

Supplementary Information

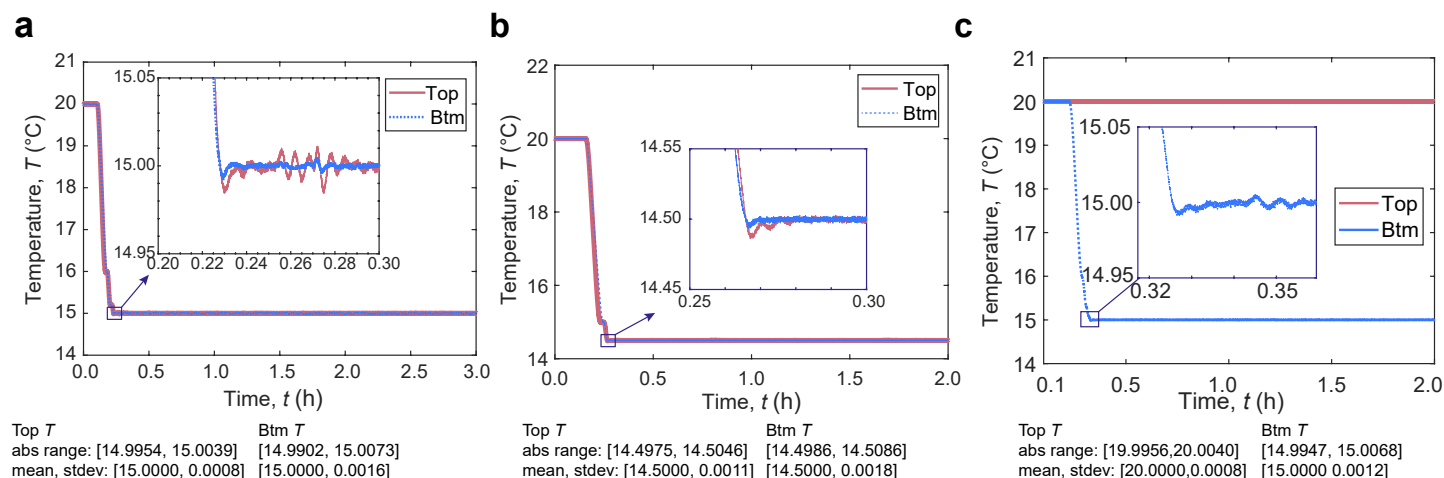
Crystallisation triggered by mass diffusion at a lower local supersaturation

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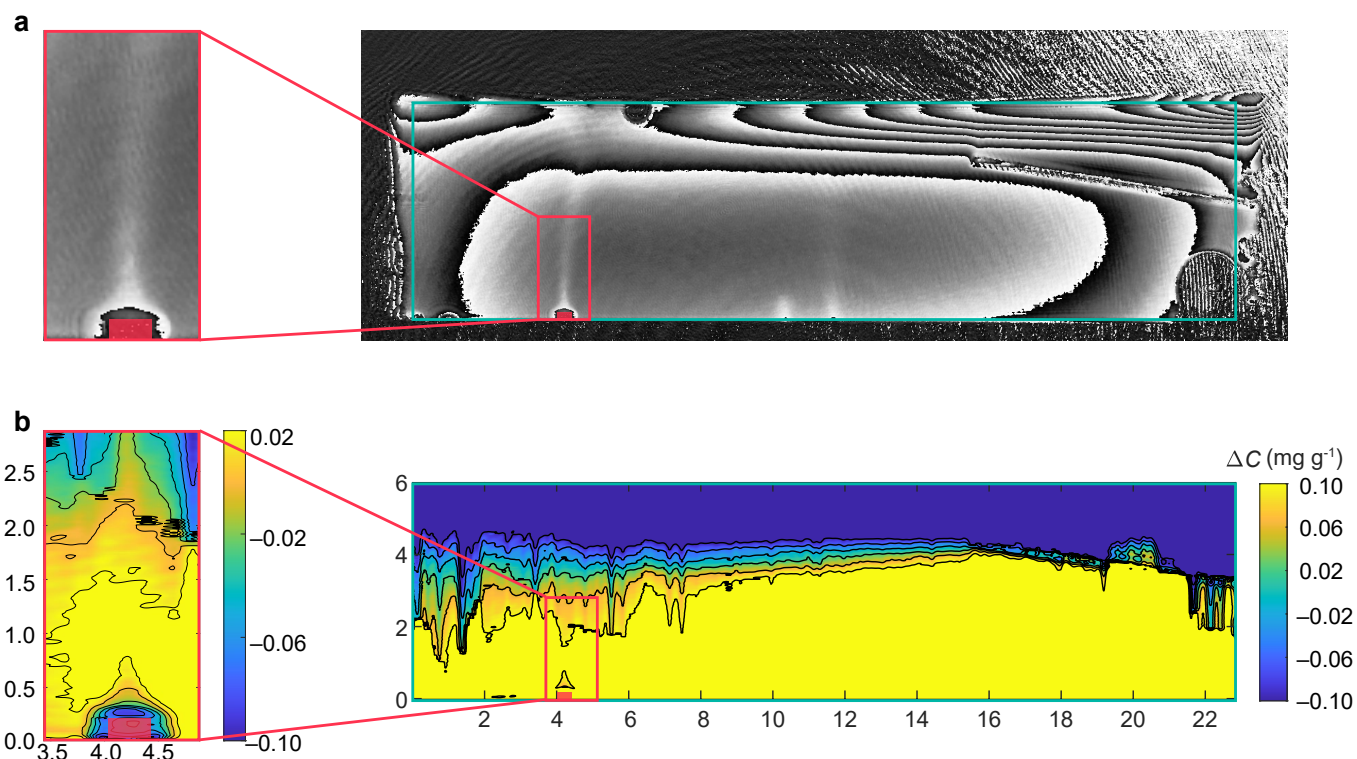
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Supplementary Figures

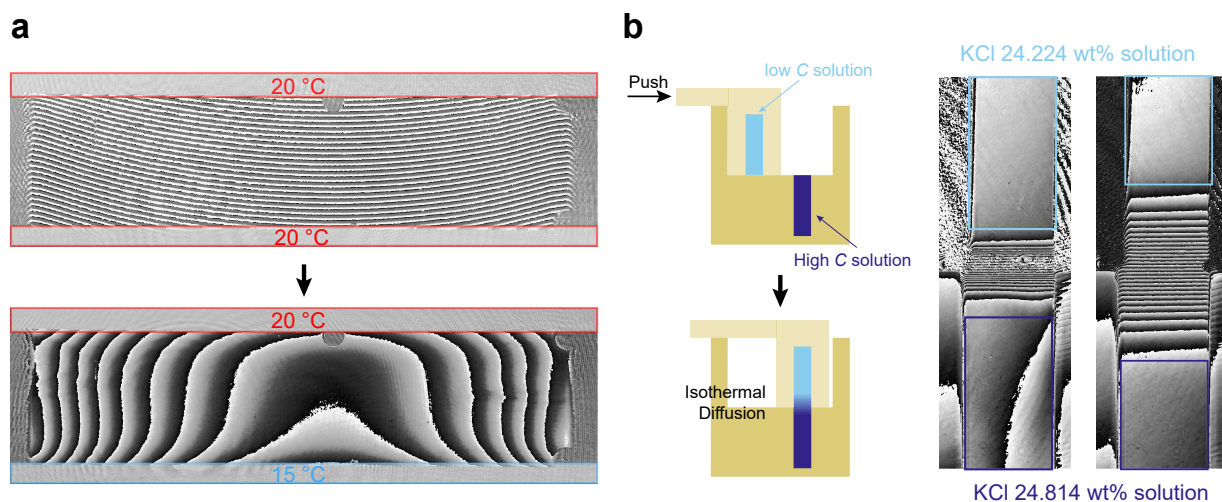
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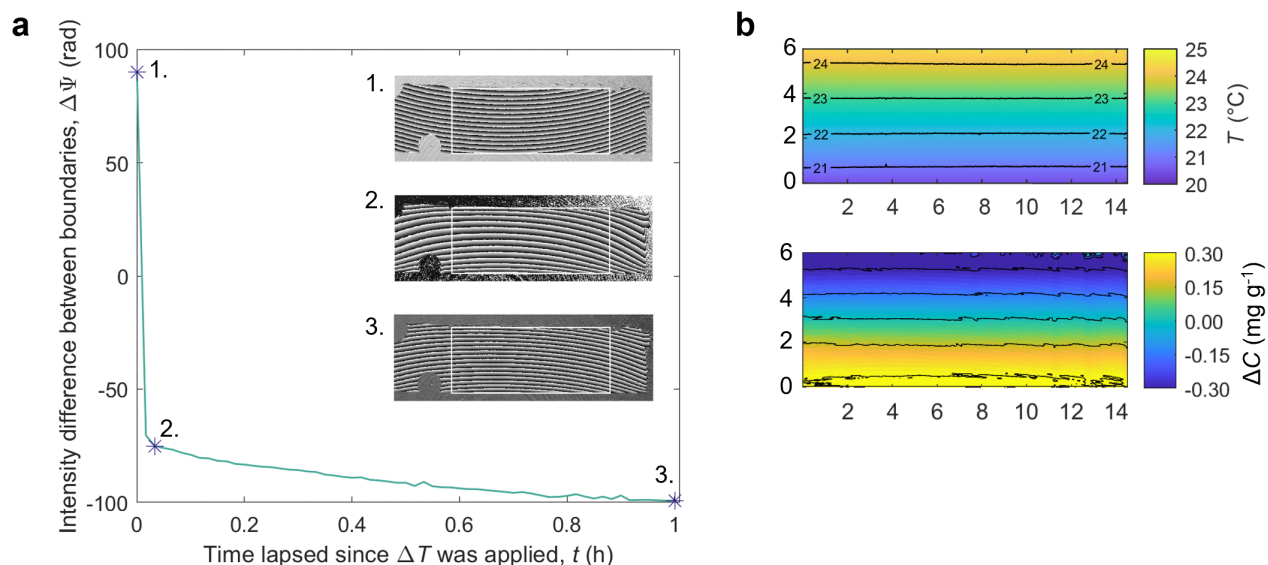
Supplementary Fig. 1 | Temperature control. **a**, Top and bottom boundaries cool from 20 °C to 15 °C. **b**, Top and bottom boundaries cool from 20 °C to 14.5 °C. **c**, Top boundary remains at 20 °C and bottom boundary cools from 20 °C to 15.0 °C.



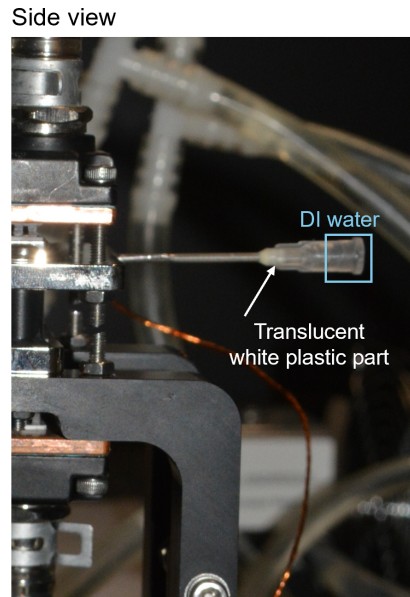
Supplementary Fig. 2 | Cooling crystallisation at 1980 s. **a**, Phase-shifted data, with one growing crystal magnified on the left. **b**, Concentration profile, with the same growing crystal magnified. Solute depletion is evident near the growing crystal, generating a rising plume that reduces the solute concentration in the upper half of the cell during active crystal growth.



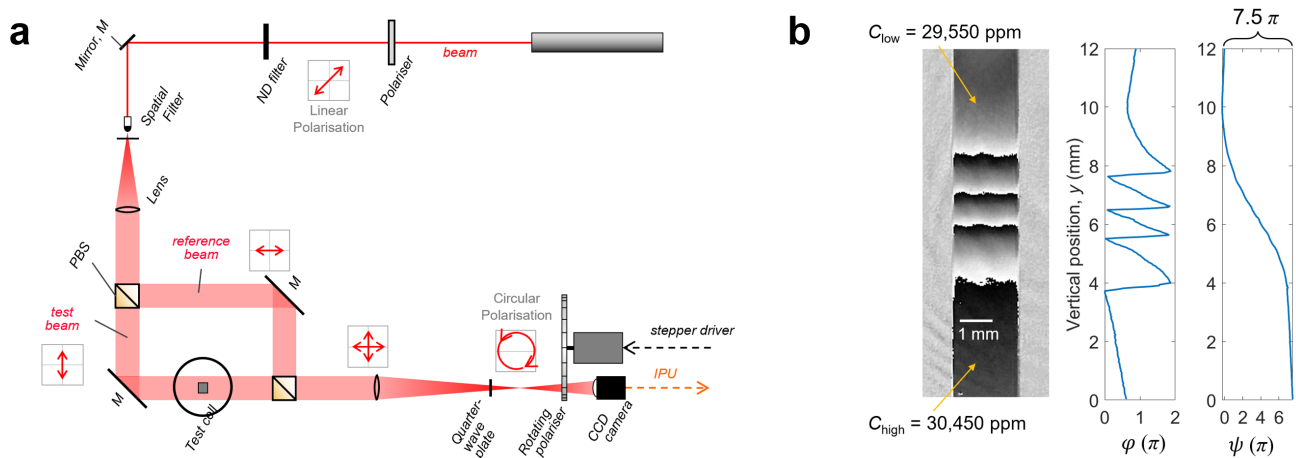
Supplementary Fig. 3 | Contrast factor measurement. **a**, Contrast factor $\delta\Psi/\delta T_{C=C_0}$ is measured by applying a known temperature difference to solution $C = C_0$. The boundary temperatures are measured with thermistors and $\delta\Psi/\delta T_{C=C_0}$ is assumed to be a constant when ΔT is small. Thus $\delta\Psi/\delta T_{C=C_0} = \frac{\Delta\Psi}{OP\Delta T}$ where OP is the optical path length. **b**, Contrast factor $\delta\Psi/\delta C_{T=\text{constant}}$ is measured by performing an isothermal diffusion experiment. The isothermal diffusion experiment was done through sliding the top part containing low C solution onto the bottom part containing the high C solution. Diffusion occurs from the high and low C solution interface. $C_{\text{high}} = 24.814 \text{ wt\%}$ and $C_{\text{low}} = 24.224 \text{ wt\%}$. We assume that the contrast factor at $C_0 = 25.36 \text{ wt\%}$ is the same. From the top to bottom, the concentration increases while the phase Ψ decreases, i.e., each fringe changes from bright to dark. Thus the contrast factor $\delta\Psi/\delta C$ is negative and $\delta\Psi/\delta C_{T=T_0} = \frac{\Delta\Psi}{OP\Delta C}$.



Supplementary Fig. 4 | KCl thermodiffusion without crystallisation. **a**, The unwrapped phase change Ψ as time progresses. The thermodiffusion is nearly constant after 0.8 h. Between 1 and 2 is the temperature development phase and between 2 and 3 is the concentration development phase. **b**, The temperature map is obtained by subtracting unwrapped PSI image in **a1** from **a2** and the concentration difference map is obtained by subtracting unwrapped PSI image in **a2** from **a3**. The maximum ΔC is 0.3097 mg g^{-1} and the minimum ΔC is $-0.3422 \text{ mg g}^{-1}$.



Supplementary Fig. 5 | Evaporation suppression for the crystallisation cell. Photo showing the side view with a glued needle used to suppress evaporation from the solution injection hole. The KCl solution was removed from the white plastic part by a lint-free tissue, and then the DI water was injected slightly forming a seal. The white part is translucent to confirm that there is air between the KCl solution in the needle and the injected DI water. This produced a tight seal of the crystallisation cell.



Supplementary Fig. 6 | Phase-shifting interferometry (PSI). **a**, Layout of the polarising Mach-Zehnder interferometer with rotating polariser, which is the basis for the PSI technique with a temporal phase-shifting algorithm, as described in Torres et al. *Optics and Laser in Engineering* **50**, 1287–1296 (2012). Adapted with permission of Elsevier. **b**, On the left is a typical phase-shifted data, which describes the diffusion field between a low and high concentration solution, C_{high} and C_{low} , respectively. The phase φ and the unwrapped phase ψ are plotted against the cell height on the right. The unwrapped phase map ψ connects discrete values of φ .