

Deep-Tissue Super-Resolution Microscopy using Single-Scattering Accumulation Algorithm



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Abstract

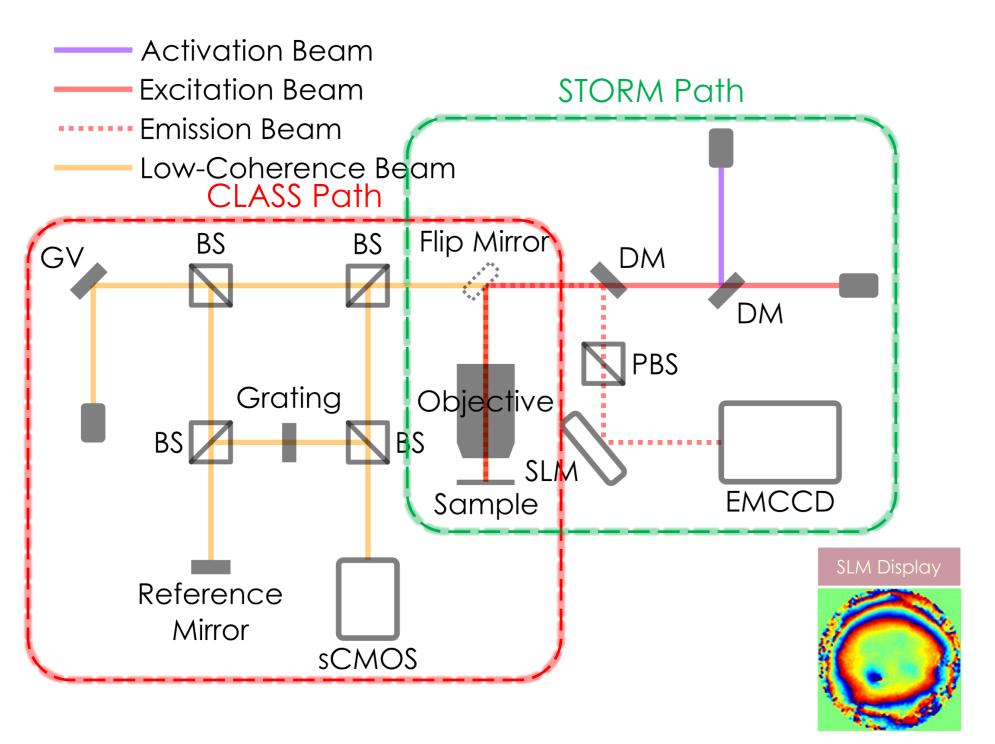
Single-molecule localization microscopy has enabled the resolving of biomolecules well beyond the diffraction-limited spatial resolution. However, its imaging depth has been too shallow to investigate thick biological tissues. A sample-induced aberration is one of the major reasons for this limitation. It gives rise to the blur and distortion of single-molecule emission images, impairing the extremely sensitive single-molecule localization process. To resolve this issue, we combined the closed-loop accumulation of single scattering (CLASS) microscopy with stochastic optical reconstruction microscopy (STORM). CLASS microscopy identifies the sample-induced aberration based on elastic backscattering, which is then applied to a spatial light modulator in the excitation beam path of STORM for aberration. We demonstrated super-resolution imaging of cell microtubules under artificially generated severe aberration consisting of Zernike modes higher than mode 20 and the STORM imaging of neurons in 100µm-thick mouse brain tissue at the depth of \sim 70µm.

Concept

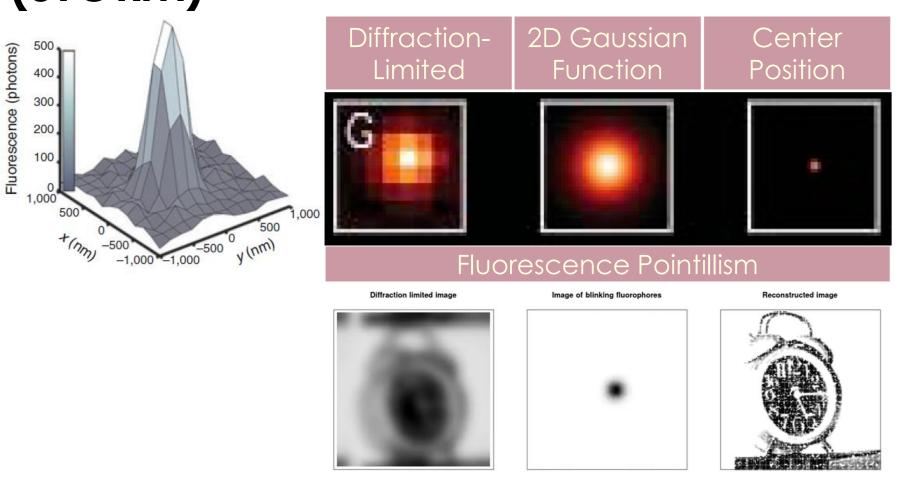
Adaptive optics-assisted super-resolution microscopy

- CLASS (closed-loop accumulation of single scattering) + STORM (stochastic optical reconstruction microscopy)
- Correcting CLASS-calculated sample-induced aberration ▶ obtaining deep-tissue STORM images
- The goal is to obtain STORM images of neurons in a mouse brain or an intact animal (e.g., zebrafish) at the depth of ~100µm

Principle



Stochastic optical reconstruction microscopy $(STORM)^{1,2}$



Collective accumulation of single-scattered waves (CASS) & Closed-loop accumulation of single scattering (CLASS)^{3,4}

 $E(\mathbf{r}; \mathbf{k}_n^{in}, \tau_0) = E_s(\mathbf{r}) + E_R(\mathbf{r})e^{-i\mathbf{k}_n^{in}\cdot\mathbf{r}}$ [interference of two beams (E-field)] where $\mathbf{r} = (x, y)$ (coordinates in camera-detected image) and \mathbf{k}_n^{in} representing nth scan angle

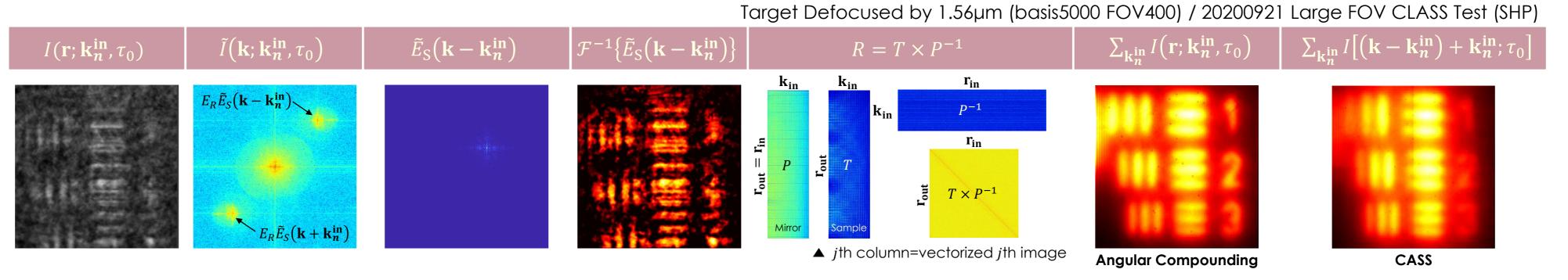
 $I(\mathbf{r}; \mathbf{k}_n^{\text{in}}, \tau_0) = \left| E(\mathbf{r}; \mathbf{k}_n^{\text{in}}, \tau_0) \right|^2 = |E_s(\mathbf{r})|^2 + |E_R(\mathbf{r})|^2 + E_S(\mathbf{r})E_R(\mathbf{r}) \left(e^{i\mathbf{k}_n^{\text{in}}\cdot\mathbf{r}} + e^{-i\mathbf{k}_n^{\text{in}}\cdot\mathbf{r}} \right) \text{ [camera-detected image (intensity)]}$ Taking Fourier transform and using the convolution theorem,

 $\mathcal{F}\{|E_{S}(\mathbf{r})|^{2} + |E_{R}(\mathbf{r})|^{2}\} = [|E_{S}(\mathbf{r})|^{2} + |E_{R}(\mathbf{r})|^{2}]\mathcal{F}\{1\} = [|E_{S}(\mathbf{r})|^{2} + |E_{R}(\mathbf{r})|^{2}]\delta(\mathbf{k})$

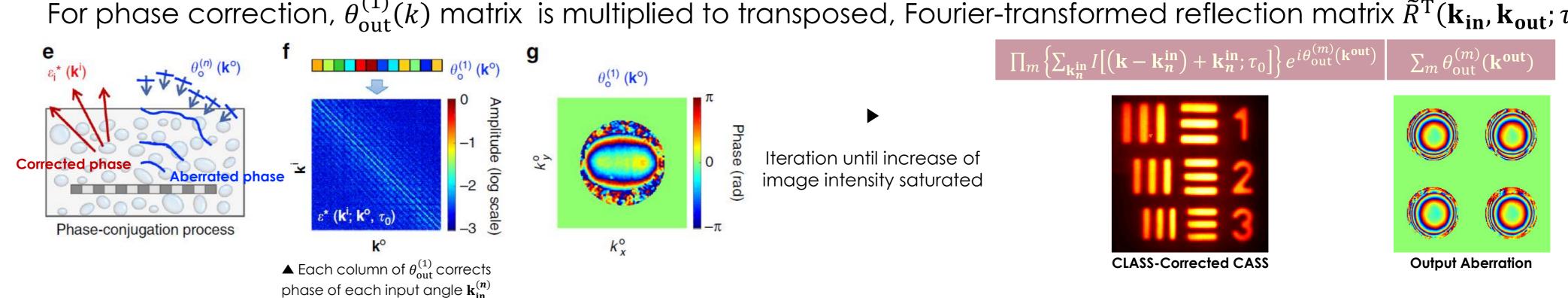
 $\mathcal{F}\left\{E_{S}(\mathbf{r})\left[E_{R}(\mathbf{r})e^{\pm i\mathbf{k}_{n}^{\text{in}}\cdot\mathbf{r}}\right]\right\} = \tilde{E}_{R}(\mathbf{k})\iint d^{2}\mathbf{r}\left[E_{S}(\mathbf{r})e^{\pm i\mathbf{k}_{n}^{\text{in}}\cdot\mathbf{r}}\right]e^{-i\mathbf{k}\cdot\mathbf{r}} = \tilde{E}_{R}(\mathbf{k})\iint d^{2}\mathbf{r}\left[E_{S}(\mathbf{r})e^{-i\left(\mathbf{k}\mp\mathbf{k}_{n}^{\text{in}}\right)\cdot\mathbf{r}}\right] = \tilde{E}_{R}(\mathbf{k})\tilde{E}_{S}(\mathbf{k}\mp\mathbf{k}_{n}^{\text{in}})$

Then,

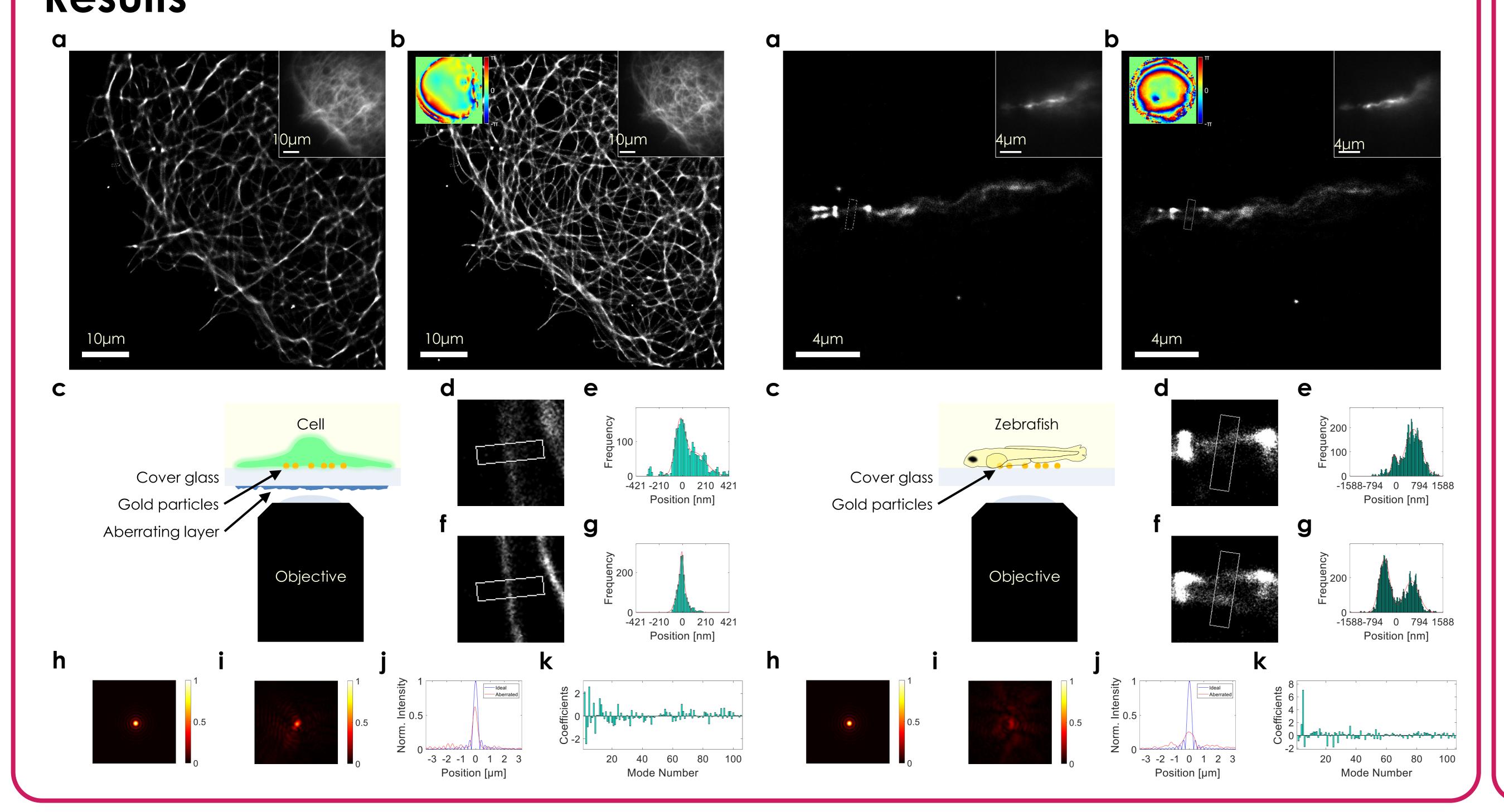
 $\mathcal{F}\{I(\mathbf{r};\mathbf{k}_n^{\text{in}},\tau_0)\} \equiv \tilde{I}(\mathbf{k};\mathbf{k}_n^{\text{in}},\tau_0) = [|E_S(\mathbf{r})|^2 + |E_R(\mathbf{r})|^2]\delta(\mathbf{k}) + \tilde{E}_R(\mathbf{k})[\tilde{E}_S(\mathbf{k} - \mathbf{k}_n^{\text{in}}) + \tilde{E}_S(\mathbf{k} + \mathbf{k}_n^{\text{in}})] \text{ (Fourier-transformed image)}$ where $\mathbf{k} = (k_x, k_y)$



For phase correction, $\theta_{\text{out}}^{(1)}(k)$ matrix is multiplied to transposed, Fourier-transformed reflection matrix $\tilde{R}^{\text{T}}(\mathbf{k_{in}}, \mathbf{k_{out}}; \tau_0)$



Results



References

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- 2. E. Betzig et al., Science 313, 5793 (2006)
- 3. Q-H. Park and W. Choi et al., Nature Photonics 9, 4 (2015)
- 4. K. H. Kim & W. Choi et al., Nature Communications 8, 2157 (2017)