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## Controlling the thermally-driven crystallization of DNA-coated nanoparticles with formamide†

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DNA-coated nanoparticles, also known as programmable atom equivalents (PAEs), facilitate the construction of materials with nanoscopic precision. Thermal annealing plays a pivotal role by controlling DNA hybridization kinetics and thermodynamics, which ensures the formation of intended structures. While various design handles such as particle size, DNA design, and salt concentration influence the stability of the DNA duplexes linking PAEs in a lattice, their influence on the system's melting temperature ( $T_m$ ) often follows complicated trends that make rational tuning of self-assembly challenging. In this work, the denaturant formamide is used to precisely tune the thermal response of PAEs. Our results reveal a clear and predictable trend in the PAEs' response to formamide, enabling rational control over the  $T_m$  of a diverse set of PAE systems. Unlike adjustments made through alterations to PAE design or solution parameters such as ionic strength, formamide achieves its temperature shift without impacting the kinetics of assembly. As a result, PAEs can be rapidly crystallized at ambient temperatures, producing superlattices with similar quality to PAE crystals assembled through standard protocols that use higher temperatures. This study therefore positions formamide as a useful tool for enhancing the synthesis of complex nanostructures under mild conditions.

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## Introduction

Colloidal self-assembly is a powerful method for creating materials with intricate nanoscale features, which can exhibit unique properties for applications in structural materials, electronics, and photonics.<sup>1–11</sup> The primary advantage of self-assembly is its ability to produce these features through a rational, programmable approach using building block design and processing conditions to control the formation of different structural motifs.<sup>12–18</sup> In particular, DNA hybridization provides a robust method for customizing interparticle interactions for a wide range of particle sizes, shapes, and compositions.<sup>19–24</sup> The highly ordered superlattices generated by DNA-directed nanoparticle crystallization have resulted in the moniker of “programmable atom equivalents” (PAEs) to describe these DNA-grafted colloids.<sup>25–27</sup>

The thermal response of PAEs is crucial for controlling particle interactions, thereby enabling the assembly of diverse crystalline structures through thermal annealing.<sup>28–30</sup> While a variety of design handles (*e.g.* solution ionic strength, DNA design, particle size, particle shape) have been studied as a means to tailor PAEs' thermal response,<sup>31–33</sup> modulation of

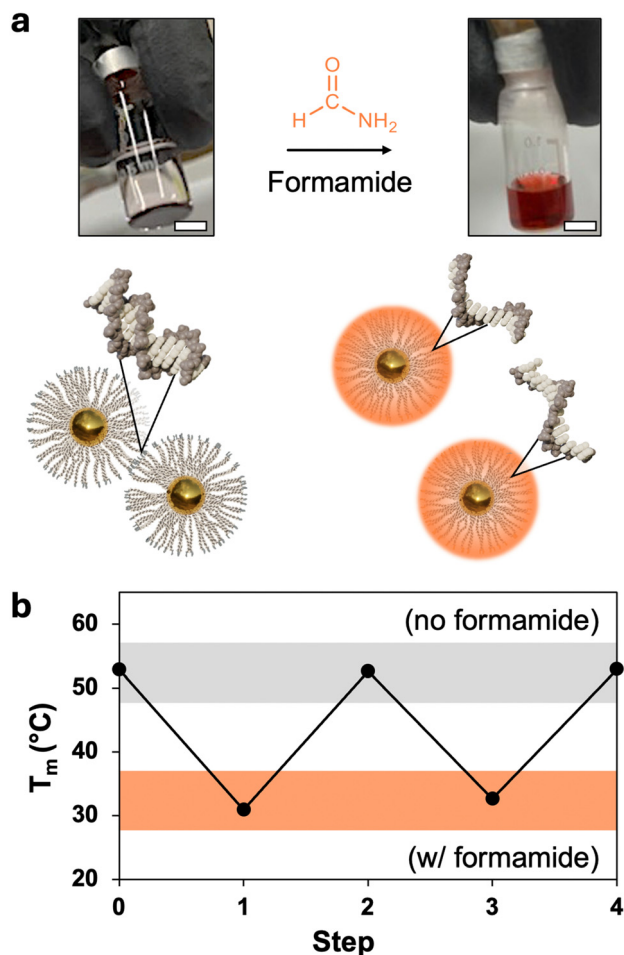
these parameters often leads to complex assembly phenomena that can be difficult to rationally and finely control in laboratory experiments. For example, ionic strength affects the system's melting temperature ( $T_m$ ),<sup>34</sup> but also changes the interaction pair potential,<sup>32</sup> as well as the crystallization kinetics and final crystal quality.<sup>35</sup> Particle size/curvature and DNA grafting density also affect  $T_m$ , but these cannot be easily manipulated after PAEs are synthesized.<sup>36</sup> Thus, it would be beneficial to have a singular design handle that could be applied post-synthetically to modulate DNA interactions and thermal response in a straightforward manner.

In this work, we examine how formamide, a known denaturant for DNA, can be used to predictably control the thermodynamics and kinetics of PAE assembly. Formamide disrupts DNA's hydration shell, interfering with DNA duplexation, thereby lowering PAE dissociation temperature.<sup>37</sup> Importantly, increasing formamide concentration linearly lowers the temperature at which DNA strands dissociate, regardless of DNA nucleobase sequence. Thus, formamide could be an excellent addition to the toolbox of PAE self-assembly as a means of more finely controlling assembly behavior. However, dense brushes of DNA grafted to a nanoparticle surface can alter the local ion distribution around DNA duplexes and also the local concentration of DNA duplexes between particles.<sup>34</sup> These structural differences make it unclear if formamide will have the same reliability in altering PAEs'  $T_m$ . By investigating how varying concentrations of formamide affect the melting transitions of

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**Fig. 1** Formamide serves as a denaturant for PAEs. (a) PAEs are held together at room temperature via hybridized DNA strands, leading particles to assemble and form a darkly colored sediment. Formamide dehybridizes those bonds, leading the particles to dissociate and form a stable suspension, exhibited by the characteristic red color of single gold nanoparticles. Scale bars 1 cm. (b)  $T_m$  of PAE assemblies drops with the addition of formamide but removing formamide allows  $T_m$  to return to its original value. This cycle can be repeated, and indicates that formamide is non-destructive to the DNA-brush on the nanoparticles.

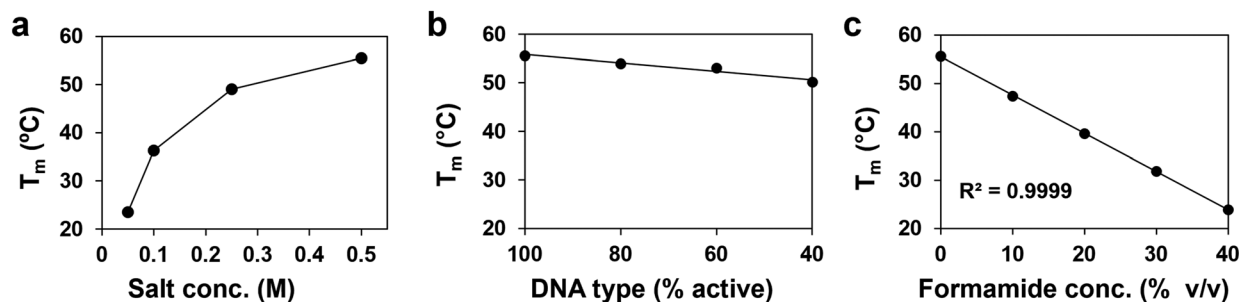
PAEs with different particle and DNA designs, we demonstrate that it provides significantly enhanced level of control over the

thermal response of PAEs compared to existing methods. This enhanced control then enables the formation of high-quality crystal structures at significantly lower temperatures than are typically required for PAE crystallization.

## Results and discussion

Formamide is a denaturant that is widely used in molecular biology to reduce the melting temperature of DNA (Fig. 1a).<sup>38</sup> Its mild yet effective action allows for the controlled separation of genetic material while preserving the integrity of nucleic acids,<sup>39</sup> making it a useful tool for applications such as Southern blots, polyacrylamide gel electrophoresis (PAGE), and polymerase chain reactions (PCR).<sup>40,41</sup> It is also reliable enough in its effect on DNA hybridization to be used as a substitute for thermal annealing in the complex folding process of DNA-origami nanostructures.<sup>42,43</sup> This work demonstrates that formamide similarly serves as a denaturant for PAEs, which is an effective means to increase our control over PAE self-assembly.

Adding formamide to PAE assemblies comprising gold nanoparticle cores bound through hybridized DNA results in their rapid dissolution into a homogeneous nanoparticle suspension at temperatures significantly lower than their native  $T_m$  (Fig. 1a). Crucially, the nucleobases are kept intact during denaturation, allowing the nanoparticles to return to their original binding capacity upon formamide removal (Fig. 1b). While other processing conditions can similarly be used to reversibly affect DNA binding to disperse or assemble PAEs, formamide potentially offers much finer control over self-assembly than existing methods. In particular, formamide is noteworthy because DNA duplex  $T_m$  decreases linearly with increasing concentrations of formamide, and this behavior is consistent across various DNA designs, with trends ranging from 0.6 to 0.8 °C per volume percentage of formamide (% v/v).<sup>37</sup> In comparison, the impact of variables such as ionic strength on DNA can have more complicated, non-linear effects. Existing predictive models can provide general information to tune  $T_m$  using these design handles,<sup>44</sup> but precise modulation remains challenging. These complications are further exacerbated with PAE assembly due to the larger



**Fig. 2** Methods to tune PAE  $T_m$ . (a)  $T_m$  is plotted over the concentration of sodium chloride, displaying a nonlinear trend. (b)  $T_m$  is plotted over variations in the linker loadings. With increasing concentrations of non-participatory dummy strands, the percent of active binding DNA on the PAEs drops, leading to a lower  $T_m$ . While the trend is linear, the range is narrow. (c)  $T_m$  is plotted over volume fraction of formamide. The trend is linear, and the range of temperatures that can be selected is wide.



number of design variables and the high local concentration of polyanionic DNA strands around the nanoparticle core.<sup>32</sup> Adopting this ability to tune thermal response is essential for PAEs, as there is currently no method to precisely set the  $T_m$  of a suspension without extensive empirical testing of conditions. For example, like free DNA duplexes, PAE  $T_m$ s respond non-linearly to solution ionic strength, and trends can vary significantly between systems (Fig. 2a).<sup>34</sup> The grafting density of DNA strands on each PAE can also be used to alter  $T_m$ . In practice, PAE assembly is typically induced by hybridizing “linker” strands to the DNA that is covalently bonded to the nanoparticle core (Fig. S1, ESI<sup>†</sup>). Thus, the number of DNA interactions between any given pair of bonded PAEs can be controlled by altering the ratio of linker strands to PAEs. PAEs can also be co-loaded with both linker strands and non-participatory “dummy” strands, which lack the unpaired sticky ends necessary to form DNA duplexes between PAEs. Reduction of linker density also lowers PAEs'  $T_m$ , but offers only a narrow window of control (Fig. 2b), and the quality of crystals can drop significantly with low loading of active linkers. In contrast to both of these design handles (ionic strength and linker

density), formamide allows  $T_m$  to be tuned over a wide window ( $>30$  °C), and follows a linear and predictable trend (Fig. 2c). In principle, this linearity allows any new  $T_m$  to be selected without meticulously optimizing assembly conditions, if the original  $T_m$  is known (0% v/v formamide).

While this initial result is promising for the use of formamide as a means to modulate PAE assembly thermodynamics, it is important to note that PAEs exhibit unique multivalency-driven thermal behaviors that are affected by the number and arrangement of the numerous oligonucleotides grafted to the nanoparticles' surfaces.<sup>34</sup> Such effects could alter the slope or disrupt the linearity of the PAEs' formamide response. Thus, more quantitative examination of formamide's effects on various systems must also be performed in order for this approach to yield a useful tool for modulating PAE assembly.

Experiments were conducted to investigate multivalency effects by varying core sizes and grafting density of PAEs. Two particle sizes were prepared (15 and 35 nm) (Fig. S2, ESI<sup>†</sup>), containing approximately 200 and 900 DNA linkers at maximum grafting density, respectively.<sup>45,46</sup> The grafting density of PAEs was then modified independently of particle size by co-loading PAEs with controlled ratios of active linkers with dummy linkers, ultimately providing a wide variety of multivalent architectures to explore. Notably, with formamide, all systems demonstrated a similar reliable linear trend in  $T_m$  reduction across a wide temperature range (Fig. 3a). Additionally, the slope of these plots was measured to be very similar across all systems, showing a decrease of approximately  $0.8 \pm 0.02$  °C per volume percent of formamide (Fig. 3b).

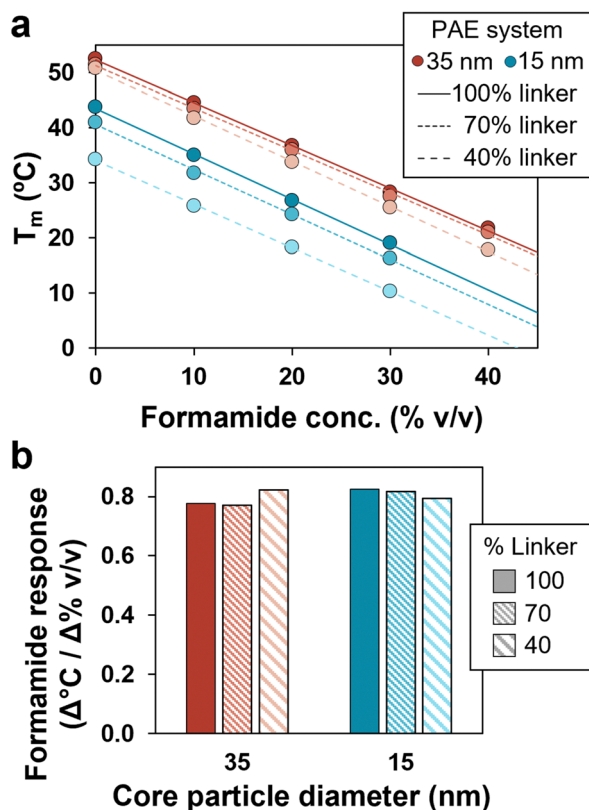


Fig. 3 Formamide's effect on diverse PAE systems. (a)  $T_m$  is plotted over the volume fraction of formamide for a variety of PAEs. Traces in red comprise PAEs with gold cores measuring 35 nm in diameter, while traces in blue comprise 15 nm gold cores. Additionally, the percentage of active linker strands is varied from 100% to 40%, represented by dashed lines. All traces display a linear trend. (b) The response to formamide (change in  $T_m$  over the change in volume fraction of formamide) is given by the slope of the traces in “a” and is plotted for each PAE system. Despite differences in multivalency for each system, the response to formamide is similar.

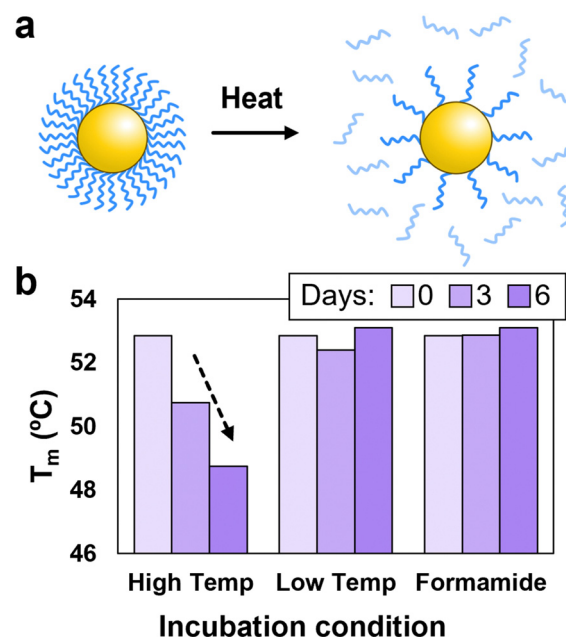


Fig. 4 PAE viability in different environments. (a) Schematic of DNA anchors desorbing from gold cores during heating. (b)  $T_m$  is plotted for PAEs incubated in different conditions over the course of 6 days. While PAEs held at elevated temperatures (60 °C) display a significant drop in  $T_m$  in this time, those held at room temperature with or without formamide remain constant.

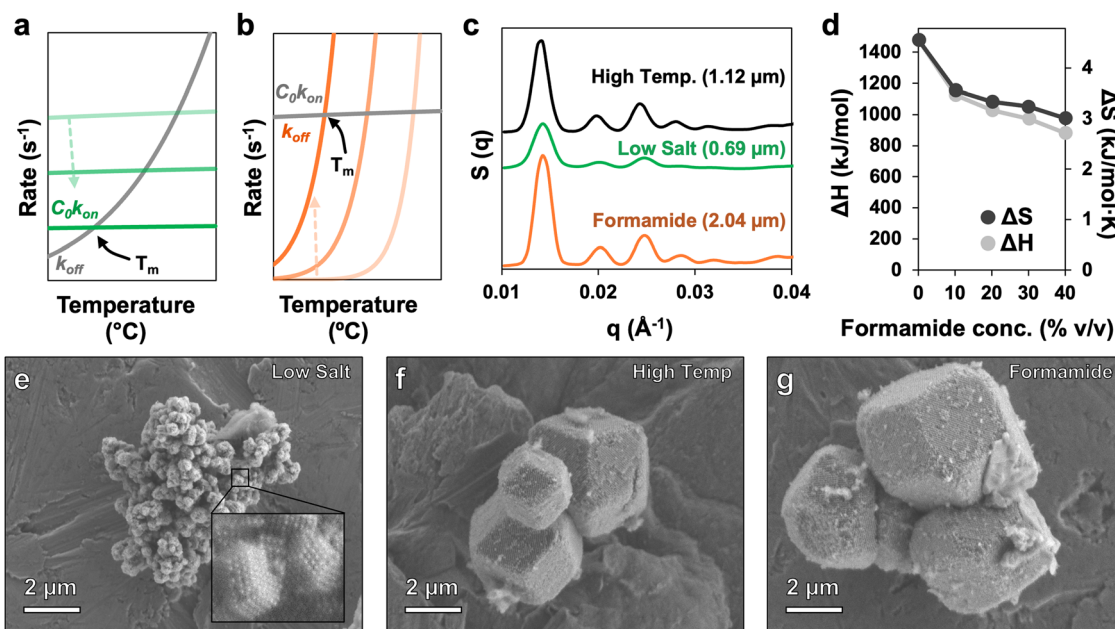


This formamide response is consistent with prior examinations of formamide's effects on free DNA strands.<sup>37</sup> These trends were also observed for different DNA designs, including DNA linkers of different lengths, and both binary and unary PAE assemblies (Fig. S3, ESI†). These results confirm that the formamide response is consistent across multiple PAE designs, emphasizing its utility as a tool for tuning PAE self-assembly.

To further demonstrate the advantages of using formamide to modulate PAE assembly, it is important to consider the potential benefits associated with conducting assembly at lower temperatures. Typically, to grow large single crystals, PAEs must undergo slow cooling to prevent excessive nucleation.<sup>28,31</sup> This process often requires maintaining the particles at elevated temperatures for extended periods,<sup>47</sup> which can result in the detachment of DNA grafts thereby degrading the DNA brush and diminishing assembly quality (Fig. 4a).<sup>45,48</sup> An indication of this degradation is a drop in  $T_m$  after the PAEs are held at elevated temperatures for significant periods of time, which is attributed to a reduced number of linkers binding the particles together. For example, when PAEs were incubated at 60 °C, a 4 °C decrease in  $T_m$  was observed after six days (Fig. 4b). In contrast, PAEs maintained at room temperature display a consistent  $T_m$  over the same time period. Similarly, PAEs incubated in 25% v/v formamide (at room temperature) show no decrease in  $T_m$ , indicating that formamide does not degrade DNA over extended periods. Therefore,

formamide could alleviate DNA desorption issues that would be expected during standard slow cooling experiments to produce single crystals, as the addition of formamide would permit assembly to be conducted at temperatures where DNA grafts remain stable. However, it is important to note that formamide can oxidize into formic acid which can cleave nucleobases,<sup>49</sup> thus requiring the use of fresh formamide stored in at low temperatures (additional details in ESI,† Section S1.5).

As noted above, solution ionic strength would be expected to induce many of the same effects observed from the addition of formamide, albeit with a non-linear correlation between solution salt concentration and  $T_m$ . Nevertheless, even if the correlation between solution salt concentration and PAE  $T_m$  were meticulously characterized, formamide still provides a major advantage because it only alters DNA hybridization thermodynamics. Conversely, lowering PAE  $T_m$  by altering solution ionic strength drastically changes both assembly thermodynamics and assembly kinetics, which has adverse effects on the quality of the resulting PAE crystals. The  $T_m$  of a given PAE system is the temperature at which the rate of DNA association ( $C_0k_{on}$ ) and dissociation ( $k_{off}$ ) are equal, where  $C_0$  is the initial concentration of unhybridized DNA, and  $k_{on}$  and  $k_{off}$  are the rate constants of association and dissociation, respectively.<sup>35,36</sup> One of the effects of low ionic strength on DNA is to decrease the rate of  $k_{on}$ ,<sup>35,50</sup> which slows down



**Fig. 5** Utility of formamide for PAE assembly. (a) Heuristic image showing the trends of  $C_0k_{on}$  (green) and  $k_{off}$  (grey) at various salt concentrations (light green: high salt, dark green: low salt).  $T_m$  is represented by the intersection of  $C_0k_{on}$  and  $k_{off}$  for each trace; decreasing  $C_0k_{on}$  with lower salt concentrations leads to a drop in  $T_m$ . At lower  $T_m$ s, rates for  $k_{off}$  and  $C_0k_{on}$  drop, leading to poor crystallization kinetics.<sup>35,36,50</sup> (b) Heuristic image showing the trends of  $C_0k_{on}$  (grey) and  $k_{off}$  (orange) at various formamide concentrations (light orange: low formamide, dark orange: high formamide). Increasing  $k_{off}$  with higher formamide concentrations leads to a drop in  $T_m$ . At all  $T_m$ s, rates for  $k_{off}$  and  $C_0k_{on}$  remain high, allowing for rapid reconfiguration and large crystal growth.<sup>51</sup> (c) 1D SAXS patterns for traditional high temperature annealing of PAEs (black;  $T_m = 48$  °C), low temperature annealing using a low salt concentration to lower  $T_m$  (green;  $T_m = 25$  °C), and low temperature annealing using formamide to lower  $T_m$  (orange;  $T_m = 25$  °C). All samples cooled from 60 °C to 10 °C at a rate of 0.1 °C min<sup>-1</sup>. Included is the average crystallite size determined via Williamson–Hall analysis. (d)  $\Delta H$  and  $\Delta S$  for PAE binding at varied formamide concentration determined via Van't Hoff analysis. A drop in  $\Delta H$  indicates broadening of the melt curve at higher formamide concentrations, which can help facilitate the growth of larger crystals. (e)–(g) SEM images of crystals formed via low temperature annealing with low salt, traditional high temperature annealing, and low temperature annealing using formamide, respectively.



particle reconfiguration around  $T_m$  (Fig. 5a). Consequently, even with careful control of  $T_m$  by altering solution salt concentration, assembling high-quality crystals at lower temperatures remains challenging due to sluggish particle kinetics. In contrast, evidence suggests that formamide's denaturation effects are due mainly to the destabilization the double helix,<sup>37,38</sup> that is, increasing  $k_{\text{off}}$  without significantly affecting  $k_{\text{on}}$ .<sup>51</sup> Therefore, the kinetics of assembly are expected to remain high across all formamide concentrations, even at lower  $T_m$ s (Fig. 5b).

To further investigate the effects of kinetics, annealing experiments were conducted to assess the quality of crystals formed under different processing conditions, including traditional high temperature annealing and lower temperature annealing where  $T_m$  adjustments were made using either ionic strength or formamide. SAXS data confirm that while the control "high temperature" and "formamide" samples have a high degree of crystallinity, the "low salt" sample displays only a few broad peaks indicating a lower quality of crystallization (Fig. 5c and Fig. S4, S5, ESI†). Williamson–Hall analysis of SAXS patterns also indicate that assembling crystals under low salt conditions produces smaller crystallites, while assembly in the presence of formamide produces crystallites that actually exceed the size of crystals assembled using standard high-temperature methods. One possibility for the relative increase in crystal size under formamide conditions is broadening of the melt curve, which provides particles more time to nucleate and assemble into fewer, larger crystals. Melt broadening, characterized by a decrease in  $\Delta H$ , has been previously observed in formamide-treated DNA systems,<sup>37</sup> while a similar trend emerged for PAEs, as determined by Van't Hoff analysis of melt curves for different formamide concentrations (Fig. 5d and Fig. S6, ESI†). High resolution SEM images indeed show that crystals formed in low salt conditions are substantially smaller compared to the other annealing methods (Fig. 5e). Additionally, both the standard high temperature method and assembly with formamide-tuned  $T_m$  successfully produced crystals with well-defined rhombic dodecahedral habits, confirming their higher quality indicated by SAXS (Fig. 5f–g). While the low salt sample clearly does display some order, and likely could be improved in quality by increasing the duration of cooling, formamide provides a far more robust and simple means to achieve crystallization at ambient conditions, without the need to extensively optimize PAE assembly protocols.

## Conclusions

The use of formamide to tune PAE  $T_m$  represents a significant advance for the field of self-assembly. While numerous colloidal interactions can be used to modulate DNA-driven assembly thermodynamics, formamide offers a rare design handle that allows precise and rational control. The predictability and consistency of  $T_m$  reduction with formamide, as opposed to the non-linear effects observed with other variables such as ionic strength, present a clear advantage in designing and

implementing controlled self-assembly processes. This research enhances our understanding of PAE behavior under varying thermal conditions and offers a practical method to achieve rapid crystallization at ambient temperatures. Such conditions not only prolong the lifespan of traditional PAE systems utilizing gold-thiol bonds but could also be crucial for utilizing PAEs with DNA ligands bound through weaker chemistries that are even more sensitive to elevated temperatures.<sup>52,53</sup> Additionally, this methodology may be similarly applicable for controlling the interactions of DNA-coated colloidal particles on the micron scale.<sup>22</sup> High-temperatures currently pose challenges for detailed analysis using higher-magnification oil immersion optics, which act as heat sinks. In contrast, room temperature assembly could facilitate the use of these lenses for enhanced imaging at higher magnifications. Precise  $T_m$  control also enables applications for sensing, wherein functional nanostructures respond programmatically to thermal fluctuations near biologically relevant temperatures.

## Author contributions

All authors designed experiments and analyzed data. TH and SW synthesized materials and collected data. All authors wrote and edited the manuscript.

## Data availability

The data supporting this article have been included as part of the ESI.†

## Conflicts of interest

There are no conflicts of interest to declare.

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## References

- 1 F. Li, S. F. Liu, W. Liu, Z. W. Hou, J. Jiang and Z. Fu, *et al.*, 3D printing of inorganic nanomaterials by photochemically bonding colloidal nanocrystals, *Science*, 2023, **381**(6665), 1468–1474.
- 2 T. Hueckel, X. Luo, O. F. Aly and R. J. Macfarlane, Nanoparticle Brushes: Macromolecular Ligands for Materials Synthesis, *Acc. Chem. Res.*, 2023, **56**(14), 1931–1941.
- 3 T. Hueckel, G. M. Hocky and S. Sacanna, Total synthesis of colloidal matter, *Nat. Rev. Mater.*, 2021, **6**(11), 1053–1069.



- 4 Z. Cai, Z. Li, S. Ravaine, M. He, Y. Song and Y. Yin, *et al.*, From colloidal particles to photonic crystals: advances in self-assembly and their emerging applications, *Chem. Soc. Rev.*, 2021, **50**(10), 5898–5951.
- 5 S. Dhulipala, D. W. Yee, Z. Zhou, R. Sun, J. E. Andrade and R. J. Macfarlane, *et al.*, Tunable Mechanical Response of Self-Assembled Nanoparticle Superlattices, *Nano Lett.*, 2023, **23**(11), 5155–5163.
- 6 R. L. Li, C. J. Thrasher, T. Hueckel and R. J. Macfarlane, Hierarchically Structured Nanocomposites *via* a “Systems Materials Science” Approach, *Acc. Mater. Res.*, 2022, **3**(12), 1248–1259.
- 7 L. Z. Zornberg, D. J. Lewis, A. Mertiri, T. Hueckel, D. J. D. Carter and R. J. Macfarlane, Self-Assembling Systems for Optical Out-of-Plane Coupling Devices, *ACS Nano*, 2023, **17**(4), 3394–3400.
- 8 D. J. Lewis, D. J. D. Carter and R. J. Macfarlane, Using DNA to Control the Mechanical Response of Nanoparticle Superlattices, *J. Am. Chem. Soc.*, 2020, **142**(45), 19181–19188.
- 9 M. A. Boles, M. Engel and D. V. Talapin, Self-Assembly of Colloidal Nanocrystals: From Intricate Structures to Functional Materials, *Chem. Rev.*, 2016, **116**(18), 11220–11289.
- 10 S. C. Glotzer and M. J. Solomon, Anisotropy of building blocks and their assembly into complex structures, *Nat. Mater.*, 2007, **6**(8), 557–562.
- 11 Z. Li, Q. Fan and Y. Yin, Colloidal Self-Assembly Approaches to Smart Nanostructured Materials, *Chem. Rev.*, 2022, **122**(5), 4976–5067.
- 12 T. Hueckel, G. M. Hocky, J. Palacci and S. Sacanna, Ionic solids from common colloids, *Nature*, 2020, **580**(7804), 487–490.
- 13 J. Opdam, R. Tuinier, T. J. Hueckel, T. Snoeren and S. Sacanna, Selective colloidal bonds via polymer-mediated interactions, *Soft Matter*, 2020, **16**(32), 7438–7446.
- 14 R. Klajn, K. J. M. Bishop and B. A. Grzybowski, Light-controlled self-assembly of reversible and irreversible nanoparticle suprastructures, *Proc. Natl. Acad. Sci. U. S. A.*, 2007, **104**(25), 10305–10309.
- 15 A. M. Kalsin, M. Fialkowski, M. Paszewski, S. K. Smoukov, K. J. M. Bishop and B. A. Grzybowski, Electrostatic Self-Assembly of Binary Nanoparticle Crystals with a Diamond-Like Lattice, *Science*, 2006, **312**(5772), 420–424.
- 16 P. F. Damasceno, M. Engel and S. C. Glotzer, Predictive Self-Assembly of Polyhedra into Complex Structures, *Science*, 2012, **337**(6093), 453–457.
- 17 P. J. M. Swinkels, Z. Gong, S. Sacanna, E. G. Noya and P. Schall, Visualizing defect dynamics by assembling the colloidal graphene lattice, *Nat. Commun.*, 2023, **14**(1), 1524.
- 18 C. J. Thrasher, F. Jia, D. W. Yee, J. M. Kubiak, Y. Wang and M. S. Lee, *et al.*, Rationally Designing the Supramolecular Interfaces of Nanoparticle Superlattices with Multivalent Polymers, *J. Am. Chem. Soc.*, 2024, **146**(16), 11532–11541.
- 19 R. J. Macfarlane, B. Lee, M. R. Jones, N. Harris, G. C. Schatz and C. A. Mirkin, Nanoparticle Superlattice Engineering with DNA, *Science*, 2011, **334**(6053), 204–208.
- 20 Y. Zhang, F. Lu, K. G. Yager, D. van der Lelie and O. Gang, A general strategy for the DNA-mediated self-assembly of functional nanoparticles into heterogeneous systems, *Nat. Nanotech.*, 2013, **8**(11), 865–872.
- 21 W. B. Rogers, W. M. Shih and V. N. Manoharan, Using DNA to program the self-assembly of colloidal nanoparticles and microparticles, *Nat. Rev. Mater.*, 2016, **1**(3), 1–14.
- 22 Y. Wang, Y. Wang, X. Zheng, É. Ducrot, J. S. Yodh and M. Weck, *et al.*, Crystallization of DNA-coated colloids, *Nat. Commun.*, 2015, **6**(1), 7253.
- 23 A. Hensley, W. M. Jacobs and W. B. Rogers, Self-assembly of photonic crystals by controlling the nucleation and growth of DNA-coated colloids, *Proc. Natl. Acad. Sci. U. S. A.*, 2022, **119**(1), e2114050118.
- 24 L. Ding, X. Chen, W. Ma, J. Li, X. Liu and C. Fan, *et al.*, DNA-mediated regioselective encoding of colloids for programmable self-assembly, *Chem. Soc. Rev.*, 2023, **52**(16), 5684–5705.
- 25 D. Samanta, W. Zhou, S. B. Ebrahimi, S. H. Petrosko and C. A. Mirkin, Programmable Matter: The Nanoparticle Atom and DNA Bond, *Adv. Mater.*, 2022, **34**(12), 2107875.
- 26 P. A. Gabrys, L. Z. Zornberg and R. J. Macfarlane, Programmable Atom Equivalents: Atomic Crystallization as a Framework for Synthesizing Nanoparticle Superlattices, *Small*, 2019, **15**(26), 1805424.
- 27 C. Zhang, R. J. Macfarlane, K. L. Young, C. H. J. Choi, L. Hao and E. Auyeung, *et al.*, A general approach to DNA-programmable atom equivalents, *Nat. Mater.*, 2013, **12**(8), 741–746.
- 28 E. Auyeung, L. Ting, A. J. Senesi, A. L. Schmucker, B. C. Pals and M. O. de la Cruz, *et al.*, DNA-mediated nanoparticle crystallization into Wulff polyhedra, *Nature*, 2014, **505**(7481), 73–77.
- 29 R. J. Macfarlane, B. Lee, H. D. Hill, A. J. Senesi, S. Seifert and C. A. Mirkin, Assembly and organization processes in DNA-directed colloidal crystallization, *Proc. Natl. Acad. Sci. U. S. A.*, 2009, **106**(26), 10493–10498.
- 30 R. J. Macfarlane, M. R. Jones, B. Lee, E. Auyeung and C. A. Mirkin, Topotactic Interconversion of Nanoparticle Superlattices, *Science*, 2013, **341**(6151), 1222–1225.
- 31 D. J. Lewis, L. Z. Zornberg, D. J. D. Carter and R. J. Macfarlane, Single-crystal Winterbottom constructions of nanoparticle superlattices, *Nat. Mater.*, 2020, **19**(7), 719–724.
- 32 S. E. Seo, M. Girard, M. O. de la Cruz and C. A. Mirkin, The Importance of Salt-Enhanced Electrostatic Repulsion in Colloidal Crystal Engineering with DNA, *ACS Cent. Sci.*, 2019, **5**(1), 186–191.
- 33 M. R. Jones, R. J. Macfarlane, B. Lee, J. Zhang, K. L. Young and A. J. Senesi, *et al.*, DNA-nanoparticle superlattices formed from anisotropic building blocks, *Nat. Mater.*, 2010, **9**(11), 913–917.
- 34 R. Jin, G. Wu, Z. Li, C. A. Mirkin and G. C. Schatz, What Controls the Melting Properties of DNA-Linked Gold Nanoparticle Assemblies?, *J. Am. Chem. Soc.*, 2003, **125**(6), 1643–1654.
- 35 R. J. Macfarlane, R. V. Thaner, K. A. Brown, J. Zhang, B. Lee and S. T. Nguyen, *et al.*, Importance of the DNA “bond” in



- programmable nanoparticle crystallization, *Proc. Natl. Acad. Sci. U. S. A.*, 2014, **111**(42), 14995–15000.
- 36 R. J. Macfarlane, M. R. Jones, A. J. Senesi, K. L. Young, B. Lee and J. Wu, *et al.*, Establishing the Design Rules for DNA-Mediated Programmable Colloidal Crystallization, *Angew. Chem.*, 2010, **122**(27), 4693–4696.
- 37 R. D. Blake and S. G. Delcourt, Thermodynamic Effects of Formamide on DNA Stability, *Nucleic Acids Res.*, 1996, **24**(11), 2095–2103.
- 38 L. Levine, J. A. Gordon and W. P. Jencks, The relationship of structure to the effectiveness of denaturing agents for deoxyribonucleic acid, *Biochemistry*, 1963, **2**, 168–175.
- 39 B. L. McConaughy, C. D. Laird and B. J. McCarthy, Nucleic acid reassociation in formamide, *Biochemistry*, 1969, **8**(8), 3289–3295.
- 40 Y. L. Sun, Y. Z. Xu and P. Chambon, A simple and efficient method for the separation and detection of small DNA fragments by electrophoresis in formamide containing agarose gels and Southern blotting to DBM-paper, *Nucleic Acids Res.*, 1982, **10**(19), 5753–5763.
- 41 G. Sarkar, S. Kapelner and S. S. Sommer, Formamide can dramatically improve the specificity of PCR, *Nucleic Acids Res.*, 1990, **18**(24), 7465.
- 42 R. Jungmann, T. Liedl, T. L. Sobey, W. Shih and F. C. Simmel, Isothermal Assembly of DNA Origami Structures Using Denaturing Agents, *J. Am. Chem. Soc.*, 2008, **130**(31), 10062–10063.
- 43 Z. Zhang, J. Song, F. Besenbacher, M. Dong and K. V. Gothelf, Self-assembly of DNA origami and single-stranded tile structures at room temperature, *Angew. Chem., Int. Ed.*, 2013, **52**(35), 9219–92123.
- 44 R. Owczarzy, Y. You, B. G. Moreira, J. A. Manthey, L. Huang and M. A. Behlke, *et al.*, Effects of Sodium Ions on DNA Duplex Oligomers: Improved Predictions of Melting Temperatures, *Biochemistry*, 2004, **43**(12), 3537–3554.
- 45 J. Lee and S. Lee, Non-Invasive, Reliable, and Fast Quantification of DNA Loading on Gold Nanoparticles by a One-Step Optical Measurement, *Anal. Chem.*, 2023, **95**(3), 1856–1866.
- 46 S. J. Hurst, A. K. R. Lytton-Jean and C. A. Mirkin, Maximizing DNA Loading on a Range of Gold Nanoparticle Sizes, *Anal. Chem.*, 2006, **78**(24), 8313–8318.
- 47 S. Lee, H. A. Calcaterra, S. Lee, W. Hadibrata, B. Lee and E. Oh, *et al.*, Shape memory in self-adapting colloidal crystals, *Nature*, 2022, **610**(7933), 674–6749.
- 48 N. Bhatt, P. J. J. Huang, N. Dave and J. Liu, Dissociation and Degradation of Thiol-Modified DNA on Gold Nanoparticles in Aqueous and Organic Solvents, *Langmuir*, 2011, **27**(10), 6132–6317.
- 49 M. S. Lowenthal, E. Quittman and K. W. Phinney, Absolute quantification of RNA or DNA using acid hydrolysis and mass spectrometry, *Anal. Chem.*, 2019, **91**(22), 14569–14576.
- 50 J. G. Wetmur and N. Davidson, Kinetics of renaturation of DNA, *J. Mol. Biol.*, 1968, **31**(3), 349–370.
- 51 J. R. Hutton, Renaturation Kinetics and thermal stability of DNA in aqueous solutions of formamide and urea, *Nucleic Acids Res.*, 1977, **4**(10), 3537–3555.
- 52 K. L. Young, M. B. Ross, M. G. Blaber, M. Rycenga, M. R. Jones and C. Zhang, *et al.*, Using DNA to Design Plasmonic Metamaterials with Tunable Optical Properties, *Adv. Mater.*, 2014, **26**(4), 653–659.
- 53 C. Chen, X. Luo, A. E. K. Kaplan, M. G. Bawendi, R. J. Macfarlane and M. Bathe, Ultrafast dense DNA functionalization of quantum dots and rods for scalable 2D array fabrication with nanoscale precision, *Sci. Adv.*, 2023, **9**(32), eadh8508.

