

Supplementary Information

Acoustically Shaped DNA-Programmable Materials

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Supplementary Methods

Acoustic Transducer

A Y+128° X-propagation lithium niobate (LiNbO₃) wafer (500 μm thick) was coated with a layer of AZ5214 photoresist (MicroChem, Newton, MA) through spin-coating. The coated wafer underwent lithography using a maskless aligner (MLA 150-Heidelberg) for precise patterning and later treated with a photoresist developer (Microchem AZ726) for development. Subsequently, an e-beam evaporator (Bestec) was used to deposit a double metal layer (Cr/Au, 100Å/1000Å) onto the wafer. Following this procedure, a lift-off process was employed to create the interdigital transducers (IDTs).

Crystallites Orientations

Images of before and after the acoustic field was applied were used to assess the orientation of crystals, as seen in Supplementary Figure 2. The angle measured was from the parallel to the capillary wall. The chosen angle was always between 0°-45° as shown in Figure 2a. Analysis was performed using *Fiji*¹ (*version 1.54f*).

Small-Angle X-ray Scattering (SAXS) of Assembled Structures

The SAXS measurements were performed at the Complex Materials Scattering (CMS) beamline at the National Synchrotron Light Source II (NSLS-II) at Brookhaven National Laboratory (BNL) in Upton, NY. The 2D scattering data was collected on area detectors located downstream of the sample's position. Area images were integrated into a one-dimensional (1D) $I(q)$ scattering curve as a function of the scattering vector q , where $q = 4\frac{\pi}{\lambda} \sin\left(\frac{\theta}{2}\right)$ with λ and θ being the wavelength of the incident X-rays and the full scattering angle, respectively. The resultant 1D curves spanned

22 roughly 0.04 nm^{-1} to 1 nm^{-1} with a resolution of 0.002 nm^{-1} . The structure factor $S(q)$ was obtained
 23 by dividing $I(q)$ by the corresponding particle form factor $P(q)$.

Beamline	11-BM CMS
Photon Energy (keV)	13.5
Horizontal \times Vertical Beam size ($\mu\text{m} \times \mu\text{m}$)	200 x 200
Approximate Flux (photons/sec)	10^{11}
Sample-to-Detector Distance (m)	5.05
Detector Manufacturer	Dectris
Detector Model	Pilatus 1M
Detector Pixel Size ($\mu\text{m} \times \mu\text{m}$)	172 x 172

24 **Supplementary Table 1.** X-ray beam characteristics and setup details at the CMS (NSLS-II) beamline

25 In this work, we implement modeling of the presented analysis using the ScatterSim software
 26 package, a python package that implements a scattering formalism for superlattices^{2,3}. This
 27 formalism can generically model the scattering from arbitrary anisotropic nano-objects within the
 28 unit cell of a regular superlattice. We used this library to perform the modeling for the simple cubic
 29 crystal, with the code being available for download on Git Hub³.

30 **Supplementary Discussion**

31 **Acoustic Field Considerations**

32 SAWs are generated by IDTs fabricated with photolithography to convert electrical signals
 33 into acoustic waves through the piezoelectric effect of a substrate. The acoustic force applied on
 34 particles is dependent on the acoustic contrast between the medium and particles, which is
 35 sufficient for most systems of particles in aqueous media. The SSAW-induced pressure generates
 36 a force F_{ac} , applied on small particles, has a form:

$$37 \quad F_{ac} = \frac{4}{3} \pi r^3 E_{ac} q \sin(2kx) \phi$$

38 where r is the radius of the particle, E_{ac} is the acoustic energy density, q is the wave number, x is
 39 the distance of the particle from the node and ϕ is the acoustic contrast factor between the particle
 40 and the medium (water).

41 Acoustic contrast factor

42 To estimate the acoustic contrast factor ϕ of our material, we first consider a hypothetical case
43 when DNA origami fills entire volume of the frame. The density ratio of DNA to water is
44 approximately $\rho_{DNA}/\rho_{H_2O} \approx 1.7$ which corresponds to

45
$$\phi_{DNA} = \frac{5\rho_{DNA} - 2\rho_{H_2O}}{2\rho_{DNA} + \rho_{H_2O}} - \frac{\beta_{DNA}}{\beta_{H_2O}} \approx 1.5 - \frac{\beta_{DNA}}{\beta_{H_2O}}$$

46 where β_{DNA}/β_{H_2O} is the compressibility ratio of DNA to water. Since we know, based on
47 experimental evidence, that gold particles and DNA frames are pushed to the same locations
48 (Supplementary Figure 4), we can conclude that ϕ_{DNA} is positive. Hence, the compressibility ratio
49 $\frac{\beta_{DNA}}{\beta_{H_2O}}$ is between 0 and 1.5. Plotting a particle edge length threshold (L^*) vs the compressibility
50 ratio (Supplementary Figure 5) shows that even for energy of 200 J/m^3 (See ‘Acoustic Force
51 Considerations’), individual particles need to be well above a threshold of 30 nm in order for the
52 acoustic force to push them efficiently.

53 Acoustic Force Considerations

54 The force applied by the waves on particles is dependent on the position of the particle, where at
55 specific $n\pi$ locations called the ‘nodes’, the force will be reduced to zero. Small particles with
56 positive acoustic contrast around the nodes are pushed into the nodes, where they will stay for as
57 long as the wave is applied. As estimated above, the contrast factors for individual DNA frames
58 as well as for the larger self-assembled structures are expected be close to the volume fraction of
59 DNA in them, *i.e.* $\phi \sim 0.1$. For a particle to be affected by the SSAW, its energy difference between
60 the node and antinode positions, $VE_{ac}\phi$, must exceed kT . To estimate the particles size threshold
61 (L^*) that will be affected by the acoustic force, the acoustic energy density (E_{ac}) must be estimated
62 first. By using polystyrene beads of $1 \mu\text{m}$ (Supplementary Movie 5) and calculating the velocity
63 of the beads under the acoustic force, we estimated that the acoustic energy density is $E_{ac} \approx$
64 50 J/m^3 (for more details on the calculation, see Supplementary Table 2). This corresponds to the
65 particle size range of $L \gtrsim \left(\frac{kT}{E_{ac}\phi}\right)^{\frac{1}{3}} \approx 80 \text{ nm}$ (Supplementary Figure 5). This means that individual
66 DNA frames ($\sim 30 \text{ nm}$) should not be affected, while the crystalline particles larger than the critical
67 nucleus typically are.

Parameter	Description	Equation	Value
ρ (kg/m^3)	Density of polystyrene particles	-	1050
ϕ_w	Acoustic contrast of polystyrene in water	-	0.193
r (m)	Radius of polystyrene particles	-	0.5e-6
η ($N \cdot s/m^2$)	Viscosity	-	6.5e-4
d (m)	Distance particles travel (from movie S5)	-	75e-6
t (s)	Time for particles to travel (from movie S5)	-	18
v (m/s)	Velocity	d/t	4.16e-6
F_{drag} (N)	Drag force	$6\pi\eta r v$	2.55e-14
m (kg)	Particle mass	$\frac{4}{3}\pi r^3 \rho$	5.5e-16
$F_{calculated}$ (N)	Total force on particles	$m \cdot a$	1.27e-22
F_{ac} (N)	Acoustic force on particles	$F_{calculated} + F_{drag}$	2.55e-14
q (m^{-1})	Wavenumber	$1/\lambda$	5000
E_{ac} (J/m^3)	Acoustic Energy Density	$\frac{3}{4\pi r^3 q \sin(2kx) \phi_w} \cdot F_{ac}$	50.5

68 **Supplementary Table 2.** Acoustic energy density calculation table. The acoustic force calculation was performed for
69 microparticles with a known diameter, density, compressibility and acoustic contrast factor⁴ in a capillary tube filled
70 with water (0.05 x 1 mm). IDT 19.34 MHz and Amplitude of 20Vpp. The velocity and acceleration of the particles
71 were calculated by measuring the time it takes for the particles to cover a known distance. We used the time it takes
72 all dispersed particles to reach the nodes and considered half the node distance as travel distance.

73 Theoretical Modelling Parameters

74 The table below (Supplementary Table 3) lists the parameters used for modeling. Out of them,
75 Γ was used as a tuning parameter for the unmodified nucleation and growth theory that describes
76 the no-wave data. Two additional fitting parameters are specific to the infusion model. The rest of
77 the parameters were determined based on the known geometrical and thermodynamic properties
78 of the building blocks, and the experimental conditions. In particular, the diffusion coefficient was
79 determined by Einstein relationship (by setting hydrodynamic radius to a):

80 $D = \frac{kT}{6\pi\eta a} = 5.9 \cdot 10^{-12} \frac{m^2}{s}$. The sequence-specific hybridization entropy of each DNA duplex

81 was found to be $160 \frac{cal}{K \cdot mol}$. There are 4 DNA duplexes in each bond, and 3 bonds per octahedral
82 building block in the cubic lattice, which gives:

$$83 \quad \Delta S/kT = \frac{12 \cdot 160 \frac{cal}{K \cdot mol}}{RT} = 3.0 K^{-1}.$$

Parameter	a	c_0	D	$\Delta S/kT$	Γ	ε	κ
Value	40 nm	40 nM	$5.9 \cdot 10^{-12} \frac{m^2}{s}$	$3.0 K^{-1}$	7.5 (fitting)	400 (fitting)	$0.0005 s^{-1}$ (fitting)

84 **Supplementary Table 3.** Parameters for the theoretical model used in the study.

85 **Supplementary Tables**

86 **Internal DNA Staples**

Sequence Name	Sequence
OCT-Staple-1	GCCTTGAATCTTTCCGGAACCGCTCCAGAGCCAGAGCCGCCAGCATT
OCT-Staple-2	ATAAGGCGCCAAAAGTTGAGATTTAGGATAACGGACCAGTCA
OCT-Staple-3	GATGGTTGGGAAGAAAAATCCACCAGAAATAATTGGGCTTGA
OCT-Staple-4	GGACGTTTAATTTTCGACGAGAAACACCACCACTAATGCAGAT
OCT-Staple-5	ATTTTAAGAAGCTGGCTTGAATTATCAGTGA
OCT-Staple-6	TCAGAACCGCCACCCTCTCAGAGTATTAGC
OCT-Staple-7	TCAGAGCGCCACCACATAATCAAAATCAGAACGAGTAGTATG
OCT-Staple-8	AACCAGACGCTACGTTAATAAAACGAACATACCACATTCAGG
OCT-Staple-9	GTTTGCCTATTACAGGCAGGTCAGACGCCACCACACCACC
OCT-Staple-10	ACATAACTTGGCCCTAACTTTAATCATTGCATTATAACAACATTATTACAGGTAG
OCT-Staple-11	CATGTCACAAACGGCATTAAATGTGAGCAATTCGCGTTAAAT
OCT-Staple-12	AAGATTGTTTTTAACCAAGAAACCATCGACCCAAAAACAGG
OCT-Staple-13	CAGCTCATATAAGCGTACCCCGGTTGATGTGTCGGATTCTCC
OCT-Staple-14	GTTAAAATTCGCATTATAAACGTAAACTAG
OCT-Staple-15	AGCACCATTACCATTACAGCAAATGACGGGA
OCT-Staple-16	GTCACCAGAGCCATGGTGAATTATACCAATCAGAAAAGCCT
OCT-Staple-17	CCGACTTATTAGGAACGCCATCAAAAATGAGTAACAACCCCA
OCT-Staple-18	AATTATTGTTTTCATGCCTTTAGCGTCAGATAGCACGGAAAC
OCT-Staple-19	CTTCGCTGGGCGCAGACGACAGTATCGGGGCACCGTCGCCATTCAGGCTGCGCA
OCT-Staple-20	ACAAAGAAATTTAGGTAGGGCTTAATTGTATACAACGGAATC
OCT-Staple-21	TGACCTACTAGAAAAAGCCCCAGGCAAAGCAATTTTCATCTTC
OCT-Staple-22	ATAATTAATTTAAAAAACTTTTTCAAACCTTTTAACAACGCC
OCT-Staple-23	AGGCGTTAAATAAGAAGACCGTGTGCGCAAG
OCT-Staple-24	TGCCGGAAGGGGACTCGTAACCGTGCATTATATTTTAGTTCT
OCT-Staple-25	CTCCAGCCAGCTTTCCCCTCAGGACGTTGG
OCT-Staple-26	CAGTTTGAATGTTTAGTATCATATGCGTAGAATCGCCATAGC
OCT-Staple-27	TGTAGATATTACGCGGCGATCGGTGCGGGCGCCATCTTCTGG
OCT-Staple-28	AACATGTACGCGAGTGGTTTAAAATACCTAAACACATTCTTACCAGTATAAAGC
OCT-Staple-29	CGCTGGTGCTTTCCTGAATCGGCCAACGAGGGTGGTGATTGCCCTTACCAGCCT
OCT-Staple-30	AACAAAAATAACTAGGTCTGAGAGACTACGCTGAGTTTCCCT
OCT-Staple-31	AACAGTACTTGAAAACATATGAGACGGGTCTTTTTTAATGGA
OCT-Staple-32	TAGAATCCATAAATCATTTAACAATTTCTCCCGGCTTAGGTT
OCT-Staple-33	GTAAATCGTCGCTATTGAATAACTCAAGAA
OCT-Staple-34	TTGCGTATTGGGCGCCCGGGGTGCGCTC
OCT-Staple-35	TTTACCAGCATTAAAGTCGGGAAACCTGATTTGAATTACCCA
OCT-Staple-36	GCCAGCTAGGCGATAGCTTAGATTAAGACCTTTTTAACCTGT
OCT-Staple-37	ACTGCCCTTGCCCGTTGCAGCAAGCGGCAACAGCTTTTTCT
OCT-Staple-38	GGGTTATTTAATTACAATATATGTGAGTAATTAATAAGAGTCAATAGTGAATTT
OCT-Staple-39	TCCAAATCTTCTGAATTATTTGCACGTAGGTTTAACGCTAACGAGCGTCTTTCC

OCT-Staple-40	CAGATATTACCTGAATACCAAGTTACAATCGGGAGCTATTTT
OCT-Staple-41	ACGCGAGGCTACAACAGTACCTTTTACAAATCGCGCAGAGAA
OCT-Staple-42	GCACCCAGCGTTTTTATCCGGTATTCTAGGCGAATTATCA
OCT-Staple-43	AAGCCTTAAATCAAGACTTGCGGAGCAAAT
OCT-Staple-44	ATTGCGTAGATTTTCAAAACAGATTGTTTG
OCT-Staple-45	TGAATATTATCAAAAATAATGGAAGGGTTAATATTTATCCCAA
OCT-Staple-46	CCTACCAACAGTAATTTTATCCTGAATCAAACAGCCATATGA
OCT-Staple-47	GATTATAAAGAAACGCCAGTTACAAAATTTACCAACGTCAGA
OCT-Staple-48	TTTCAATAGAAGGCAGCGAACCTCCCGATTAGTTGAAACAATAACGGATTCCGC
OCT-Staple-49	GATATTCTAAATTGAGCCGGAACGAGGCCCAACTGGCGCATAGGCTGGCTGAC
OCT-Staple-50	GGTTGATTTTCCAGCAGACAGCCCTCATTTCGTCACGGGATAG
OCT-Staple-51	AGTACCGAATAGGAACCCAAACGGTGTAACTCAGGAGGTTT
OCT-Staple-52	CAAGCCCCACCCTTAGCCCGAATAGGACGATCTAAAGTTT
OCT-Staple-53	CAGAGCCACCACCCTCTCAGAACTCGAGAG
OCT-Staple-54	AAGGGAACCGAAGTACGAGCAGCGGTATCAT
OCT-Staple-55	GGACAGAGTTACTTTGTGCGAAATCCGCGTGTATCACCGTACG
OCT-Staple-56	GCTCCATTGTACCGTAACACTGAGTTAGTTAGCGTAACT
OCT-Staple-57	CGCTGAATTACCCTAATCTTGACAAGACAGACCATGAAAGA
OCT-Staple-58	TGTCGTCATAAGTACAGAACCGCCACCCATTTTCACAGTACAACTACAACGCC
OCT-Staple-59	ATGACCACTCGTTTGGCTTTTGCAAAAGTTAGACTATATTCATTGAATCCCCT
OCT-Staple-60	GTAATACGCAAACATGAGAGATCTACAACACTAGCTGAGGCCGG
OCT-Staple-61	AGAACCCCAAATCACCATCTGCGGAATCGAATAAAAATTTTT
OCT-Staple-62	AGACAGTTCATATAGGAGAAGCCTTTATAACATTGCCTGAGA
OCT-Staple-63	GTAAAGATTCAAAGGCCTGAGTTGACCCT
OCT-Staple-64	GGTAATAGTAAATGTAAGTTTTACTAT
OCT-Staple-65	GTCCAATAGCGAGAACCAGACGACGATTTCAACGCAAGGGA
OCT-Staple-66	CCAAAATACAATATGATATTCAACGTTAGGCTATCAGGTAA
OCT-Staple-67	CATAACCTAAATCAACAGTTCAGAAAACGTCATAAGGATAGC
OCT-Staple-68	GTCTGGATTTTGCCTTTTAAATGCAATGGTGAGAAATAAATTAATGCCGGAGAG
OCT-Staple-69	GGGCGACCCCAAAAGTATGTTAGCAAACATAAAGAGTCACAATCAATAGAAAAT
OCT-Staple-70	TATAAAGCATCGTAACCAAGTACCGCACCGGCTGTAATATCC
OCT-Staple-71	CAACATGATTTACGAGCATGGAATAAGTAAGACGACAATAAA
OCT-Staple-72	CATCCTATTACGCTAAAAGGTAAGTAAAAAGCAAGCCGTTT
OCT-Staple-73	GATAAGTCCTGAACAACTGTTTAAAGAGAA
OCT-Staple-74	TAAAGGTGGCAACATAGTAGAAAATAATAA
OCT-Staple-75	AGACACCTTACGCAGAACTGGCATGATTTCTGTCCAGACAA
OCT-Staple-76	CTCCTTAACGTAGAAAACCAATCAATAATTCATCGAGAACAGA
OCT-Staple-77	CGGAATAATTCAACCGAGCCAAAGACTTATTTTAAACGCAA
OCT-Staple-78	TTATTTTTACCACAATGCAGAACGCGGAAAAATCTTTCCTTATCATTCCAAG
OCT-Staple-79	CAGCCTTGGTTTTGTATTAAGAGGCTGACTGCCTATATCAGA
OCT-Staple-80	GGAAGCGCCACAAACAGTTAATGCCCCGACTCCTCAAGATA
OCT-Staple-81	GAGATAACATTAGAAGAATAACATAAAAAGGAAGGATTAGGA
OCT-Staple-82	GTCAGAGGGTAATTGAGAACACCAAAATAG

OCT-Staple-83	AAGTTTTAACGGGGTCGGAGTGTAGAATGG
OCT-Staple-84	CAGTGCCTACATGGGAATTTACCGTTCCACAAGTAAGCAGAT
OCT-Staple-85	AGCGTCACGTATAAGAATTGAGTTAAGCCCTTTTTAAGAAAG
OCT-Staple-86	AAAGCGCCAAAGTTTATCTTACCGAAGCCCAATAATGAGTAA
OCT-Staple-87	TGCTAAACAGATGAAGAAACCACCAGAATTTAAAAAAGGCT
OCT-Staple-88	GAGAATAGAGCCTTACCGTCTATCAAATGGAGCGGAATTAGA
OCT-Staple-89	CCAAAAGGAAAGGACAACAGTTTCAGCGAATCATCATATTCC
OCT-Staple-90	TTCACGTTGAAAACTTTGCGAATGGGATTT
OCT-Staple-91	GTCCACTATTAAGAACCAGTTTTGGTTCC
OCT-Staple-92	TCAAAGGGAGATAGCCCTTATAAATCAAGACAACAACCATCG
OCT-Staple-93	ATAGCCCGCGAAAATAATTGTATCGGTTTCGCCGACAATGAGT
OCT-Staple-94	GAAATCGATAACCGGATACCGATAGTTGTATCAGCTCCAACG
OCT-Staple-95	ATTAAGTATAAAGCGGCAAGGCAAAGAACTAATAGGGTACC
OCT-Staple-96	CACGACGAATTCGTGTGGCATCAATTCCTTTAGCAAAAATTACG
OCT-Staple-97	CAGGTCGACTCTAGAGCAAGCTTCAAGGCG
OCT-Staple-98	TAACTGTTTAGCTATTTTCGCATTCATTC
OCT-Staple-99	GAGTCGTTGTAAACGCCAGGGTTTTCCAAAGCAATAAAGCC
OCT-Staple-100	CGCGAGCTTAGTTTTTCCCAATTCTGCGCAAGTGTAAGCCCT
OCT-Staple-101	AGTAGATTGAAAAGAATCATGGTCATAGCCGGAAGCATAAGT
OCT-Staple-102	CATATAACTAATGAACACAACATACGAGCTGTTTCTTTGGGG
OCT-Staple-103	ATGTTTTGCTTTTGATCGGAACGAGGGTACTTTTTCTTTGATAAGAGGTCATT
OCT-Staple-104	AGAAGCAACCAAGCCAAAAGAATACACTAATGCCAAAACCTC
OCT-Staple-105	GAGGAAGCAGGATTCGGGTAAAATACGTAAAACACCCCCAG
OCT-Staple-106	AACAGGTCCCGAAATTGCATCAAAAAGATCTTTGATCATCAG
OCT-Staple-107	TCAAAGCGAACCAGACCGTTTTATATAGTC
OCT-Staple-108	GCTTTGAGGACTAAAAGCAACGGGGAGTT
OCT-Staple-109	AAGTTTCAGACAGCCGGGATCGTCAACCTTCTGTAGCTCAAC
OCT-Staple-110	CAGCGAACATTAAGAGAGTACCTTTACTGAATATAATGAA
OCT-Staple-111	AAAGGCCAAATATGTTAGAGCTTAATTGATTGCTCCATGAGG
OCT-Staple-112	CGATTATAAGCGGAGACTTCAAATATCGCGGAAGCCTACGAAGGCACCAACCTA
OCT-Staple-113	GGGGTGCCAGTTGAGACCATTAGATACAATTTTCACTGTGTGAAATTGTTATCC
OCT-Staple-114	TCAGAGCTGGGTAAACGACGGCCAGTGCATCCCCGTAGTAGCATTAAACATCCA
OCT-Staple-115	TTAGCGGTACAGAGCGGGAGAATTAAGTGCCTAATTCGGAACCTATTATTCT
OCT-Staple-116	TGATTATCAACTTTACAATAAAGGAATCCAAAAAGTTTGAGTAACATTATCAT
OCT-Staple-117	GTAGCGCCATTAATTGGGAATTAGAGCGCAAGGCGCACCGTAATCAGTAGCGA
OCT-Staple-118	AGCCGAAAGTCTCTCTTTTGATGATACAAGTGCCTTAAGAGCAAGAAACAATGA
OCT-Staple-119	GTGGGAAATCATATAAATATTTAAATTTGAATTTTTGTCTGGCCTTCCTGTAGCC
OCT-Staple-120	CCCACGCGCAAATGGTTGAGTGTGTTTCGTGGACTTGCTTTCGAGGTGAATTT

87 **Supplementary Table 4.** Sequence list of produced DNA origami octahedron structure. Without further strands
88 ('sticky ends'), this DNA structure will not be capable of binding to other origami.

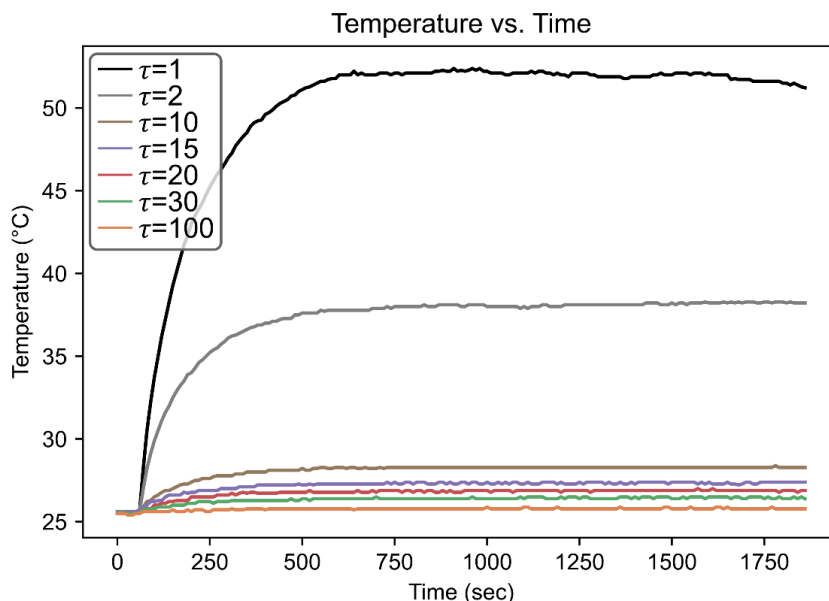
89 **External Staples ('Sticky Ends')**

Sequence Name	Sequence
Vertex A-1	AGAGCCTAATTGATTTTTGTTTAAATCCTGAAATAAAGAATTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex A-2	TGTAGCATTCACGTTAGTAAATGAAGTGCCGCGCCACCCTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex A-3	GAAACATGAAAGCTCAGTACCAGGCGAAAAATGCTGAACAAATTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex A-4	TTTGC GGAAACAATGGCAATTCATCAATCTGTATAATAATTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex B-1	AAAGATTCATCAGGAATTACGAGGCATGCTCATCCTTATGCGTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex B-2	CTTCATCAAGAGAAATCAACGTAACAGAGATTGTCAATCATTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex B-3	CAAATGCTTTAAAAAATCAGGCTTTAAGAGCAGCCAGAGGGTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex B-4	AAACGAAAGAGGGCGAAACAAAGTACTGACTATATTCGAGCTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex C-1	AGCTTTCATCAACGGATTGACCGTAAAATCGTATAATTTTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex C-2	ACTGTTGGGAAGCAGCTGGCGAAAAGGATAGGTCAAGATCGCATTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex C-3	GGTAGCTATTTAGAGAATCGATGAAAACATTAAATGTGTAGTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex C-4	ATAAATCATACATAAATCGGTTGTACTGTGCTGGCATGCCTGTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex D-1	TGATTGCTTTGAGCAAAAAGAAGATGAAATAGCAGAGGTTTTGTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex D-2	AACGGGTATTAAGGAATCATTACGCCAGTAATCAACAATTTTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex D-3	CAACGCTAACAGCAGAGGCATTTCAATCCAATGATAAATTTTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex D-4	ATCAAAATCATATATGTAATGCTGAACAAACACTTGCTCTTTTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex E-1	GGCCCTGAGAGAAGCAGGCGAAAATCATTGCGTAGAGGCGTTTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex E-2	CTTAAACAGCTTATATATTCGGTCGCTTGATGGGGAACAAGATTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex E-3	GCTCACAATTCGTGAGCTAACTCACTGGAAGTAATGGTCAATTTTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex E-4	TTTGC GGATGGCCAACTAAAGTACGGGCTGCAGCTACAGAGTTTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex F-1	GACAGGAGGTTGAAACAAATAAATCCGCCCCCTCCGCCACCCTTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex F-2	CAGAATCAAGTTTCGGCATTTTCGGTTAAATATATCACCAGTTTTTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)

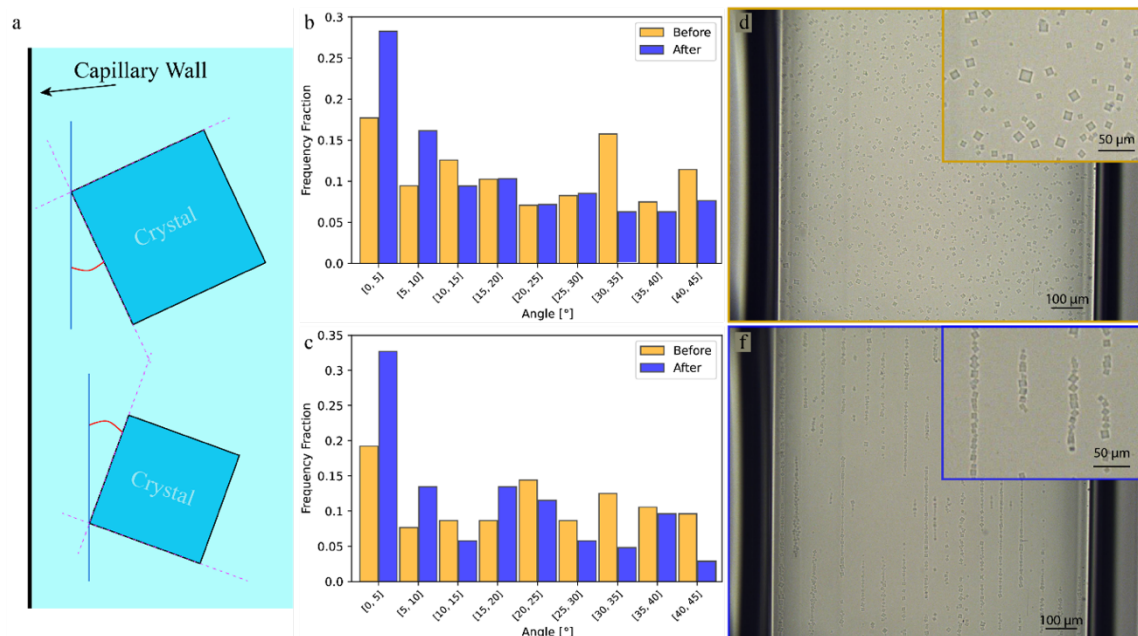
	(Binding Sequence)
Vertex F-3	TCATATGGTTTACGATTGAGGGAGGGAAACGCAATACATATTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Vertex F-4	AATAGCAATAGCACCAGAAGGAAACCTAAAGCCACTGGTAATTTTTTTTTTTTTTTTTTTTTT + (Binding Sequence)
Binding Sequence A	ACCTACAC
Binding Sequence B	GTGTAGGT

90 **Supplementary Table 5.** Externally facing sticky ends located at the 6 vertices of the octahedron origami frame.
91 There are 4 sequences (1-4) present at each vertex (A-F), where our designs keep the 'Binding Sequence' the
92 same for these 4 sequences. The Binding Sequences are listed at the end of the tables.

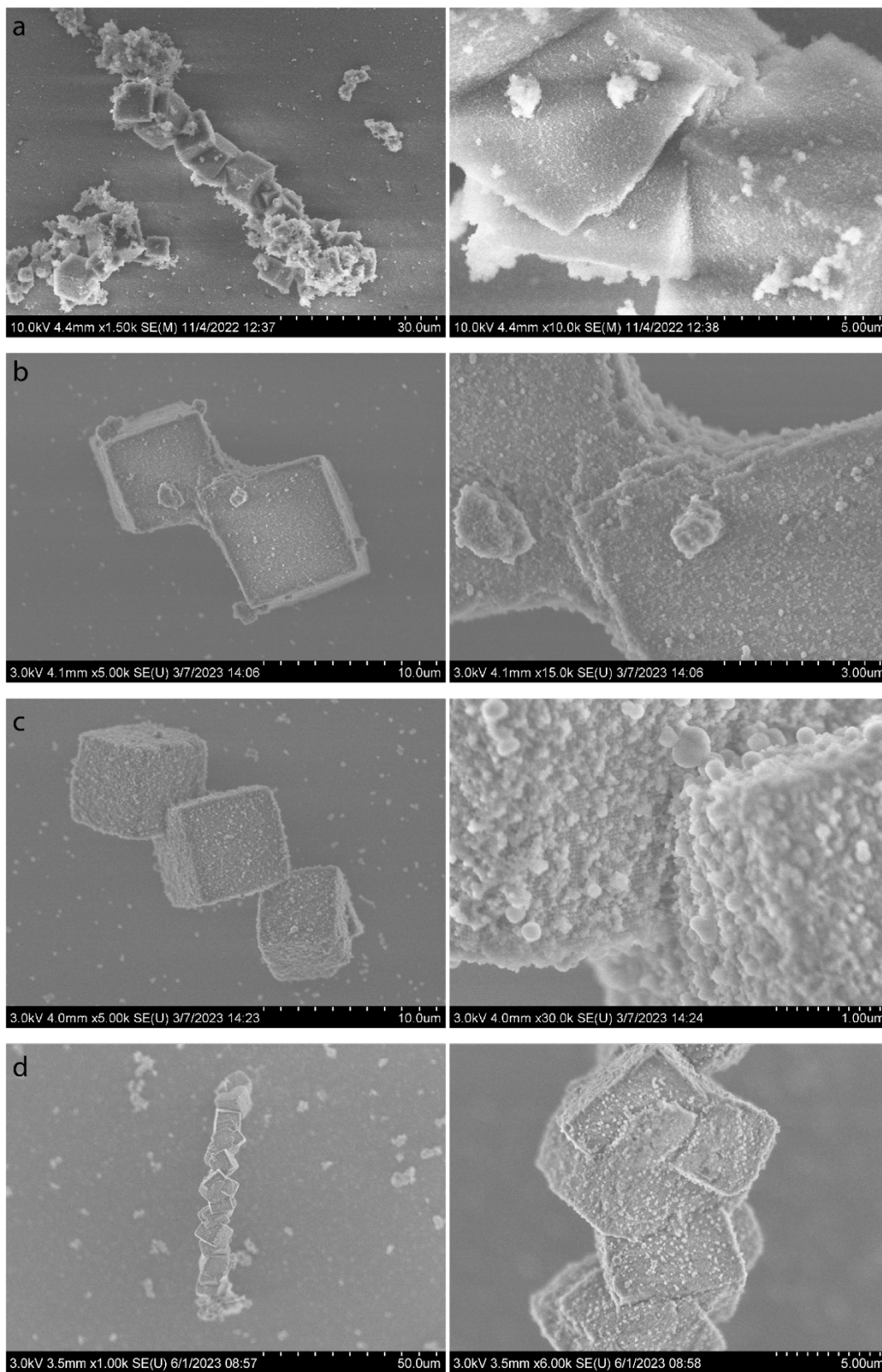
93 **Supplementary Figures**



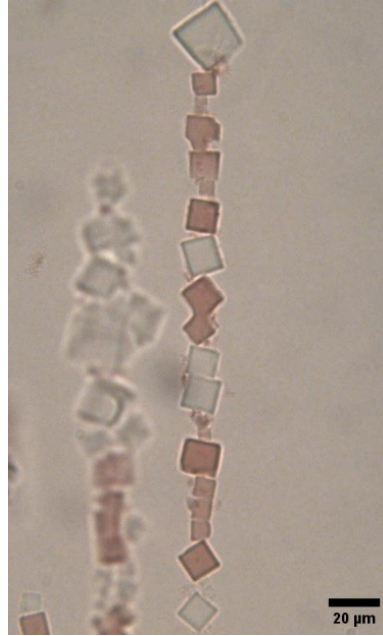
94
 95 **Supplementary Figure 1.** Temperature change induced by the acoustic waves in the active region. τ is defined
 96 as the period divided by the pulse length. $\tau=1$ means constant waves are applied. Waves were applied after 60
 97 seconds at room temperature (25.5 °C) without the use of active temperature control. τ values in the experiments
 98 conducted were between 10 – 100. Thermal paste was used to ensure proper temperature measurement of the
 99 active region.



100
 101 **Supplementary Figure 2.** Orientation of crystals before and after acoustic organization. **a.** Schematic of how
 102 the orientation of crystals was measured. The angle is always between 0° to 45°. **b,c.** Histogram of two separate
 103 experiments where the angles of individual crystals were measured before and after acoustic waves were applied.
 104 Panel **b** Before n=254, After n=223; Panel **c** Before n=104, After n=104. **d,f.** Images corresponding to the
 105 Histogram in panel **b**, before (**d**) and after (**f**) the acoustic waves. The Histogram in panel **(c)** corresponded to
 106 Supplementary Movie 2.

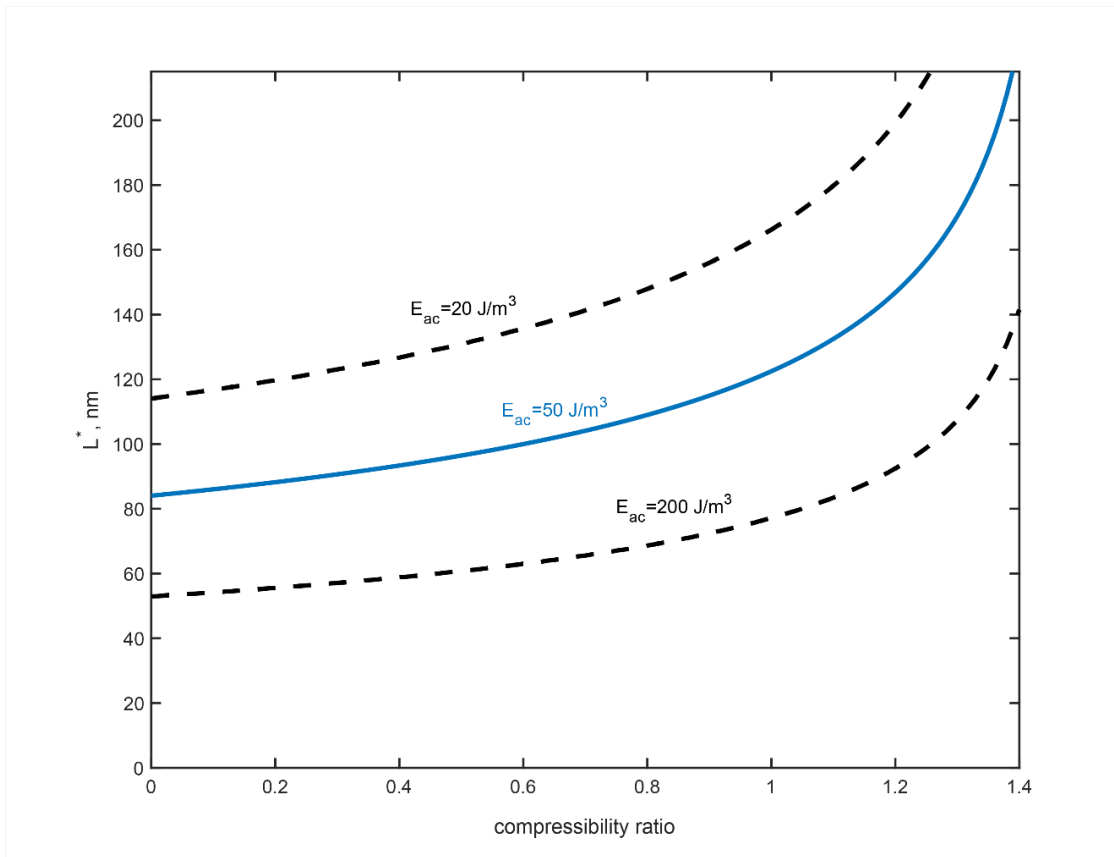


107
 108 **Supplementary Figure 3. a-d.** Additional SEM images of fused crystals after reannealing. Zoomed-out (left)
 109 and zoomed-in (right) images of the same region.



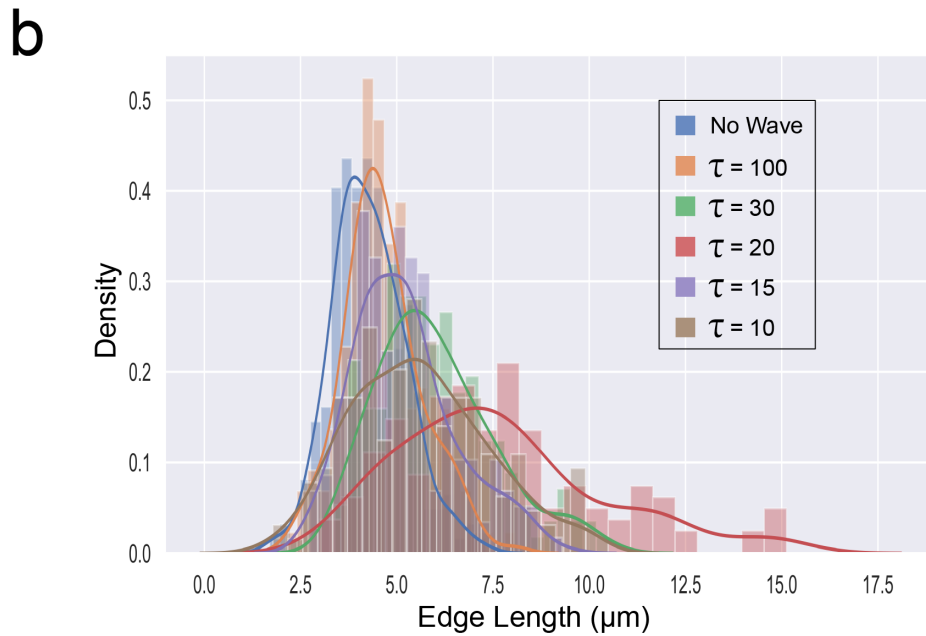
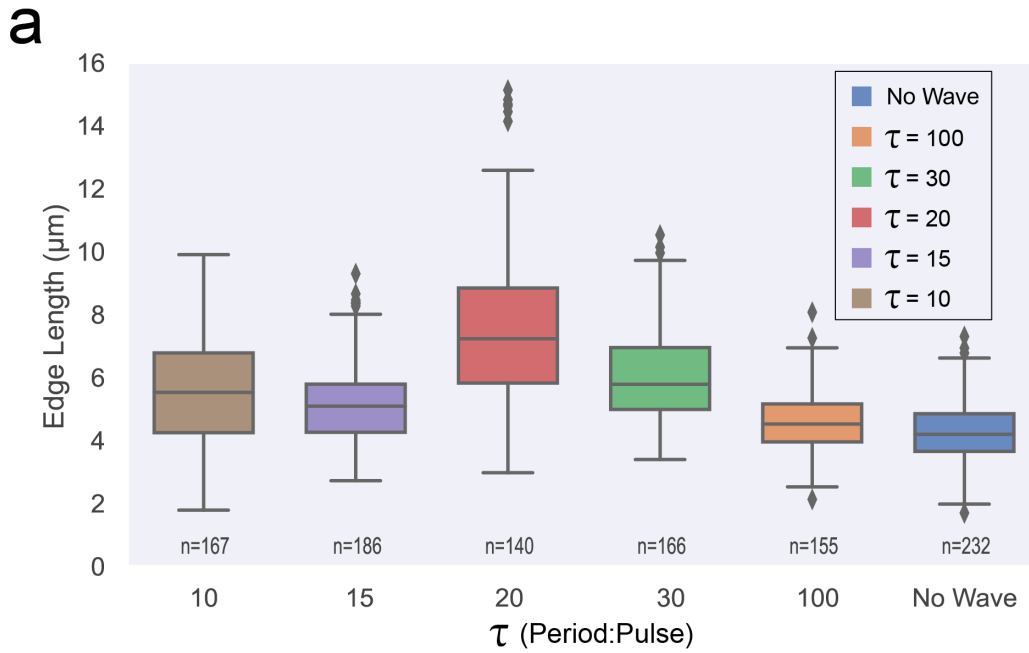
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Supplementary Figure 4. Crystals are arranged linearly by acoustic field. Crystals filled with AuNPs are red, while empty crystals are clear.



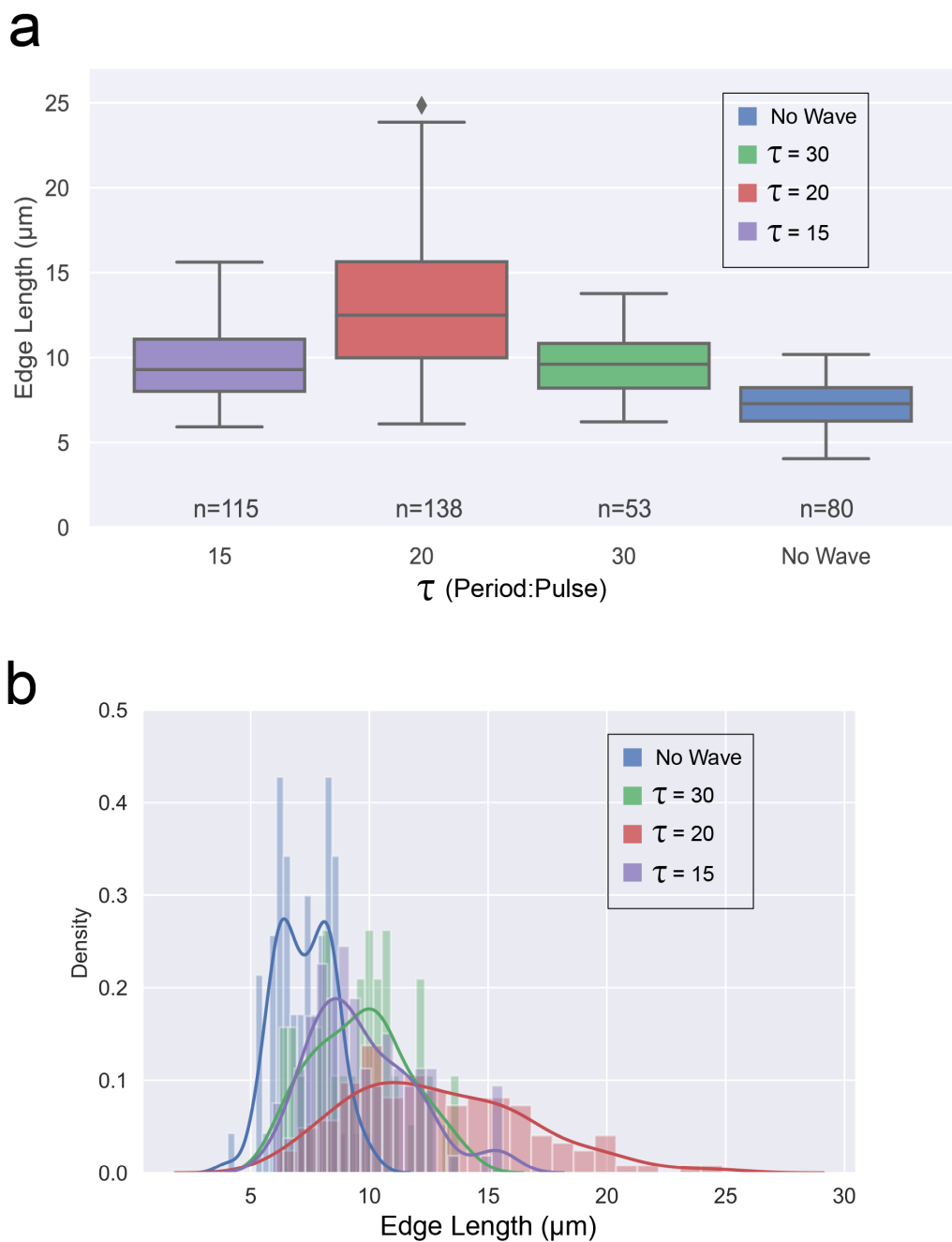
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Supplementary Figure 5. Edge length threshold (L^*) vs the compressibility ratio (β_{DNA}/β_{H_2O}) calculated for a range of acoustic energy densities between 20 and 200 J/m^3 , where 50 J/m^3 (blue) is the estimated energy density for this study (Supplementary Table 2).



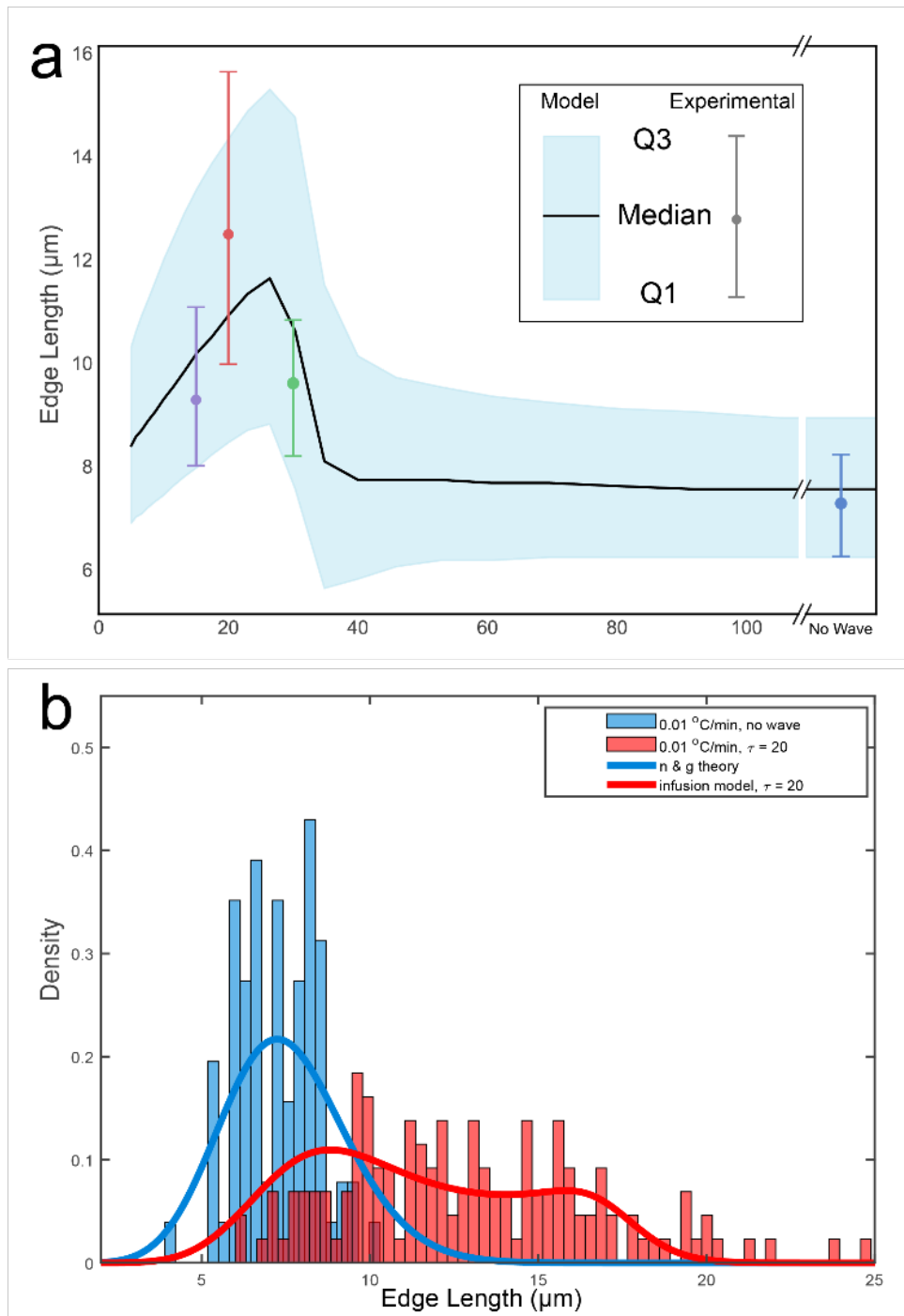
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118 **Supplementary Figure 6. a.** Edge length box plot and distribution of crystals after formation with acoustic wave
 119 pulses with a *fast* temperature decrease rate of 0.03 °C/min. **b.** Edge length distribution of crystals after
 120 formation with acoustic wave pulses with a *fast* temperature rate of 0.03 °C/min.



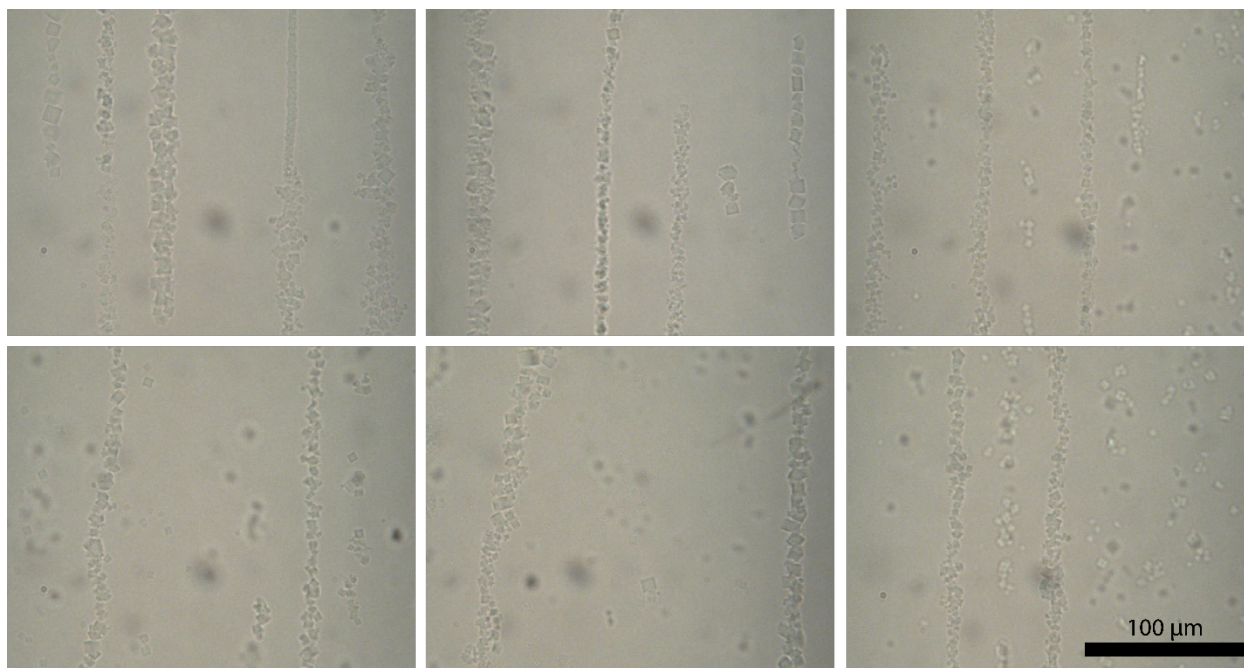
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122 **Supplementary Figure 7. a.** Edge length box plot and distribution of crystals after formation with acoustic wave
 123 pulses with a 'slow' temperature decrease rate of 0.01 °C/min. **b.** Edge length distribution of crystals after
 124 formation with acoustic wave pulses under a 'slow' temperature rate of 0.01 °C/min.

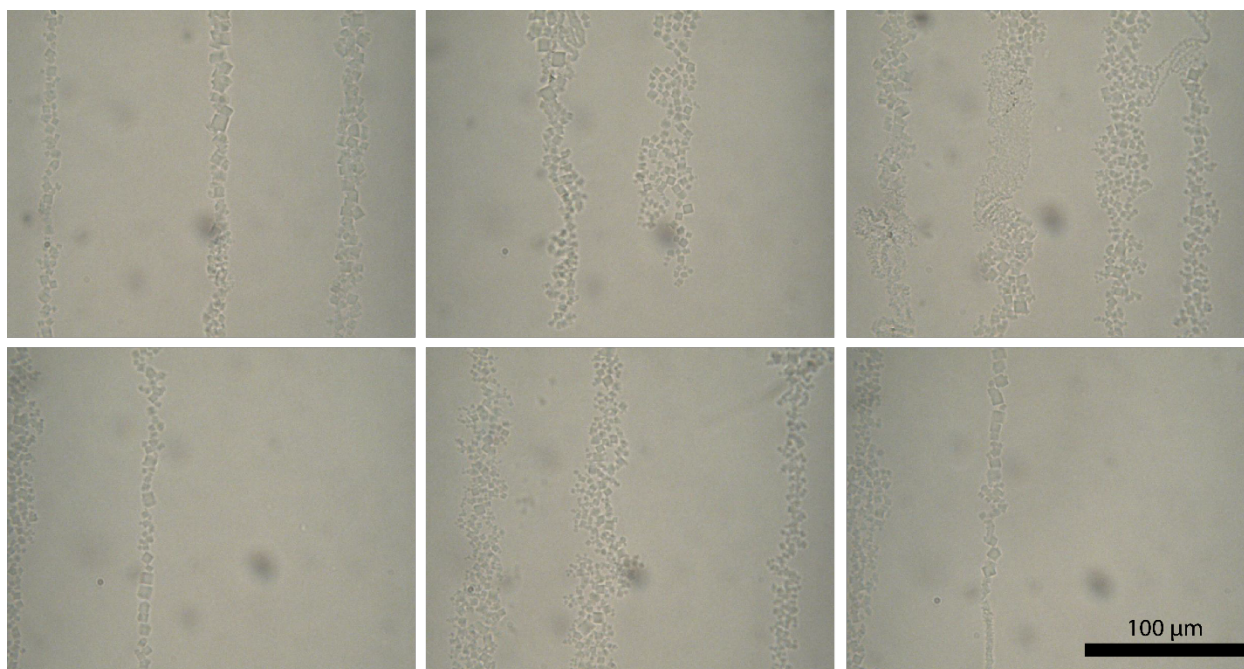


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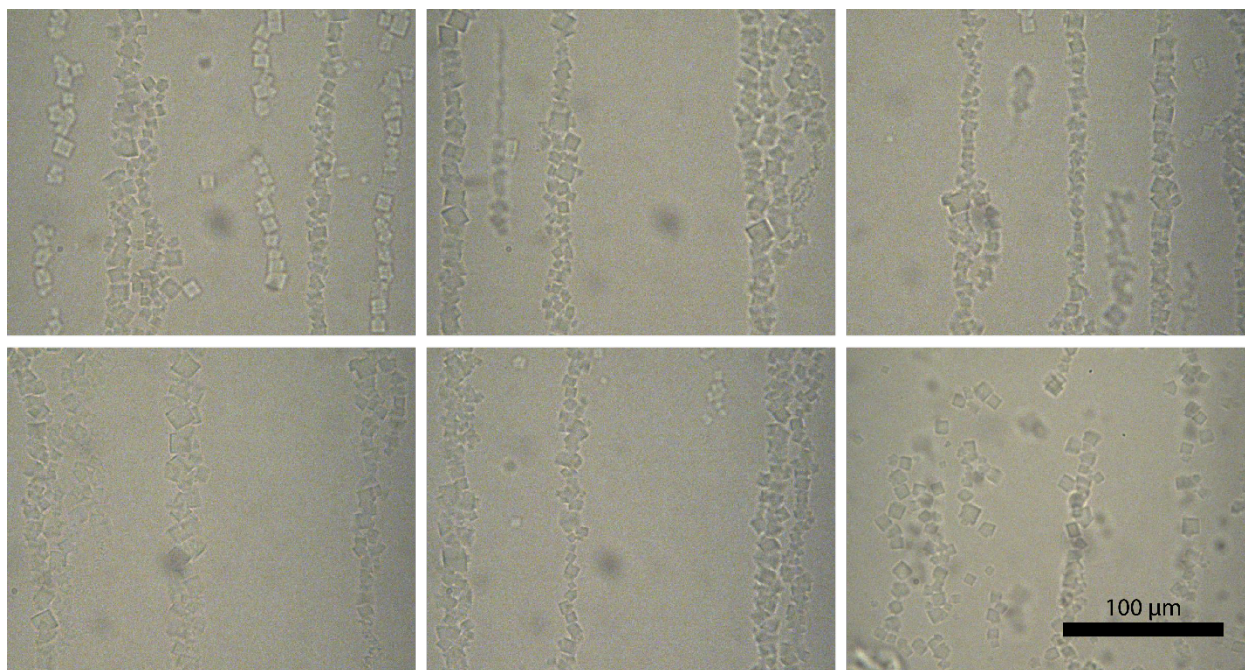
126 **Supplementary Figure 8. a.** Summary of crystal edge lengths vs. τ as calculated from the model and observed
 127 by the experimental data with a 'slow' temperature ramp of 0.01 °C/min. **b.** Histogram of crystal size distribution
 128 with no acoustic waves (blue) and with $\tau = 20$ (red) for thermal ramp down of 'slow' 0.01 °C/min. The nucleation
 129 and growth theoretical fit (blue line) and the infusion model (red line) take into consideration the effect of the
 130 acoustic field.



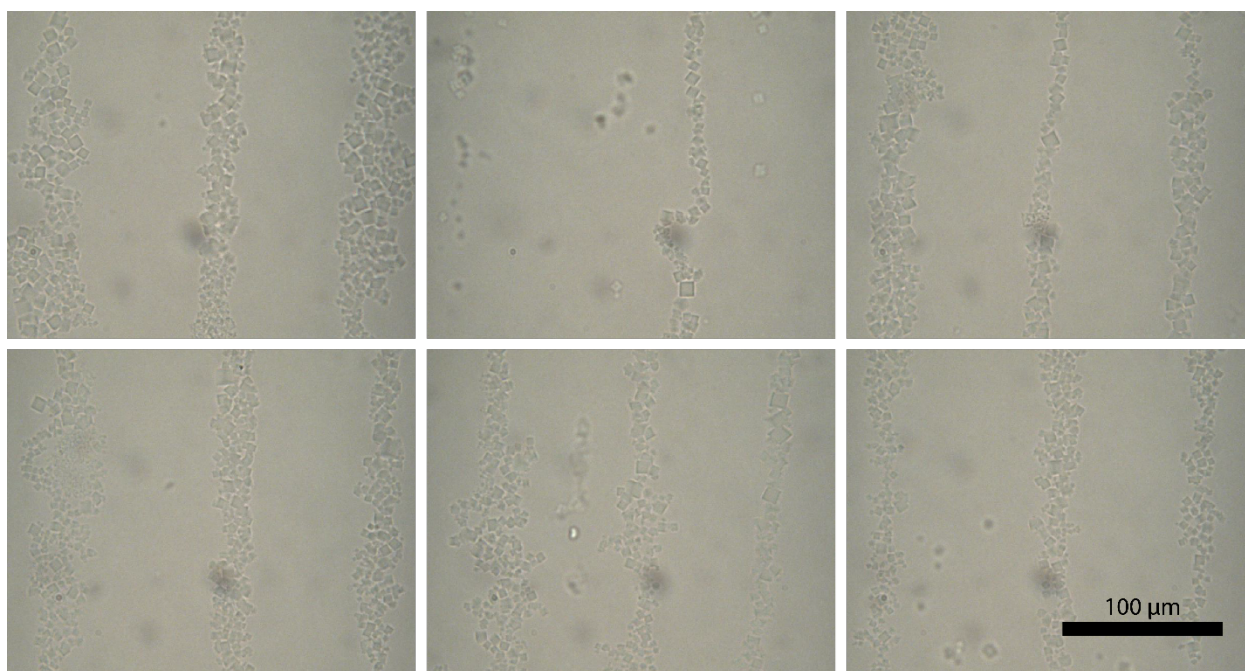
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 132 **Supplementary Figure 9.** Microscopy images of crystals after formation with acoustic wave pulses with a ‘fast’
 133 temperature decrease rate of 0.03 °C/min with an acoustic pulse every 500 ms ($\tau = 10$).



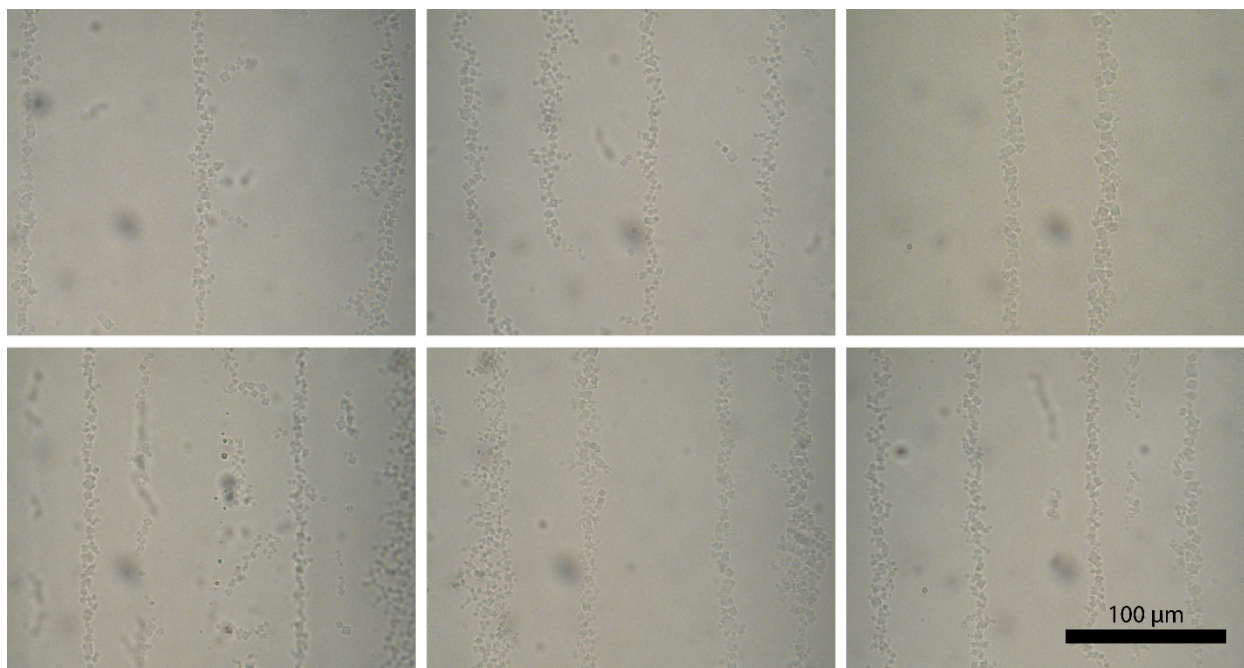
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 135 **Supplementary Figure 10.** Microscopy images of crystals after formation with acoustic pulses with a ‘fast’
 136 temperature decrease rate 0.03 °C/min with an acoustic pulse every 750 ms ($\tau = 15$).



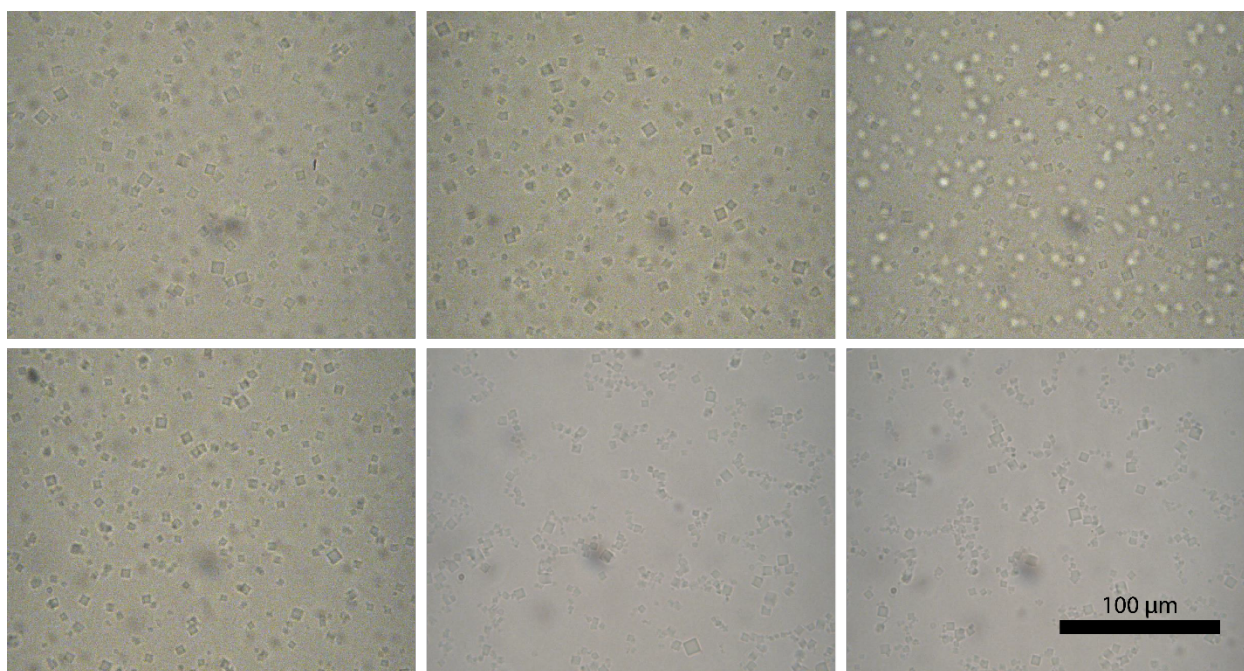
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 138 **Supplementary Figure 11.** Microscopy images of crystals after formation with acoustic wave pulses with a
 139 '*fast*' temperature ramp of 0.03 °C/min with an acoustic pulse every 1000 ms ($\tau = 20$).



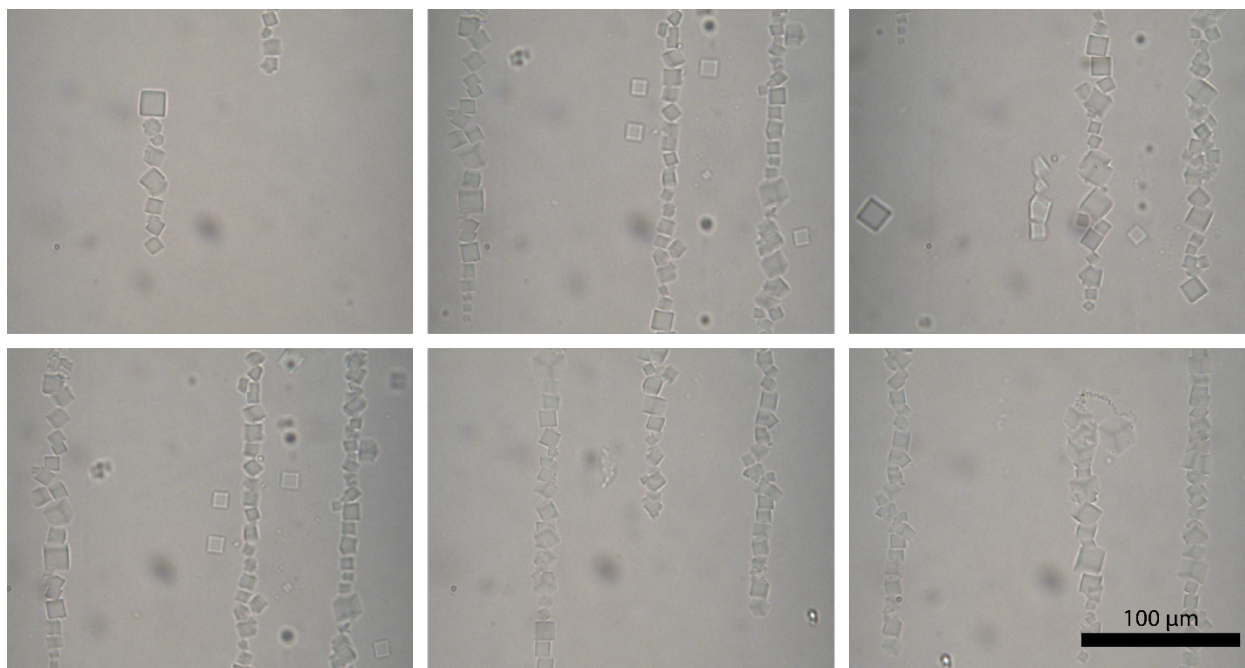
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 141 **Supplementary Figure 12.** Microscopy images of crystals after formation with acoustic wave pulses with a
 142 '*fast*' temperature decrease rate 0.03 °C/min with an acoustic pulse every 1500 ms ($\tau = 30$).



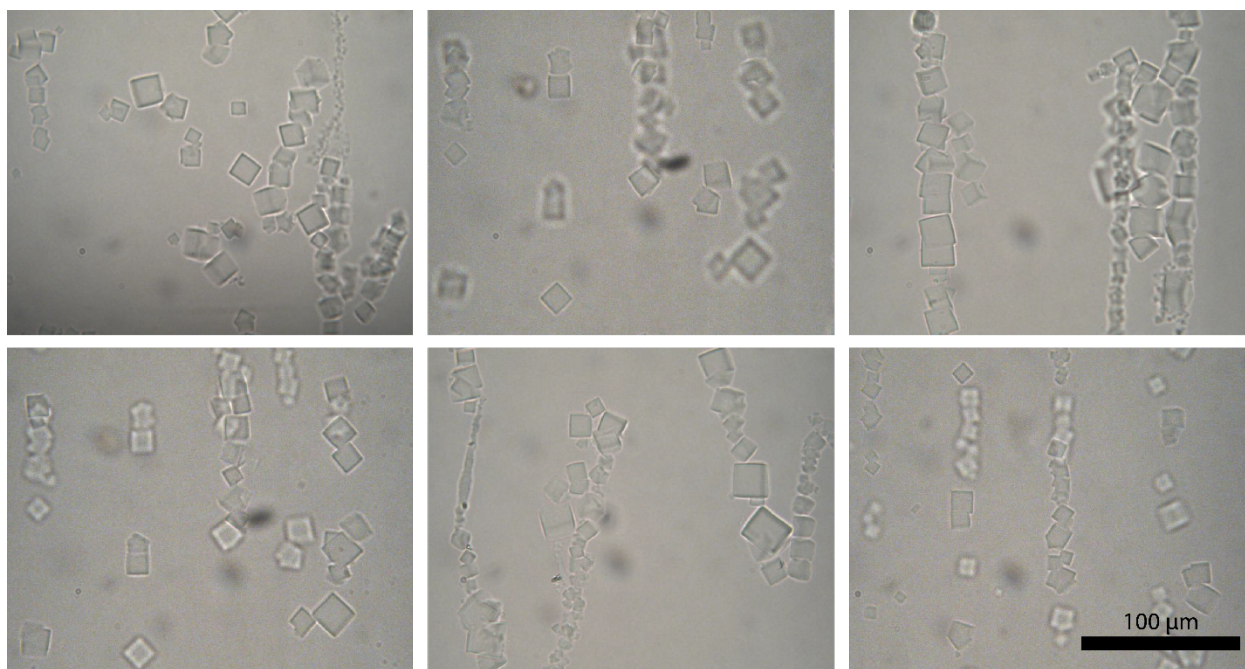
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144 **Supplementary Figure 13.** Microscopy images of crystals after formation with acoustic wave pulses with a
145 '*fast*' temperature decrease rate of 0.03 °C/min with an acoustic pulse every 5000 ms ($\tau = 100$).



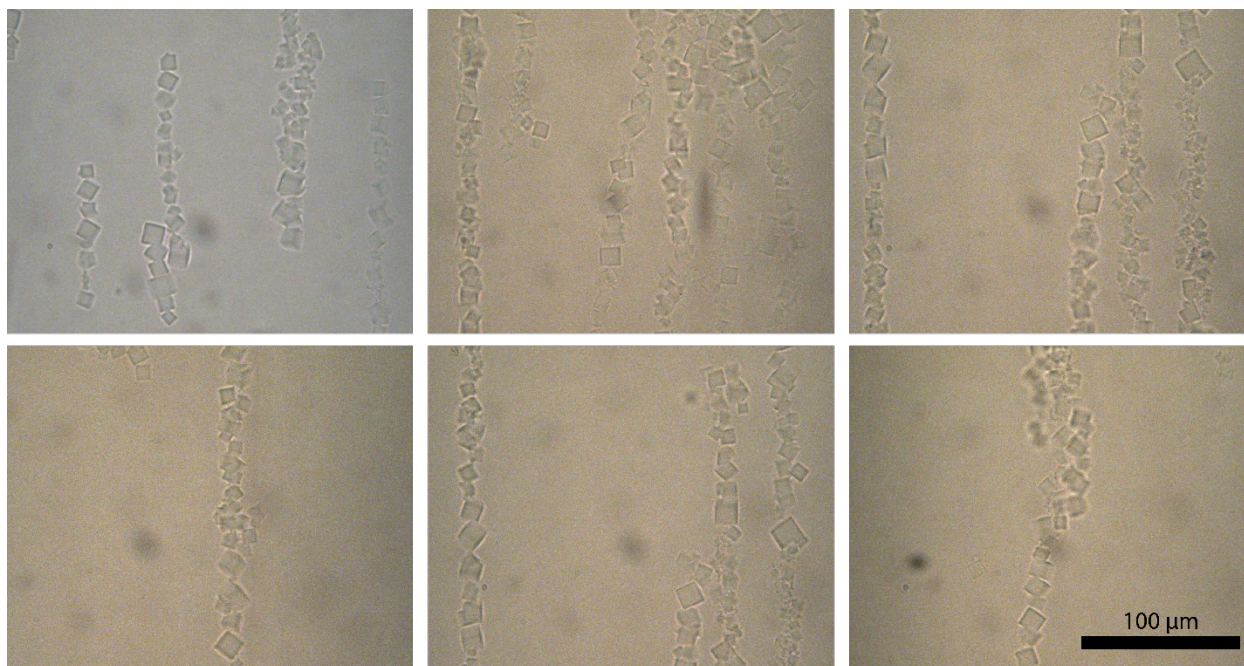
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147 **Supplementary Figure 14.** Microscopy images of crystals after formation with no acoustic wave pulses with a
148 '*fast*' temperature decrease rate of 0.03 °C/min ($\tau = \infty$).



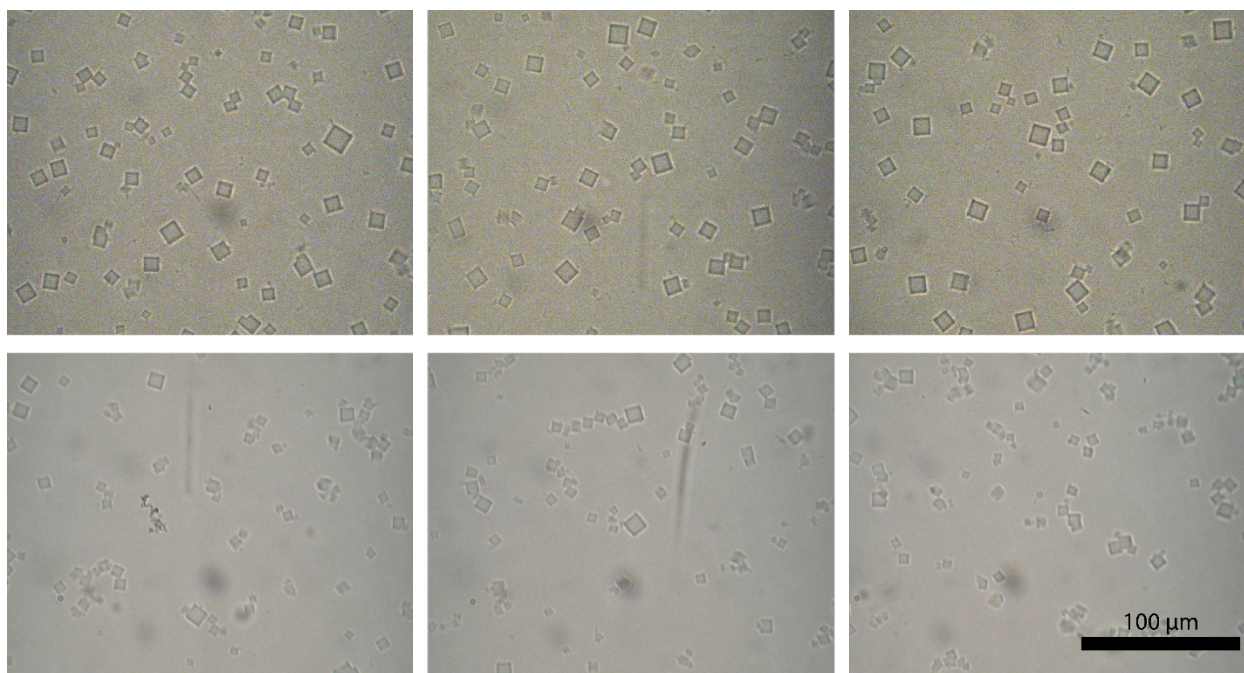
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 150 **Supplementary Figure 15.** Microscopy images of crystals after formation with acoustic wave pulses with a
 151 ‘slow’ temperature decrease rate of 0.01 °C/min with an acoustic pulse every 750 ms ($\tau = 15$).



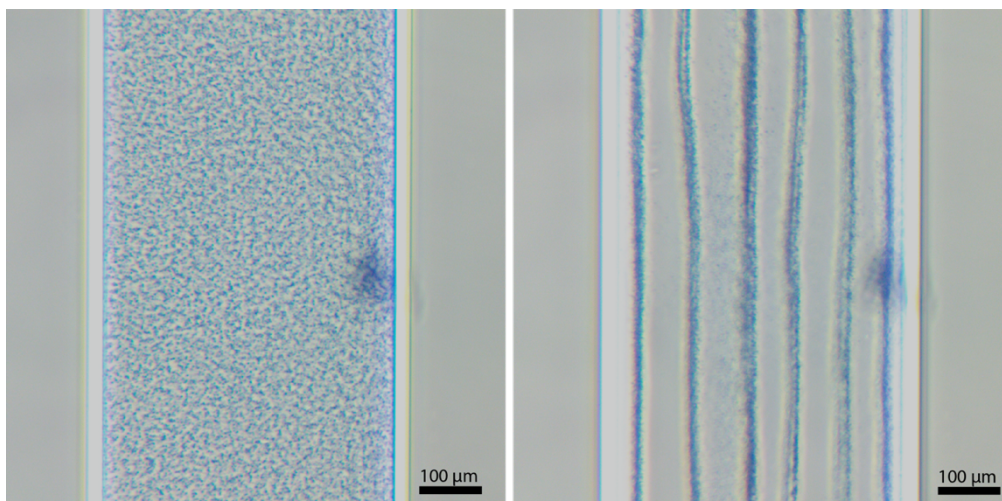
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 153 **Supplementary Figure 16.** Microscopy images of crystals after formation with acoustic wave pulses with a
 154 ‘slow’ temperature decrease rate of 0.01 °C/min with an acoustic pulse every 1000 ms ($\tau = 20$).



155
 156 **Supplementary Figure 17.** Microscopy images of crystals after formation with acoustic wave pulses with a
 157 'slow' temperature decrease rate of 0.01 °C/min with an acoustic pulse every 1500 ms ($\tau = 30$).



158
 159 **Supplementary Figure 18.** Microscopy images of crystals after formation with no acoustic wave pulses with a
 160 'slow' temperature decrease rate of 0.01 °C/min ($\tau = \infty$).



161
162 **Supplementary Figure 19.** Polystyrene beads of 1 μm diameter, in a 1 mm wide capillary. The beads are
163 dispersed in the capillary (left image) and then arranged by a pulse of 50ms acoustic waves every 1 second (τ
164 = 20). The corresponding video can be viewed in Supplementary Movies (Supplementary Movie 5).

165 **Supplementary References**

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