

COMPARING METRICS FOR INTENSITY-BASED ADAPTIVE OPTICS IN SINGLE-MOLECULE LOCALIZATION MICROSCOPY

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Recent developments in excitation modalities and labeling strategies enable single-molecule localization microscopy in deep tissue (>20 μm). However, sample-induced aberrations hamper optimal detection and localization, which can be overcome by intensity-based adaptive optics (AO). The need for fiducial markers or ‘guide stars’ in complex samples can be circumvented by using the single-molecule acquisitions themselves in combination with an appropriate metric, a measure for the quality of an acquisition. Here we compare the performance of different metrics suggested in literature [1-3] and a newly proposed metric by simulating single-molecule acquisitions in a wide variety of conditions using model-based optimization. This reveals that correction is challenging and that straightforward implementations do not guarantee improvement. We show that a series of 300 acquisitions with a combination of two metrics can robustly correct initial aberrations up to 1 rad rms below 0.3 rad rms (0.9 Strehl-ratio), restoring localization precision. The achieved aberration correction is experimentally verified on a DNA-PAINT sample by adding known aberrations that are subsequently corrected. These results reveal under which SNR conditions and aberration levels robust correction can be achieved.

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