supramolecular
43 structures (e.g., metallacycles or metal–organic cages from metal–ligand coordination bonds or
44 extended pi-pi stacking clusters) which themselves can serve as cross-links. While such structures
45 can introduce advanced functionality, the mechanical properties and stimuli-responsiveness of the
46 resulting materials are generally defined by the underlying non-covalent bonds.

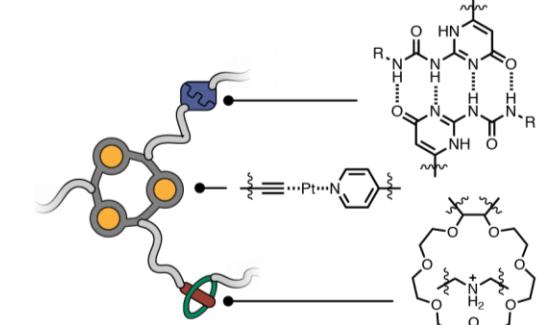
47 Researchers have combined multiple types or strengths of non-covalent bonds to construct
48 new materials with advanced functions (Figure 1A). In these examples, however, the interactions
49 used (*vide infra*) have been introduced as discrete, isolated interactions where the
50 (thermo)dynamics of one are independent of the other within the material structure.^{8–12} For
51 example, Stang and coworkers have used combinations of orthogonal metal–ligand coordination
52 (e.g., Pt²⁺-pyridine metallacycles), host–guest (e.g., ammonium/crown-ether complexes), and
53 hydrogen-bonding interactions to construct multi-responsive supramolecular polymer networks^{13–}
54¹⁶ that display self-healing behavior and stimuli-responsive gel-to-sol transitions under conditions
55 that weaken relative non-covalent bond strengths or increase dynamics (e.g., heating) or dilute
56 cross-linking functionality (e.g., addition of potassium salts to sequester host-guest cross-linking
57 sites). Since host-guest binding, hydrogen bonding, and metal–ligand coordination interactions are
58 independent in such systems, attenuating any of the interaction types leads to network
59 deconstruction. Moreover, strengthening only one non-covalent interaction types does not increase
60 the overall network stability to a competitor for the others.¹⁷ Similarly, others have reported
61 materials that display pH or solvent responsiveness made possible by a combination of host-guest
62 and electrostatic or hydrogen-bonding interactions.^{18,9} Nevertheless, while materials built from
63 host-guest interactions intrinsically have the potential for highly selective stimuli-responsive bulk

64 behaviors triggered by the presence of selective guests, such stimuli have so far been strictly
65 limited to causing cross-link displacement resulting in a decrease in stiffness or network
66 dissolution.^{19,20} We note that while host-guest binding can occur through ensemble non-covalent
67 interactions (e.g., hydrogen bonding, hydrophobic interactions, and van der Waals forces), these
68 interactions are defined by the host/guest pair and are not independently modifiable.

69 Others have combined non-covalent interactions of the same type but varied strength (e.g.,
70 a weak and a strong coordination bond) to fabricate materials (Figure 1A, right). For example,
71 Holten-Andersen and coworkers as well as Craig and coworkers have prepared end-linked or side-
72 chain linked metal–ligand coordination gels, respectively, by combining mixtures of metal ions
73 (e.g., Ni^{2+} , Zn^{2+} , Cu^{2+} , Pd^{2+} , Pt^{2+}) with a polymer-bound ligand.^{21–24} A similar concept has recently
74 been demonstrated in the context of Pd^{2+} and Pt^{2+} metal–organic cage (MOC)-crosslinked polymer
75 networks.²⁵ Nevertheless, while the combination of multiple supramolecular bond strengths
76 through mixing metals enables functions such as constant stiffness with widely variable stress
77 relaxation rates, the reliance on independent interactions limits the functions of these materials
78 similarly to those using different supramolecular interactions as described above. Overall, the use
79 of combinations of independent non-covalent interactions in series places inherent limits on the
80 functional scope of supramolecular polymer networks.

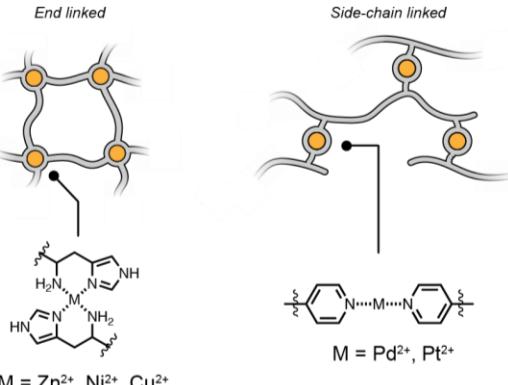
A Previous Work: Discrete Combinations of Multiple Supramolecular Interactions

Host-Guest, Metal-Ligand Coordination, and Hydrogen Bonding Interactions (Stang)



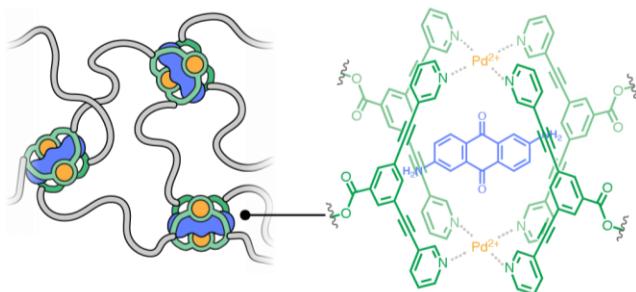
- + Broadened functionality and stimuli-responsiveness
- Traditional stoichiometric assembly limitations
- Supramolecular interactions are independent

Multiple Strengths of an Interaction Class
e.g. Metal-Ligand Coordination (Holten-Andersen, Craig)



- + Tunable dynamic properties
- Traditional stoichiometric assembly limitations
- Maintained stimuli-responsive properties

B This Work: Nested Metal-Ligand Coordination and Host-Guest Interactions



Nested Configuration Enables:

- + Modulation of global dynamics with guest strength
- + Expanded stoichiometric assembly window
- + Small-molecule triggered sol-gel transitions
- + Guest binding strength selective sol-gel transitions
- + Self-sorting driven sol-gel transition networks

Figure 1. Examples of supramolecular networks constructed with multiple interaction types or strengths. **A.** Previous examples of discrete combinations of multiple supramolecular interactions within polymer networks. Multiple distinct interaction types allow expanded functionality and stimuli-responsiveness, but ultimately suffer from the traditional limitations of supramolecular interactions in general (e.g., stoichiometric assembly limitations). Additionally, combinations of multiple strengths of the same interaction class enable exquisitely tunable dynamic properties but introduce no further network stability or stimuli-responsiveness. **B.** PolyMOC gels leverage the nested nature of metal-ligand coordination and host-guest interactions within their junctions to enable enhanced and expanded mechanical and stimuli-responsive properties.

We envisioned a new approach to controlling bulk supramolecular material properties by leveraging multiple supramolecular interactions in a nested or parallel configuration, such that the two interactions are not independent, as in the examples described above, but dependent on each

96 other (Figure 1B). In searching for a material to demonstrate this concept, we recognized that such
97 a configuration could be achieved with MOC-crosslinked polymers (polyMOCs).²⁶⁻²⁹ MOCs are,
98 in some cases, known to display pronounced host-guest binding ability,³⁰⁻³⁴ and a few reports of
99 polyMOCs have leveraged MOCs that can bind guest molecules,³⁵⁻³⁸ however, no studies have
100 explored how the bulk properties of polyMOCs can be varied by altering guest binding
101 thermodynamics, nor have they leveraged this unique nested junction structure to achieve
102 functions that cannot be obtained with traditional supramolecular networks. For example, we
103 imagined at least three functions that could potentially be uniquely achieved with nested polyMOC
104 networks: (1) guest-induced manipulation of bulk stress-relaxation dynamics, wherein small
105 molecule guests alter the metal–ligand coordination dynamics of the junction; (2) resistance to
106 dissolution under off-stoichiometry network formation conditions or in the presence of competitive
107 reagents due to synergistic stabilization of MOC junctions bound to guests; and (3) the ability to
108 control both percolation (i.e., gelation) and dissolution using selective self-sorting of MOC
109 junction driven by guest recognition. Here, we realize the synthesis of a new class of polyMOCs
110 containing acetylene-space Pd₂L₄ junctions and demonstrate each of the above functions using
111 both neutral and ionic guest molecules, establishing nested junctions as a new design principle for
112 supramolecular polymer networks.

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114 **Results & Discussion**

115 Initially, we targeted a Pd₂L₄ MOC based on bis-pyridyl ligands with acetylene spacers
116 due to its extensively studied host-guest binding properties and our previous successful translation
117 of related bis-pyridine MOCs to polyMOC materials. When the bis-*meta*-pyridine ligand **L1** is
118 mixed with 0.5 equiv. [Pd-(MeCN)₄][OTf]₂ in DMSO (5mM Pd²⁺) at room temperature,
119 supramolecular complex **MOC** is formed rapidly and quantitatively (Figure 2A). Although no
120 guest molecule is explicitly added, the triflate counterion is presumed to reside within the MOC
121 cavity (*vide infra*). Here, we note that coordinating solvents (e.g., DMSO or acetonitrile) are
122 necessary for MOC formation and subsequent polyMOC fabrication based on this MOC. MOCs
123 of this structure are known to strongly bind guests with electron-rich substituents within their
124 cavity, including (divalent) anions³⁹ or quinone-type molecules; however, previous studies have
125 largely focused on quantifying association constants within structurally similar Pd₂L₄ MOCs
126 bearing non-coordination counterions such as tetrakis[3,5-bis(trifluoromethyl)phenyl]borate

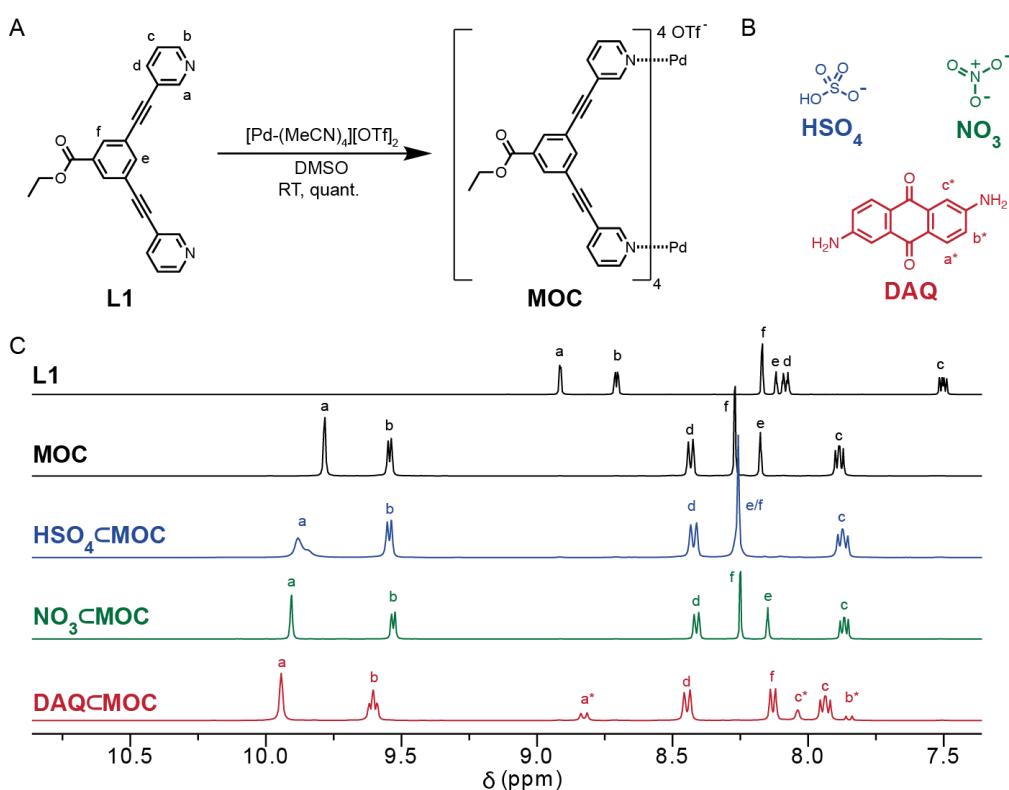
127 (BArF), cages in non-polar solvents (e.g., dichloromethane, nitromethane), or a combination
128 thereof.⁴⁰ Therefore, direct translation of previously reported extended-quinone guest molecules
129 proved unsuccessful, as either their binding strength with **MOC** is too low (in the case of
130 naphthoquinone, $K_a \sim 25 \text{ M}^{-1}$), or the guests exhibit insufficient solubility in DMSO (as was the
131 case with anthraquinone and pentacenequinone). Thus, we identified three previously unreported
132 guests for **MOC**: tetra butyl ammonium salts of hydrogen sulfate, **HSO₄**, or nitrate, **NO₃**, and 2,6-
133 diaminoanthraquinone, **DAQ** (Figure 2B). When 1 equiv of each guest is added to a 5 mM DMSO-
134 *d*₆ solution of **MOC**, distinct down-field peak shifts are observed via ¹H NMR spectroscopy
135 (Figure 2C). In line with previous reports, the interior proton resonances of **MOC** (H_a and H_b based
136 on labeling in Figure 2A) show the largest shifts upon guest addition, indicative of binding *within*
137 the **MOC** cavity. Association constants, K_a , for guest binding were quantified through titration
138 experiments and are listed in Table 1 (see Supplementary Figures 1-6). The K_a measurements fit
139 well to a 1:1 binding isotherm and span approximately three orders of magnitude, with binding
140 strength increasing in the order: **HSO₄** < **NO₃** < **DAQ**. To our knowledge, the association constant
141 of $62,000 \pm 2,000 \text{ M}^{-1}$ for **DAQ** is the largest reported for any Pd₂L₄ MOC in DMSO by three
142 orders of magnitude. While we could not quantify the binding strength of the triflate anion within
143 the cavity of this cage due to its inherent presence in the palladium salt used, we performed density
144 functional theory calculations of its binding strength using the B3LYP-D3(BJ) method,^{41,42} which
145 predict that the triflate anion will bind within a guest-free **MOC** with a favorable free energy
146 difference of 128.4 kJ/mol, suggesting near quantitative guest occupancy (triflate or otherwise)
147 under the conditions explored in this work.

148 To further probe the nature of these guest binding interactions within **MOC**, single-crystal
149 X-ray diffraction studies of **MOC** and its three host-guest complexes were performed. The crystal
150 structures of **MOC**, **HSO₄⊂MOC**, **NO₃⊂MOC**, and **DAQ⊂MOC** are shown in Figure 3. In all
151 cases, the added guest molecules reside within the cavity of the **MOC**, despite the variable ionic
152 nature of the guests explored. Consistent with ¹H NMR studies, hydrogen bonding interactions are
153 observed between guest molecules and interior proton H_a , with bond distances ranging from 2.35
154 to 2.75 Å for all added guests, suggesting that binding of these guests is not exclusively driven by
155 partial ion-exchange. Interestingly, guest orientation was consistent with that of **MOC** packing for
156 both **OTf⊂MOC** and **DAQ⊂MOC** (i.e., guest molecules were oriented in a consistent fashion
157 throughout the crystal) perhaps due to their size relative to the **MOC** cavity; however, the position

158 of **NO₃** and **HSO₄** within their respective hosts was observed to be split between ‘top’ and ‘bottom’
 159 hydrogen bonding sites, as judged by equal electron density in both positions, but with relative
 160 intensity ~half that of the MOC structure, consistent with 1:1 binding as observed from titration
 161 experiments. Despite having the highest measured K_a , **DAQ** shows the longest hydrogen bonding
 162 distances; complementary ArH··· π interactions are observed between H_e of the MOC and the
 163 flanking aryl rings of **DAQ**. This ArH··· π interaction is hypothesized to drive the stronger binding
 164 and larger ligand distortion and desymmetrization of the cavity in **DAQ**·**MOC**, as compared to
 165 **MOC**, **HSO₄·MOC** and **NO₃·MOC** as judged by the angles between pyridine ligands (an
 166 average deviation from 90° of 1.86° as compared to 1.03°, 1.07°, and 0.36°, respectively. See
 167 Figures S36-40 for more details). Such desymmetrization illustrates the flexural ability of MOCs
 168 to change their structure to selectively bind guests.⁴³

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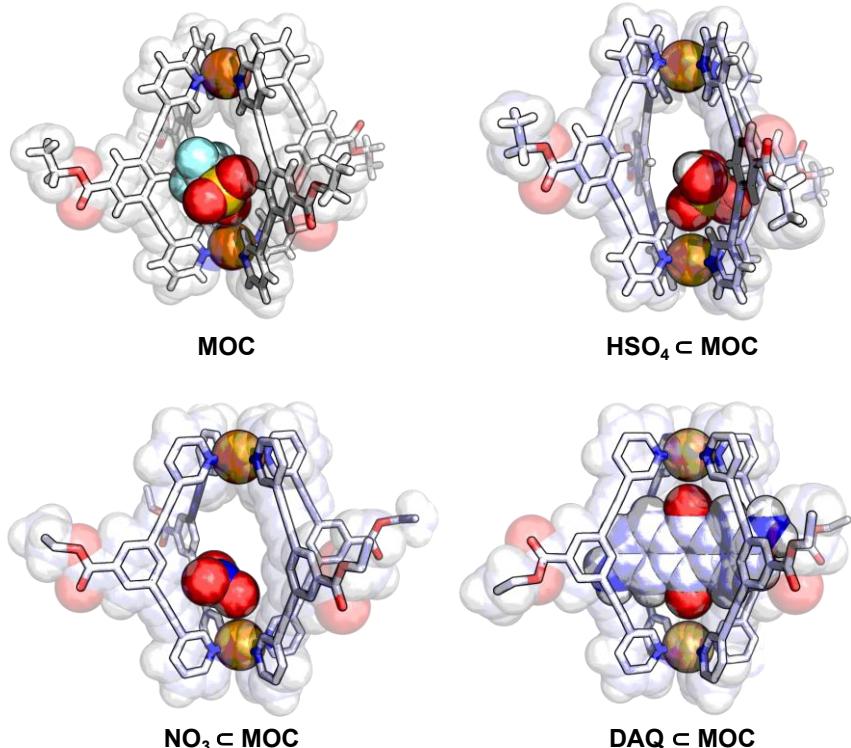


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172 **Figure 2.** **A.** Synthesis of **MOC-OTf** from **L1** and $[\text{Pd}-(\text{MeCN})_4][\text{OTf}]_2$ in DMSO. **B.** Chemical
 173 structures of the three guests used in this study: tetrabutylammonium salts of hydrogen sulfate
 174 (HSO_4^-), and nitrate (NO_3^-), and 2,6-diaminoanthraquinone (**DAQ**). **C.** Partial ^1H NMR spectra of
 175 **L1**, **MOC-OTf**, $\text{HSO}_4 \cdot \text{MOC}$, $\text{NO}_3 \cdot \text{MOC}$, and **DAQ**·**MOC** for 1 equiv. of guest added per
 176 cage (full spectra are shown in Supplementary Figures 25-28). Distinct down-field peak-shifts

177 relative to **L1** indicate successful MOC formation and subsequent guest-binding within the MOC
178 cavity.

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183 **Figure 3.** X-ray crystal structures of MOCs illustrating the presence of guest within the cavity:
184 **MOC**, **HSO₄⊂MOC**, **NO₃⊂MOC**, and **DAQ⊂MOC**. Counter ions and solvent molecules have
185 been removed for clarity.

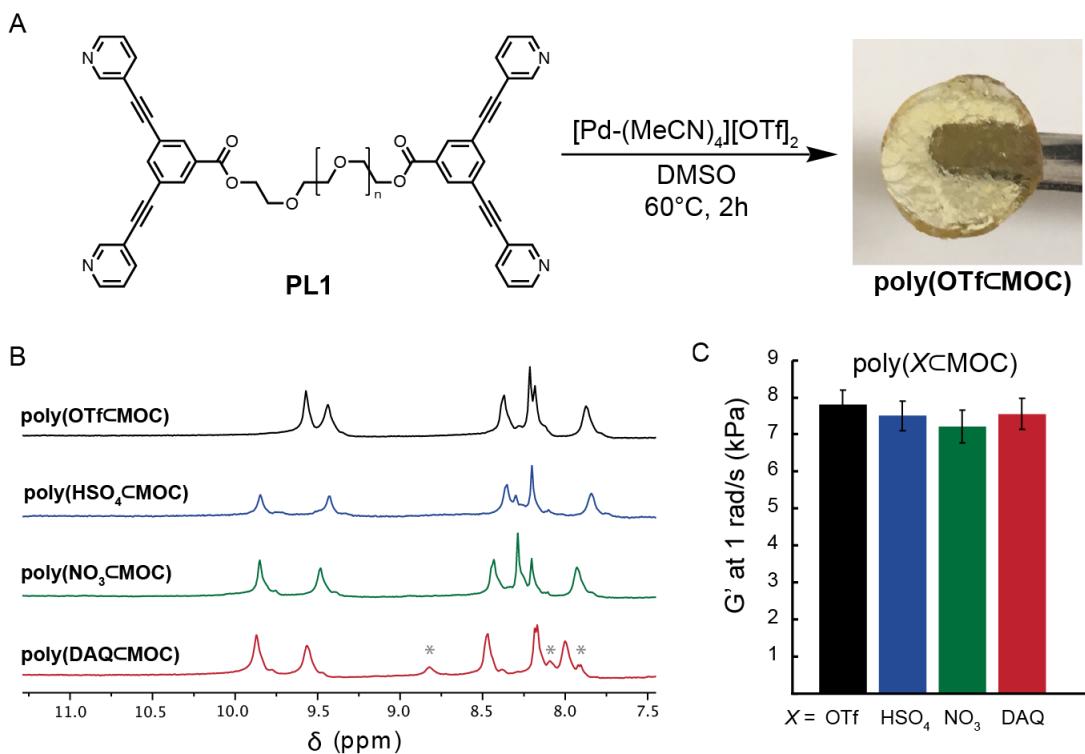
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187 **Table 1.** Measured Association Constants for guests with **MOC**.

Guest	Association Constant (M ⁻¹)
HSO ₄	1,000 ± 500
NO ₃	6,000 ± 1,000
DAQ	62,000 ± 2,000

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190 Having characterized the binding of **HSO₄**, **NO₃**, and **DAQ**, we next sought to verify that the same
191 host-guest interactions exist in an analogous polyMOC gel. A series of four polyMOC gels were
192 fabricated from polymer ligand **PL**—which contains **L1** ligands appended to the ends of a linear

193 poly(ethylene glycol) chain ($M_n = 4,600$ Da)—in DMSO (5 wt.% **PL**), with or without the
 194 inclusion of guest (Figure 4A). Gels were annealed for 2 h at 60 °C with inclusion of sufficient
 195 equivalents of each guest to achieve >95% binding occupancy (see Supplementary Table S1). This
 196 procedure yielded robust, free-standing materials (Figure 4A) in all cases, which we refer to as
 197 **poly($X \subset$ MOC)**, where X indicates the molecule residing within the MOC junctions: **OTf**, **HSO₄**,
 198 **NO₃**, **DAQ**. Cross-polarization magic-angle spinning (CP-MAS) ¹H NMR spectroscopy of the
 199 polyMOC gels show complete MOC assembly in all cases and distinct down-field resonance shifts
 200 when guests are added, as compared to the parent **poly(OTf \subset MOC)** (Figure 4B). The relative
 201 peak shifts from CP-MAS ¹H NMR are in-line with those observed in solution for discrete host-
 202 guest complexes, suggesting that the presence and nature of guest binding remains similar within
 203 the polyMOC gels. Moreover, resonances associated with **DAQ** are present in the
 204 **poly(DAQ \subset MOC)** spectrum, further supporting the formation of a host-guest complex effectively
 205 immobilizing **DAQ** within the MOC junctions of the material. The mechanical properties of each
 206 polyMOC gel were probed through shear rheology frequency sweep experiments. The gels
 207 exhibited near identical storage moduli (G') of 7 kPa at 1 rad/s (Figure 4C) indicating that the
 208 presence or absence of added guest molecules does not noticeably change junction assembly or
 209 network topology.⁴⁴



210 **Figure 4.** Fabrication and characterization of Pd_2L_4 polyMOC gels. **A.** A polymer ligand (**PL1**) is
211 combined with $[\text{Pd}(\text{MeCN})_4]\text{OTf}_2$ in DMSO. Annealing provides **poly(OTf**⊂**MOC**) which is
212 composed of Pd_2L_4 **MOC** junctions. Additional guest molecules can be included before annealing
213 to yield similar results. **B.** CP-MAS ^1H NMR spectra of polyMOC gels fabricated with and without
214 added guests. Relative resonance shifts of H_a within polyMOC gels are in-line with those of the
215 analogous MOCs (see Figure 1A), suggesting the guest binding remains unchanged when
216 translated to a polyMOC gel. Resonances marked with an asterisk in the **poly(DAQ**⊂**MOC**)
217 spectrum belong to cage-bound **DAQ**. Full spectra are shown in Figures S25–S28. Slight shoulder
218 peaks may be due to MOC clustering within the gel that could modify guest-binding behavior. **C.**
219 Comparison of gel shear modulus (G') as measured at 1 rad/s. Inclusion of guest has no statistically
220 significant impact on modulus. Full frequency sweep data are shown in Supplementary Figure 29.
221 Error bars represent ± 1 standard deviation.

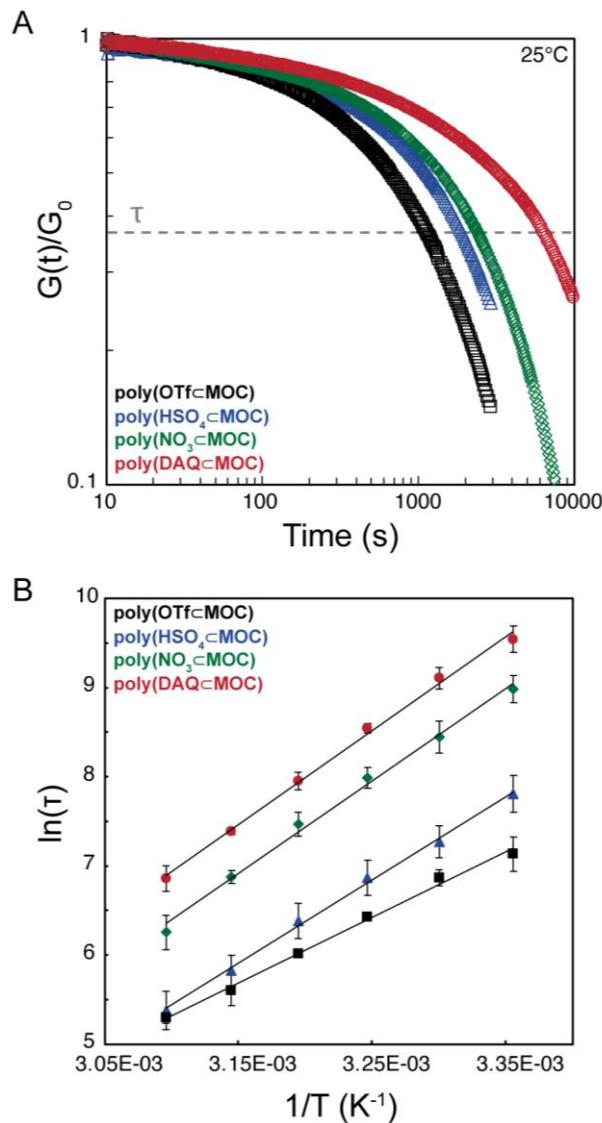
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223 Additionally, we studied the ability of pre-formed polyMOC gels to bind guests from solution.
224 When a 1 mL puck of **polyMOC-OTf** was placed into a solution of **DAQ** (10 mM in DMSO-d⁶),
225 within 60 s the surface of the gel began to darken, suggesting diffusion of **DAQ** into the gel. After
226 3 h, the gel was characterized by CP-MAS ^1H NMR; the spectrum displayed similar chemical
227 shifts to as assembled **poly(DAQ**⊂**MOC**) (Supplementary Figure 36), confirming **DAQ** diffused
228 into and formed a host-guest complex within the junctions of the network. This result suggests that
229 guest binding does not require the elevated temperature and increased ligand exchange dynamics
230 present during fabrication to enable guest binding.

231

232 Next, we sought to determine if the inclusion of guest-binding within the junctions of polyMOC
233 gels would affect their stress relaxation dynamics. Stress relaxation dictates the temporal response
234 of a material to applied forces and is a key parameter in the design of polymer networks for use as
235 adhesives, injectable biomaterials, and cell scaffolds.⁷ We reasoned that the inclusion of host-guest
236 binding may serve to stabilize the MOC junctions against the solvent- or ligand-assisted exchange
237 reactions at palladium that lead to microscopic network restructuring and ultimately macroscopic
238 stress relaxation.⁴⁵ Stress-relaxation studies conducted at 25 °C show that the presence of host-
239 guest binding leads to an increased characteristic relaxation time concomitant with the strength of
240 guest binding, up to approximately an order of magnitude greater for **poly(DAQ**⊂**MOC**) as
241 compared to **poly(OTf**⊂**MOC**) (Figure 5A). To gain further insight into how guest-binding
242 affects stress relaxation, variable temperature experiments were performed, and an Arrhenius
243 analysis was conducted (Figure 5B). The extracted activation energies for each gel are listed in
244 Table 2. Activation energy was found to trend upward with guest binding strength, indicating that

245 the critical ligand dissociation step required for stress relaxation has a higher activation barrier for
 246 more stable host-guest complexes. This result highlights the ability for host-guest binding to
 247 stabilize the metal-ligand coordination interaction through the unique nested configuration present
 248 in this system. Such behavior contrasts with the properties observed for previously reported
 249 supramolecular gels with multiple discrete interactions, where, upon incorporation of additional
 250 interaction types and strengths, the terminal relaxation mode is unaffected.⁴⁶ Further studies
 251 regarding these effects are ongoing in our laboratory.



252

253 **Figure 5.** Stress relaxation studies on polyMOC gels. **A.** Room temperature step-strain (2%)
 254 relaxation curves of **poly(OTf<MOC)**, **poly(HSO₄<MOC)**, **poly(NO₃<MOC)**, and
 255 **poly(DAQ<MOC)**. **B.** Arrhenius plot of stress relaxation time (τ) versus inverse temperature for

256 **poly(OTf⊂MOC)**, **poly(HSO₄⊂MOC)**, **poly(NO₃⊂MOC)**, and **poly(DAQ⊂MOC)**. Thermal
257 activation energies are listed in Table 2. Error bars represent the 99% confidence interval.
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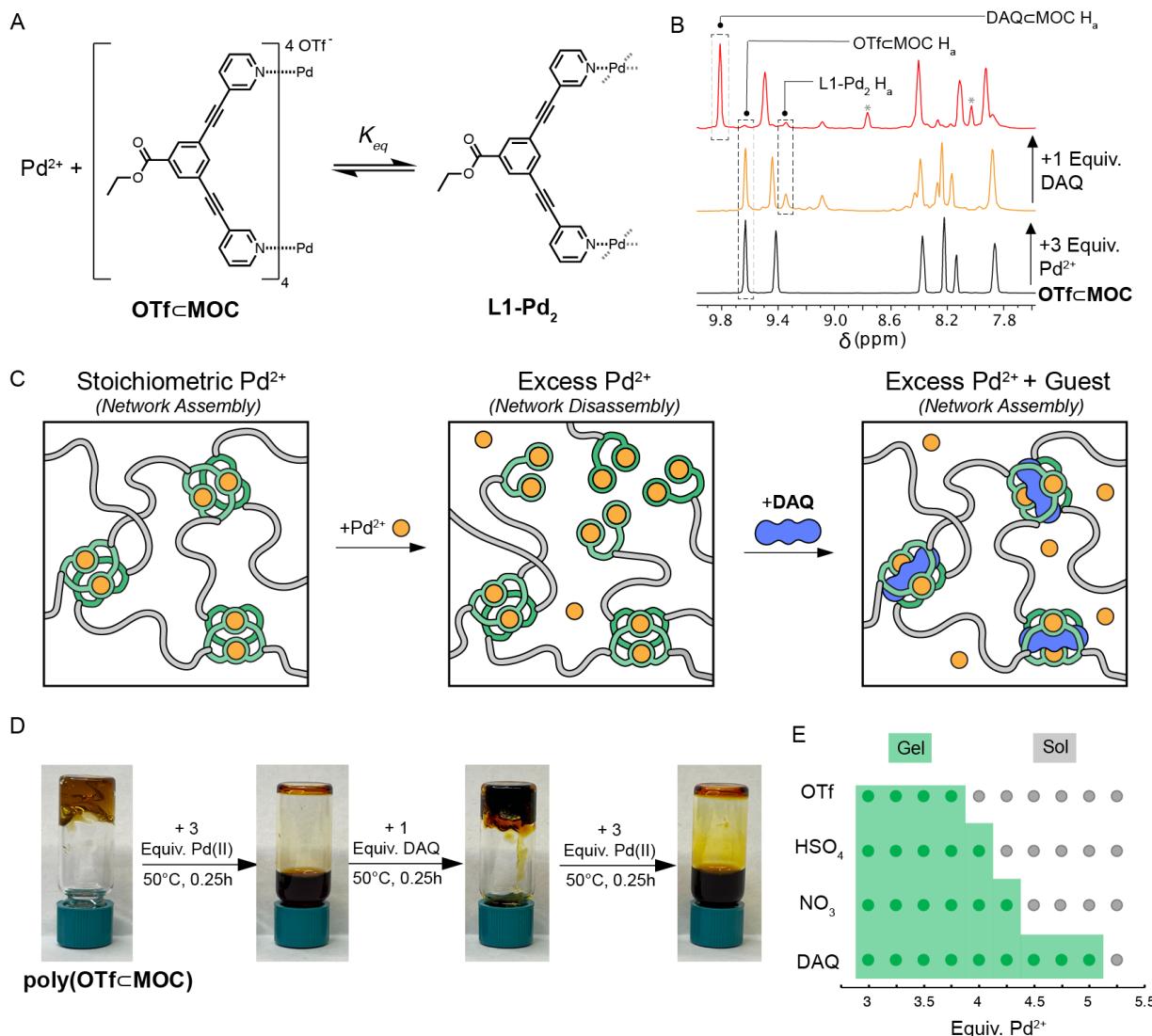
259 **Table 2.** Measured Activation Energies for Stress Relaxation in polyMOC gels.

polyMOC Gel	E _a (kJ·mol ⁻¹)
poly(OTf⊂MOC)	61 ± 3
poly(HSO ₄ ⊂MOC)	78 ± 5
poly(NO ₃ ⊂MOC)	86 ± 3
poly(DAQ⊂MOC)	88 ± 4

260
261 Having confirmed guest binding within the junctions of polyMOC gels and its ability to modulate
262 the rates of ligand-exchange reactions at the molecular level, which translate to bulk stress
263 relaxation differences, we hypothesized that guest inclusion could provide a means to stabilize
264 polyMOC gels under conditions that are typically incompatible with network formation, such as
265 in the presence of super-stoichiometric metal ion. Stoichiometry is critical for the formation of
266 both traditional and supramolecular polymer networks, and previous studies have explored the
267 impact of off-stoichiometry components on metallo-gel properties.^{47,48} With sub stoichiometric
268 metal, cross-linking density is proportional to metal ion concentration; when there is excess metal,
269 incomplete, lower-functionality metal-ligand complexes form, precluding gelation. In our
270 polyMOCs, addition of superstoichiometric Pd²⁺ would therefore be expected to drive MOC
271 disassembly and produce free ligands coordinated to two palladium ions (Figure 6A). To test the
272 potential ability of guests to stabilize MOC structures under these conditions, assembly of **MOC**
273 was attempted with 4 equiv. of [Pd(MeCN)₄][OTf]₂. When using this super-stoichiometric Pd²⁺, a
274 mixture of products is obtained as supported by ¹H NMR (Figure 6B, orange spectrum). The ¹H
275 NMR spectrum shows a combination of **MOC** (65%) and what we attribute to an **L1-Pd₂**
276 coordination structure (~35%); however, in the presence of 1 equiv **DAQ**, the ¹H NMR spectrum
277 of the mixture shows an increased intensity of the peaks attributed to **MOC** and a concomitant
278 decrease in intensity of **L1-Pd₂** (Figure 6B, red spectrum), indicative of shifting the equilibrium
279 toward MOC formation (88%). Within the context of a polyMOC gel, this guest-driven
280 stabilization in the presence excess Pd²⁺ could be expected to drive sol-gel transitions if the initial
281 degree of disassembly pushed the system below the gel point as illustrated in Figure 6C.
282 Correspondingly, the attempted fabrication of **poly(OTf⊂MOC)** with 4 equiv [Pd-

283 $(\text{MeCN})_4[\text{OTf}]_2$ resulted in a free-flowing liquid. Interestingly, addition of 1 equiv **DAQ** to this
284 liquid yielded a robust, free-standing gel with a storage modulus of 2,300 Pa. This gel could then
285 be dissolved through subsequent addition of 3 equiv Pd^{2+} . CP-MAS ^1H NMR spectroscopy of these
286 materials confirmed guest-binding (see Supplementary Figures 31-33).

287 Expanding on this process, three sequential sol-gel transitions were achieved by alternating
288 addition of **DAQ** and Pd^{2+} (Figure 6D). Both HSO_4 and NO_3 can induce a sol-gel transition in
289 polyMOC gels in the presence of excess Pd^{2+} , but in line with their lower binding affinities, the
290 extent was less pronounced than for **DAQ**. A sol-gel phase diagram for assembly of polyMOC
291 gels containing 1 equiv of each guest and between 1 and 5 equiv Pd^{2+} was constructed (Figure 6E),
292 showing that the amount of excess Pd^{2+} that can be tolerated before gelation is precluded is
293 proportional to guest binding strength. Materials with 4 equiv Pd^{2+} underwent a sol-gel transition
294 when 1 equiv of all guests were added, representing an unselective guest-sensing response. In
295 contrast, systems with 4.5 equiv Pd^{2+} only underwent a sol-gel transition in the presence of **DAQ**.
296 Addition of >10 equiv HSO_4 or NO_3 under these conditions failed to elicit a sol-gel transition,
297 illustrating the ability to engineer selective guest-triggered sol-gel transitions limited by guest



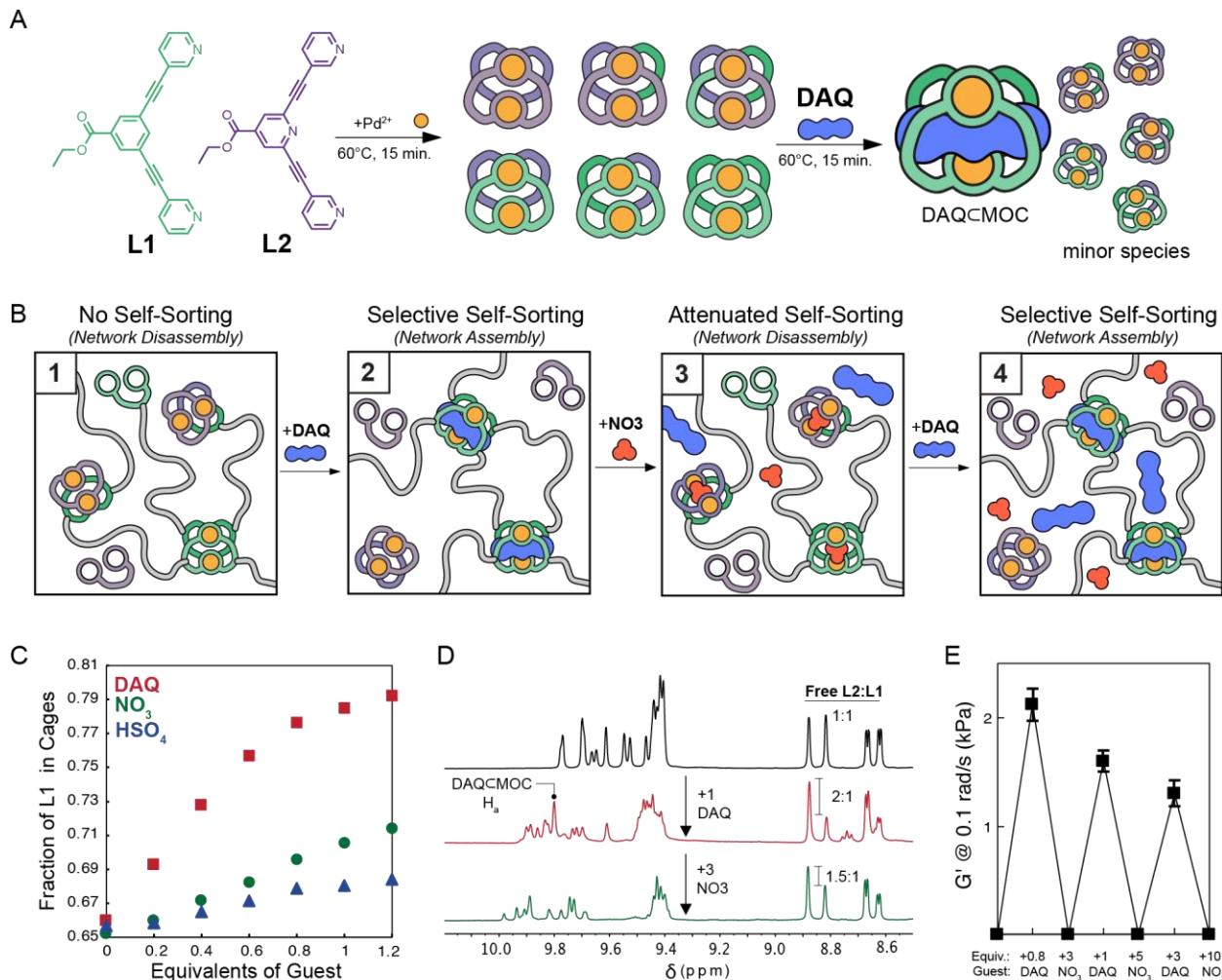
300 **Figure 6.** Effect of excess Pd^{2+} on (poly)MOC assembly **A**. Schematic of the MOC-ligand
 301 equilibrium in the presence of excess Pd^{2+} . Addition of super-stoichiometric Pd^{2+} drives the
 302 equilibrium to the right, producing free ligands coordinated to Pd (**L1-Pd**). **B.** ^1H NMR spectra of
 303 **MOC-OTf** assembled on stoichiometry (1 equiv Pd^{2+}), and with super-stoichiometric Pd^{2+} (4
 304 equiv) with and without added **DAQ**. **C.** Schematic of network degradation in the presence of
 305 super-stoichiometric Pd^{2+} and subsequent gelation upon addition of **DAQ**. Super-stoichiometric
 306 Pd^{2+} drives MOC structures to into L-Pd₂ structures, lowering cross-linking functionality and
 307 ultimately causing network degradation. Addition of **DAQ** stabilizes MOC structures in the
 308 presence of super-stoichiometric Pd^{2+} and subsequently drives MOC formation and network
 309 gelation. **D.** Photographs of sol-gel transitions caused by sequential additions of Pd^{2+} and **DAQ** to
 310 **poly(OTf^cMOC)**. **E.** Phase diagram for gelation with 1 equiv. of each guest and varying super-
 311 stoichiometric Pd^{2+} . The amount of Pd^{2+} that can be tolerated before gel degradation is proportional
 312 to guest binding strength. Gelation was assessed by the vial inversion test.
 313

Motivated by the ability to drive sol-gel transitions upon guest introduction for systems with super-stoichiometric Pd^{2+} , we sought to explore a complementary mechanism to expand the scope of selective and predictable material property changes in polyMOC systems (for example, the possibility for both sol-to-gel and gel-to-sol transitions driven by binding of the same guest, as well as reversible phase transitions when selective guest binding is attenuated). Previous reports have demonstrated that guest binding can drive self-sorting within heteroleptic MOCs formed from mixtures of chemically distinct but isostructural ligands.⁴⁹⁻⁵¹ Specifically, in the absence of guest, statistical mixtures of heteroleptic cages arise; however, when a guest is added, the equilibrium is shifted toward the cage structure that binds the guest most strongly, typically resulting in narcissistic self-sorting (i.e., homoleptic MOCs are preferentially formed). A schematic for such behavior is shown in Figure 7A for a mixture of **L1** and an isostructural endo-pyridyl variant, **L2**, that on its own will form cages with no binding affinity for **DAQ** (*vide infra*). We reasoned that the same mechanism could be leveraged to produce topological changes in polyMOC gels fabricated with mixtures of distinct polymer-bound and small-molecule ligands that on their own would form MOCs of the same architecture, but with varying abilities to bind a specific guest as illustrated in Figure 7B. If such self-sorting were sufficiently selective, sol-gel transitions could be achieved through the preferential incorporation of polymer-bound ligands into MOC structures *via* a favorable guest binding event (Figure 7B panel 1 to 2). Further, such sol-gel transitions could have the potential to be reversible upon addition of a guest that non-selectively binds cage structures from the ensemble (Figure 7B panel 2 to 3). With further addition of the selectively binding guest, another sol-gel transition may be achieved (Figure 7B panel 3 to 4); this cycle could potentially be repeatable if stronger binding or greater equivalents of each type of guest (selective versus non-selective binding) were added.

Guided by previous reports that the endo-pyridyl variant of **MOC** displays attenuated binding of quinone-type guests,⁵² we synthesized the corresponding endo-pyridyl ligand **L2** (Figure 7A) to explore its impact on MOC and polyMOC assembly in the presence of added guest and potential to enable self-sorting. Using only **L2** under standard MOC assembly conditions **N-MOC** was formed quantitatively. Titration experiments showed that unlike **MOC**, **N-MOC** displayed no affinity for **DAQ** ($K_a \ll 1 \text{ M}^{-1}$) and had a reduced affinity for both **HSO₄** and **NO₃** (association constants of $33 \pm 10 \text{ M}^{-1}$ and $2,200 \pm 400 \text{ M}^{-1}$). This difference in affinity for **DAQ** between **MOC** and **N-MOC** is sufficiently strong to drive considerable self-sorting to

345 preferentially incorporate **L1** into MOC structures when equimolar amounts of **L1** and **L2** (0.5
346 equiv total) were mixed super-stoichiometrically with 0.375 equiv [Pd(MeCN)₄][OTf]₂ in DMSO
347 (5 mM solution of Pd²⁺). For this system we first quantified the preferential incorporation of **L1**
348 into MOC structures upon guest addition using ¹H NMR spectroscopy, as H_a and H_b resonances
349 for un-bound **L1** and **L2** could easily be resolved. All guests were observed to preferentially drive
350 **L1** incorporation into MOC structures with an extent proportional to the differential guest binding
351 strength between **MOC** and **N-MOC** (Figure 7C). The greatest shift in ligand incorporation
352 occurred in the presence of >1 equiv **DAQ**, resulting in a 12% increase in the fraction of **L1** in
353 MOC structures, which corresponds to a 50% increase in the selectivity for **L1** incorporation over
354 **L2**. Interestingly, when 3 equiv **NO₃** were added to the mixed ligand system containing 1 equiv.
355 **DAQ**, the ratio of free to cage bound **L2:L1** was quickly shifted (within 15 min at room
356 temperature) from 2:1 to 1.5:1 (Figure 6C), illustrating the ability to reversibly modulate ligand
357 self-sorting *in situ* through introduction of less-selective guests.

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Figure 7. Guest-driven self-sorting in MOCs and polyMOCs containing **L1** and **L2** moieties. **A.** Schematic showing the statistical mixture of heteroleptic MOCs initially formed through mixing **L1** and **L2**. When a guest that preferentially binds one of the homoleptic MOC structures is introduced, this homoleptic MOC and subsequent host-guest complex is preferentially enriched relative to other MOC structures. **B.** Schematic of self-sorting driven sol-gel transitions within mixtures containing polymer-bound and small molecules ligands (**PL** and **L2**, respectively) with sub-stoichiometric Pd^{2+} . **C.** Fraction of **L1** incorporated into MOC structures when **L1** and **L2** are present in equimolar amounts with sub-stoichiometric Pd^{2+} (33.3% excess of bispyridine ligands). The preference to incorporate **L1** into MOC structures increases with strength and equivalents of guest added. **D.** ^1H NMR spectra showing the reversible enrichment of **L1** into MOC structures upon inclusion of **DAQ**, followed by depletion when the less selective guest, **NO₃**, is added. **E.** Reversible sol-gel transitions achieved through self-sorting behavior. An initial mixture of **PL1**, **L2**, and Pd^{2+} form a free-flowing liquid that becomes a gel upon addition of **DAQ**. Error bars represent ± 1 standard deviation.

375 Looking to translate this self-sorting effect into a polyMOC gel system, an optimized ratio of
376 polymer-bound and small-molecule ligands was required. Using Macosko-Miller theory,^{53,54} we

377 modeled the predicted network structures that could be formed through arbitrary combinations of
378 **PL**, **L2**, and **Pd**²⁺ and derived gelation criteria based on the proportions of **L2** : **PL** and of **L2** +
379 **PL** : **Pd**²⁺, and a self-sorting preference factor, denoted as $\rho_{SML:PL}$, $\rho_{L:Pd^{2+}}$, and ϕ , respectively.
380 Here, ϕ describes the preference for incorporation of polymer-bound bis-pyridine ligands into
381 MOC structures and is equal to the fraction of **L1** moieties in MOC structures divided by the
382 fraction of all ligand moieties in the mixture that are **L1**. We derived the following inequality
383 which predicts gelation to occur when:

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$$385 \left(\frac{1}{\rho_{SML:PL} + 1} \right) \frac{\phi}{2\rho_{SML:PL}} > \frac{1}{\sqrt{\frac{1 + \rho_{SML:PL}}{\frac{1}{2} \rho_{L:Pd^{2+}}} (f_w - 1)}} \quad (1)$$

386 where f_w is the weight-averaged functionality of the MOC network junction; in this case f_w
387 = 4, but we note Eq.1 is general for any value of f_w . Because $\rho_{SML:PL}$, $\rho_{L:Pd^{2+}}$, and f_w are fixed
388 according to the initial component stoichiometry and MOC system, it is clear how a shift in ligand
389 self-sorting into MOC structures (i.e., shifting the value of ϕ triggered by a small-molecule guest)
390 could cause a shift in the gelation criteria and induce a sol-gel transition. Guided by this analysis,
391 we found that a combination of 1 equiv **PL1** (two ligands per equivalent), 0.25 equiv **L2**, and 1
392 equiv [Pd(MeCN)₄][OTf]₂ (5 mM Pd²⁺ in DMSO) resulted in a free-flowing liquid even after
393 extended annealing. Upon addition of 1 equiv **DAQ** from a saturated solution in DMSO and
394 further brief annealing (15 min at 50 °C), however, the mixture underwent a sol-gel transition,
395 yielding a free-standing gel with a storage modulus of 2,100 ± 50 Pa as measured at 1 rad/s.
396 Subsequent addition of 3 equiv **NO₃** from a saturated DMSO solution caused this material to
397 transform into a free-flowing liquid. Further addition of increasing equivalents of **DAQ** and **NO₃**
398 in an alternating fashion allowed us to achieve a total of 6 sol-gel transitions in a single material
399 (Figure 6D). A marginal decrease in modulus was observed between each sol-to-gel transition,
400 which we attribute to a combination of slight dilution upon guest solution addition and a reduction
401 of the maximum extent of self-sorting in the presence of the less selective guest, **NO₃**. Nonetheless,
402 we anticipate that the results shown here to trigger a *sol-to-gel* transition for polyMOC gels
403 selectively in the presence of a suitable guest will significantly expand the utility of sol-to-gel
404 transitions as a signal transduction mechanism for sensing applications and provide a general

405 framework for the design of stimuli-responsive materials *via* guest-biding within tailor-made MOC
406 cavities in general.

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408 Herein, we have introduced a strategy to incorporate two distinct supramolecular interactions in a
409 nested configuration within a single cross-link of a polymer network based on Pd_2L_4 MOC
410 junctions. This configuration of metal-ligand coordination and host-guest binding interactions was
411 shown to enable unique functionality inaccessible with networks containing these interaction types
412 in a discrete configuration. Introduction of guest molecules enabled tuning gel relaxation time by
413 over an order of magnitude while maintaining constant stiffness. Further, the introduction and
414 modulation of guest binding strength was shown to enhance the stability of polyMOC gels in the
415 presence of super-stoichiometric Pd^{2+} and competing pyridine-containing ligands, illustrating the
416 synergistic stabilization property of two interactions in parallel within a single network junction.
417 Selective sol-gel transitions gated by guest binding strength could also be realized by simply
418 adjusting the amount of excess Pd^{2+} in the system. Finally, selective guest binding was observed
419 to drive MOC self-sorting, which could be leveraged to enable up to six sequential sol-gel
420 transitions in a single polyMOC gel material composed of a mixture of polymer-bound and small-
421 molecule ligands. A general theoretical description of this phenomena was presented which will
422 aid in the extension of these properties to other variations of polyMOC gels. Notably, guest-
423 dependent phenomena all trended proportional to guest binding strength despite their variable ionic
424 nature, suggesting that K_a serves as a general description for tuning such functions and properties
425 in the future. Altogether, these results expand the functionality and stimuli-responsive properties
426 achievable within supramolecular polymer networks through the judicious design of network
427 junction chemistry and structure. In the future, the nested supramolecular junction concept
428 introduced here may leverage the vast existing body of known MOC host-guest binding properties
429 to accelerate the creation of soft materials with exquisitely selective small-molecule-driven
430 property changes for applications ranging from sensing to biomaterials.

431

432 **Methods**

433 *Materials*

434 Solvents were purchased from Millipore Sigma and used as received unless otherwise noted. All
435 aqueous solutions were prepared with deionized water. Deuterated solvents were purchased from

436 Cambridge Isotopes Laboratories, Inc. Cambridge Isotope Laboratories, Inc. and used as received.
437 3,5-dibromobenzoic acid, ethyl-3,5,-dibromoisonicotinate, and poly(ethylene glycol) were
438 purchased from Milipore Sigma. 3-Ethynylpyridine was purchased from AmBeed Inc.
439 Tetrakis(acetonitrile)palladium(II) bis(trifluoromethanesulfonate) was purchased from TCI
440 America.

441 *Nuclear magnetic resonance spectroscopy (NMR)*

442 ^1H NMR spectra were recorded using either a 400, 500, or 600 MHz Bruker AVANCE
443 spectrometer at room temperature (25°C). The specific instrument used for each experiment is
444 indicated by the reproduced spectra or listed chemical shifts. All chemical shifts are reported as
445 parts per million (ppm) with splitting patterns designated as follows: s (singlet), d (doublet), t
446 (triplet), q (quadruplet), m (multiplet), and br (broad). Cross polarization magic-angle spinning
447 nuclear magnetic resonance (CP-MAS NMR) spectra were collected using 33.1 μL zirconia
448 spinners. Gels were made as described below, transferred to the spinners, sealed, and then allowed
449 to anneal in the spinner at the indicated time and temperature. Spectrum were collected at a
450 spinning rate of 5000 Hz on a 500 MHz Bruker AVANCE NMR spectrometer.

451 *Mass Spectrometry (MS)*

452 High-resolution mass spectrometry (HRMS) was obtained using QTOF Agilent 6545 mass
453 spectrometer coupled to an Agilent Infinity 1260 LC system running a Jet Stream ESI source. The
454 samples (in acetonitrile) were directly infused using acetonitrile as cosolvent. All measurements
455 were performed at 100 °C dry gas and source temperature and the machine was calibrated in the
456 mass range of 100-3200 Da prior the measurements. Data analysis was performed with mmass.

457 *Rheology*

458 Frequency sweep, amplitude sweep, and stress-relaxation experiments were performed using a TA
459 Instruments Discovery HR-2 rheometer with an 8 mm parallel plate geometry unless otherwise
460 noted. Samples were cased using a Teflon mold to a thickness of ~1.5 mm as squares and trimmed
461 into circular pucks with soft plastic tools once loaded on the rheometer. All samples were covered
462 in mineral oil to prevent solvent evaporation and de-swelling. Amplitude sweep experiments were
463 performed on all gels to ensure that the subsequent strain percent of frequency sweep and stress-
464 relaxation experiments were within the linear viscoelastic regime. All frequency sweep and stress-
465 relaxation experiments were performed at 1% strain. Variable temperature stress-relaxation

466 experiments were performed in 5 °C intervals from 25°C to 50°C and gels were allowed to
467 equilibrate at each temperature for 15 minutes prior to the experiment.

468 *General MOC Synthesis*

469 **L1** or **L2** (0.01 mmol, 1 equiv.) was added to a 4mL vial and dissolved in 300 μ L of DMSO-*d*₆.
470 Pd(MeCN)₄(OTf)₂ (6.27 mg, 0.005 mmol, 1.05 equiv.) was dissolved in 200 μ L of DMSO-*d*₆. The
471 palladium solution was added to the polymer ligand solution and an immediate color change from
472 red-orange to pale yellow was observed, suggesting formation of palladium-pyridine coordination
473 structures. The solution was characterized immediately by ¹H NMR. To prepare more concentrated
474 MOC solutions, <300 μ L of solvent was used to disperse **L1** or **L2**, to which a palladium stock
475 solution was added. Because the MOCs were generally more soluble in DMSO than the small-
476 molecule ligands, these mixtures would drive ligand solubilization and subsequent MOC
477 formation after stirring for ~15 minutes at room temperature.

478 *Guest Binding Studies*

479 The measurements of guest association constant with **MOC** were performed by titrating various
480 equivalents of guest compound into a solution of **MOC**, maintaining a constant concentration of
481 **MOC** throughout (5mM). All studies were done in DMSO, the same solvent used for polyMOC
482 gel fabrication. Both **NO**₃ and **HSO**₄ as guests in **MOC** and **N-MOC** displayed rapid exchange
483 with the **MOC** cavity on the NMR timescale. For these guests, the resonance shifts of one or
484 multiple protons belonging to the MOC were monitored as increasing equivalents of guest was
485 added. These resonance shifts were fit to a 1:1 binding model using a freely available tool BindFit
486 from Supramolecular.org (and P. Thordarson, *Chem. Soc. Rev.*, **2011**, *40*, 1305-1323) to calculate
487 the appropriate association constant, K_a. Because **DAQ** exhibited slow exchange with the MOC
488 cavity on the NMR timescale, a modified approach was taken. Upon addition of **DAQ**, resonances
489 associated with the guest-bound MOC, **DAQ**•**MOC**, appeared. The intensity of these resonances
490 relative to those of the un-bound MOC, as determined by integration, could be fit to a 1:1 binding
491 model to calculate the association constant.

492
493 *General PolyMOC Gel Fabrication Procedure*

494 **PL** (55 mg, 0.01 mmol, 1 equiv.) was added to a 4mL vial and dissolved in 300 μ L of DMSO-*d*₆.
495 Pd(MeCN)₄(OTf)₂ (6.27 mg, 0.0105 mmol, 1.05 equiv.) was dissolved in 200 μ L of DMSO-*d*₆. The

496 palladium solution was added to the polymer ligand solution in four portions of 50 μ L, with the
497 gel being thoroughly mixed with a spatula and vortex between each addition. After the first two
498 additions a tacky gel was observed to form. The gel was then mixed for a final time and broken
499 into small pieces before being cast between two Teflon sheets (2 cm x 2 cm x 1.5 mm) lined with
500 two pieces of polyethylene for a release liner. The gel casting was annealed at 60°C for 1 hour to
501 yield a nearly colorless, transparent, homogeneous gel (**poly(OTf \subset MOC)**).

502

503 **Data Availability**

504 The data generated or analyzed during this study are available in the article, supplementary
505 information files, or the accompanying source data file. The X-ray crystallography structures
506 reported in this study have been deposited at the Cambridge Crystallographic Data Centre
507 (CCDC), under deposition numbers 2338506, 2338512, 2338513, and 2338517. These data can be
508 obtained free of charge from The Cambridge Crystallographic Data Centre via
509 www.ccdc.cam.ac.uk/data_request/cif. All data are available from the corresponding author upon
510 request.

511

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651

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659 660 **Author Contributions Statement**

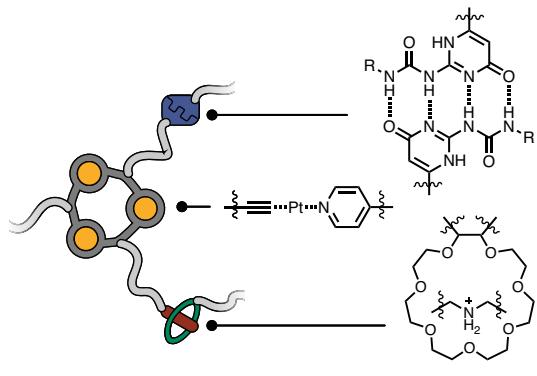
661 D.J.L., C.M.B., N.J.O., and J.A.J. conceptualized the research. D.J.L., C.M.B., E.O.B., N.J.O.,
662 Y.S.A., and J.Z. synthesized compounds. D.J.L., C.M.B., E.O.B., and N.J.O. characterized
663 materials and performed data analysis. I.K. and H.J.K. performed and advised computational
664 studies. D.J.L. and C.M.B. performed crystallographic studies. D.J.L. prepared figures. D.J.L. and
665 J.A.J. wrote the manuscript with input from all co-authors.

666 667 **Competing Interests Statement**

668 The authors declare no competing interests.
669

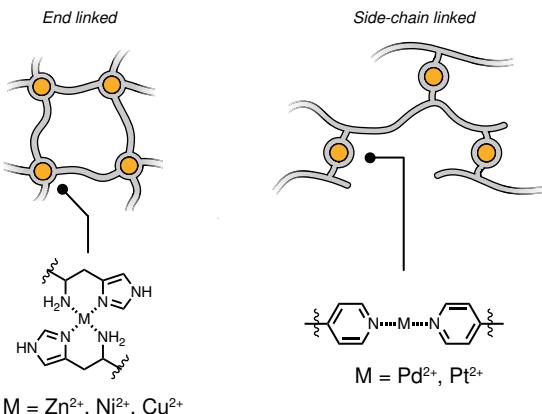
A Previous Work: Discrete Combinations of Multiple Non-Covalent Interactions

Host-Guest, Metal-Ligand Coordination, and Hydrogen Bonding Interactions (Stang)



- + Broadened functionality and stimuli-responsiveness
- Traditional stoichiometric assembly limitations
- Supramolecular interactions are independent

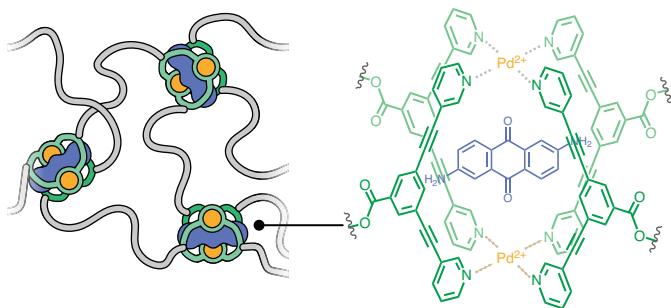
Multiple Strengths of an Interaction Class
e.g. Metal-Ligand Coordination (Holten-Andersen, Craig)



$M = \text{Zn}^{2+}, \text{Ni}^{2+}, \text{Cu}^{2+}$

- + Tunable dynamic properties
- Traditional stoichiometric assembly limitations
- Maintained stimuli-responsive properties

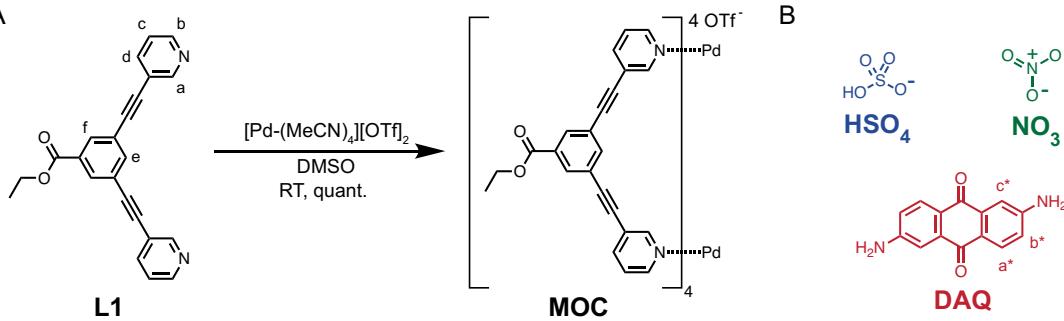
B This Work: Nested Metal-Ligand Coordination and Host-Guest Binding



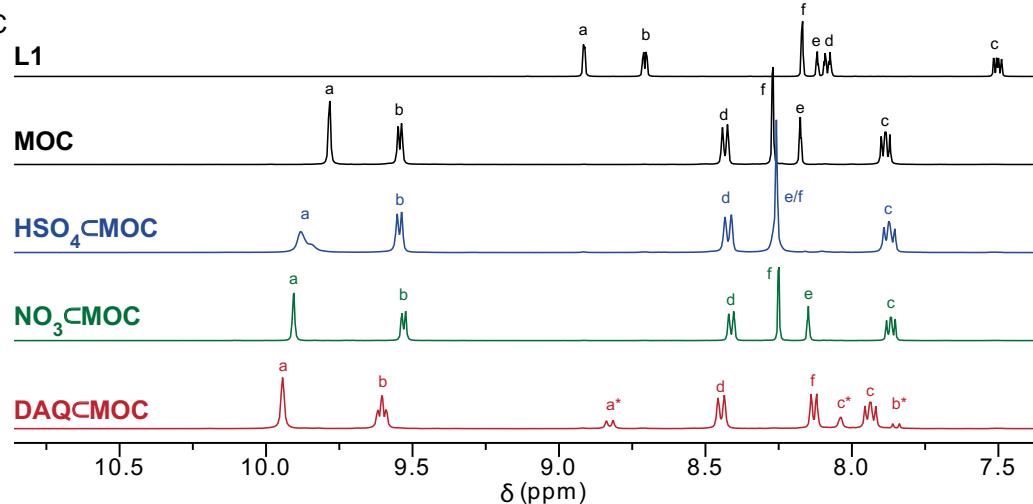
Nested Configuration Enables:

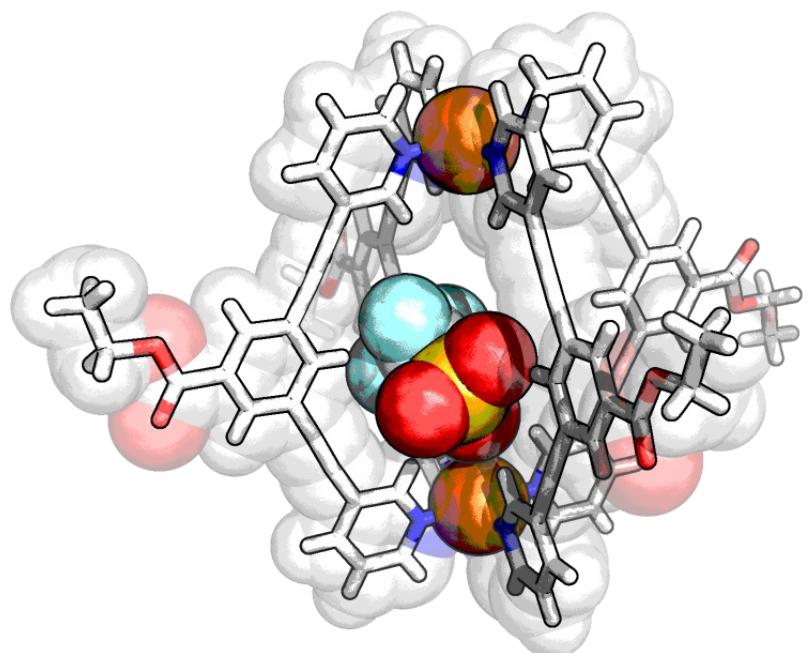
- + Modulation of global dynamics with guest strength
- + Expanded stoichiometric assembly window
- + Small-molecule triggered sol-gel transitions
- + Guest binding strength selective sol-gel transitions
- + Self-sorting driven sol-gel transition networks

A

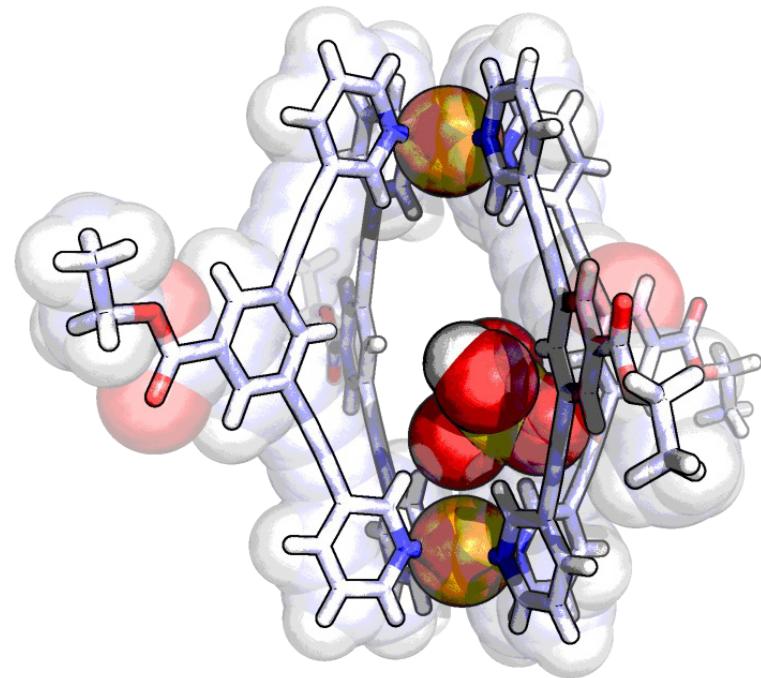


C

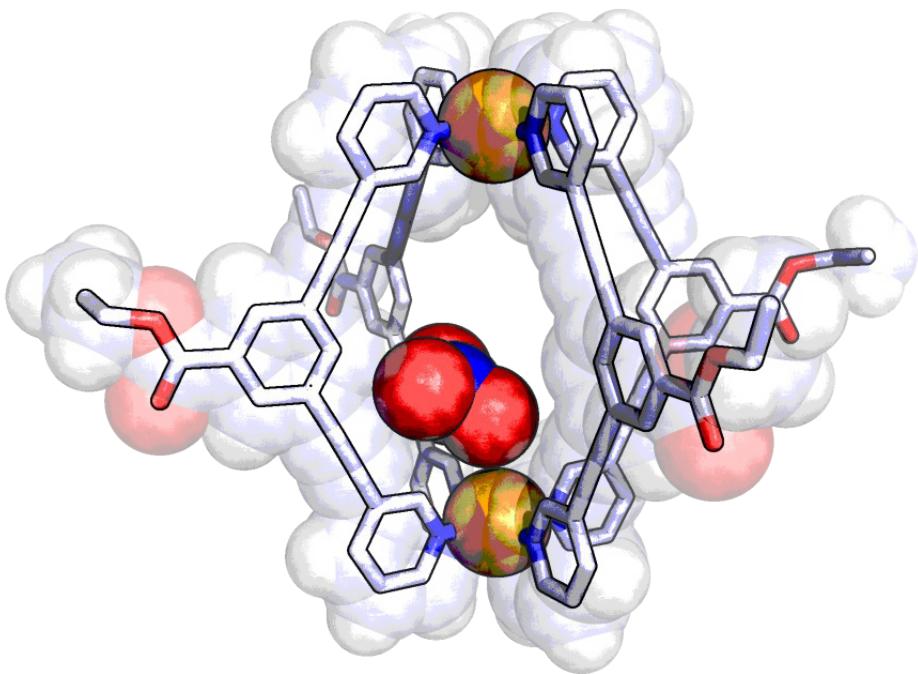




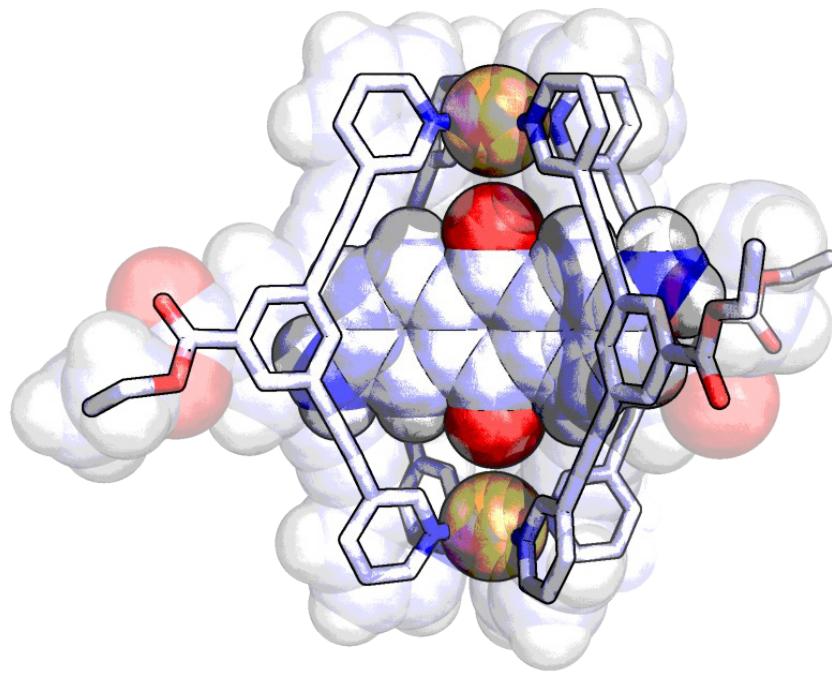
MOC



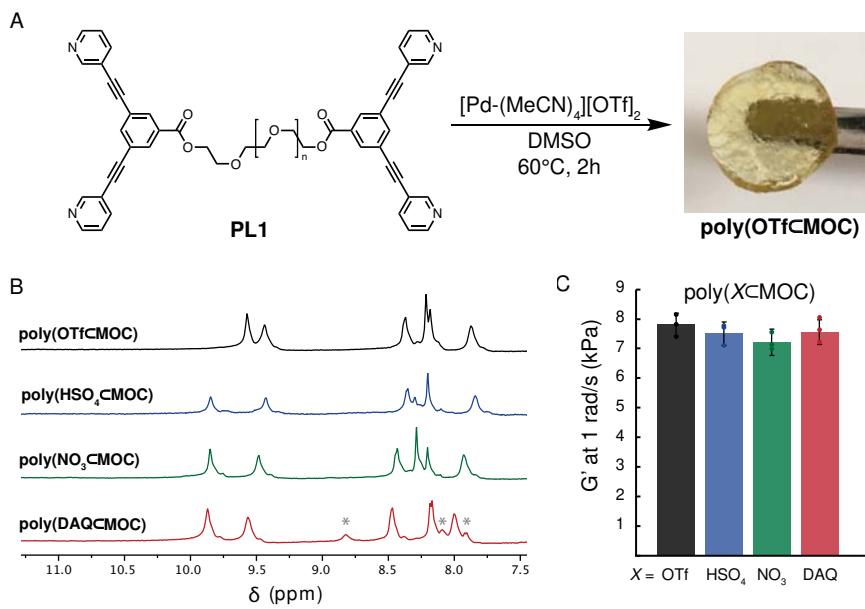
$\text{HSO}_4^- \subset \text{MOC}$

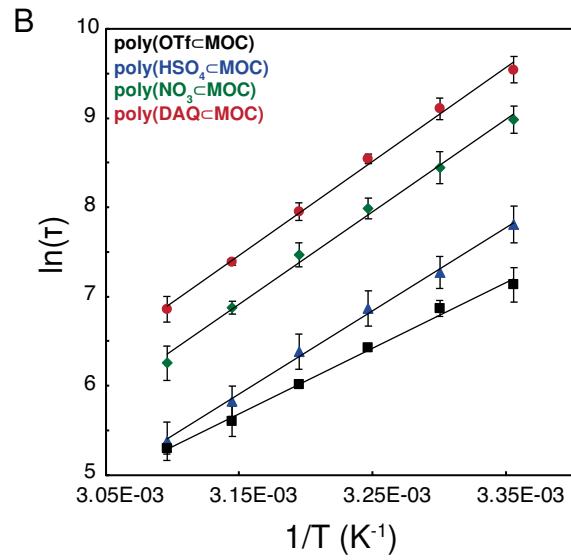
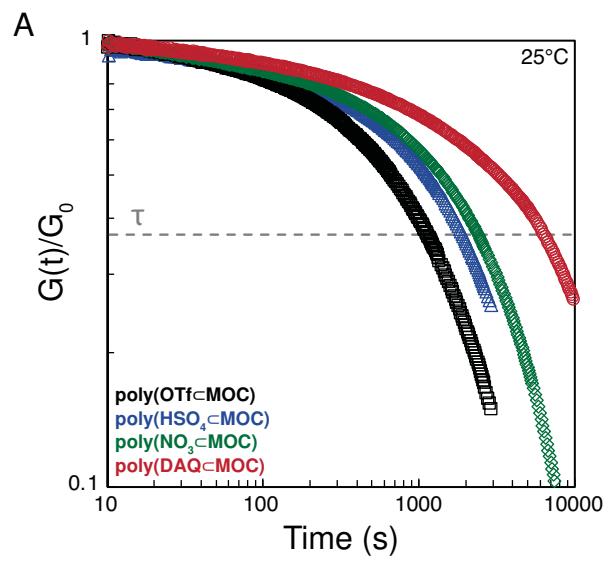


$\text{NO}_3^- \subset \text{MOC}$

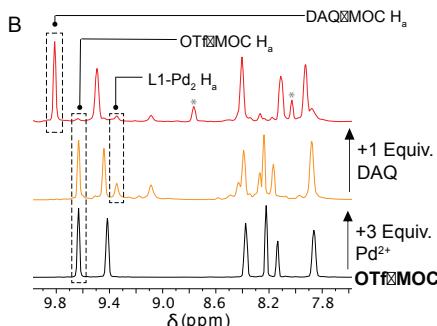
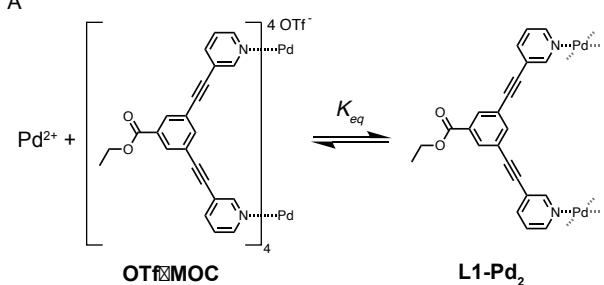


$\text{DAQ}^- \subset \text{MOC}$

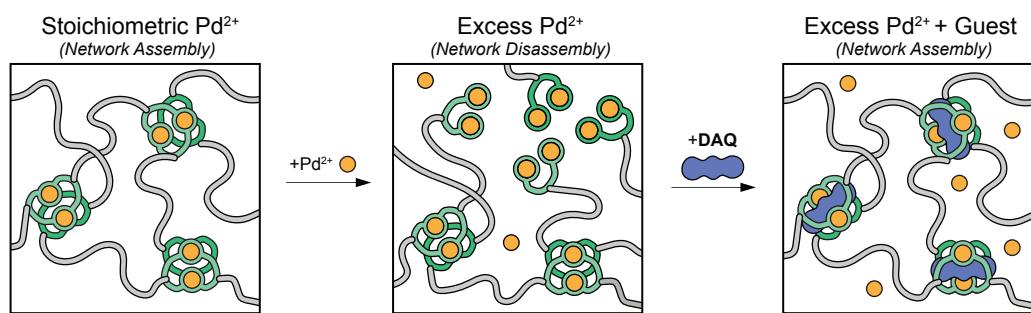




A



C



D



E

