

Supplemental Figures

Temporal dependence of shifts in mu opioid receptor mobility at the cell surface after agonist binding observed by single-particle tracking

Marissa J. Metz^{1,#}, Reagan L. Pennock^{1,+,#}, Diego Krapf^{2,3,*}, Shane T. Hentges^{1,*}

¹Department of Biomedical Sciences, Colorado State University, Fort Collins, CO, USA.

²Department of Electrical and Computer Engineering, Colorado State University, Fort Collins, CO, USA.

³School of Biomedical Engineering, Colorado State University, Fort Collins, CO, USA.

[†]Current address: Department of Neurobiology, University of Alabama at Birmingham, Birmingham, AL, USA.

*Corresponding authors: S.T.H., Colorado State University, 1617 Campus Delivery, Fort Collins, CO 80521, USA. hentgess@colostate.edu D.K., Colorado State University, 1301 Campus Delivery, Fort Collins, CO 80523, USA. diego.krapf@colostate.edu

^{#,*}These authors contributed equally to the work

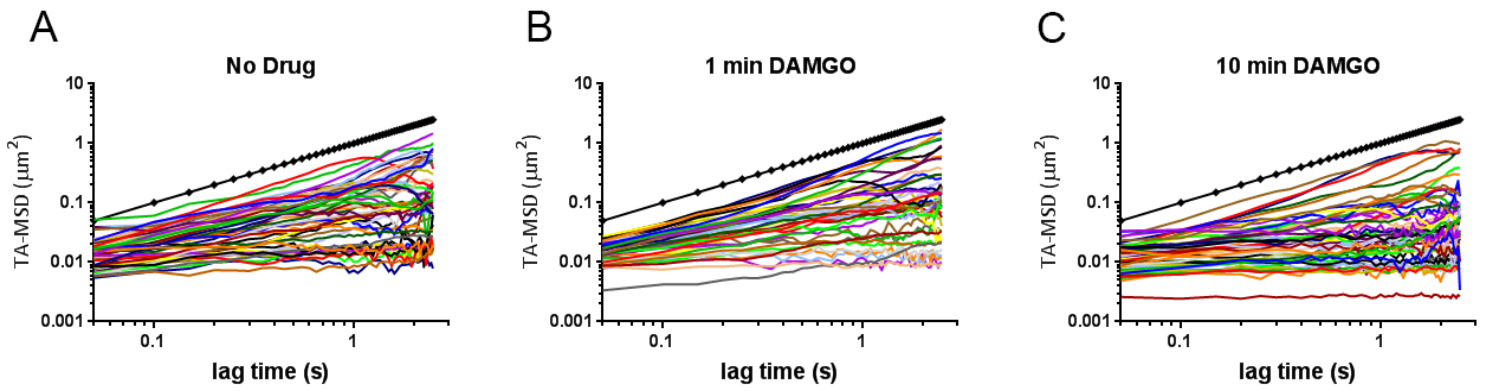


Figure S1. Individual MSDs of randomly chosen trajectories. Log-log plots of time averaged MSDs from individual tracks in the **A)** no drug, **B)** 1 min DAMGO, and **C)** 10 min DAMGO experimental conditions. Most trajectories are subdiffusive, as they lie below the slope of a simulated track of $\alpha = 1$ (black dotted line).

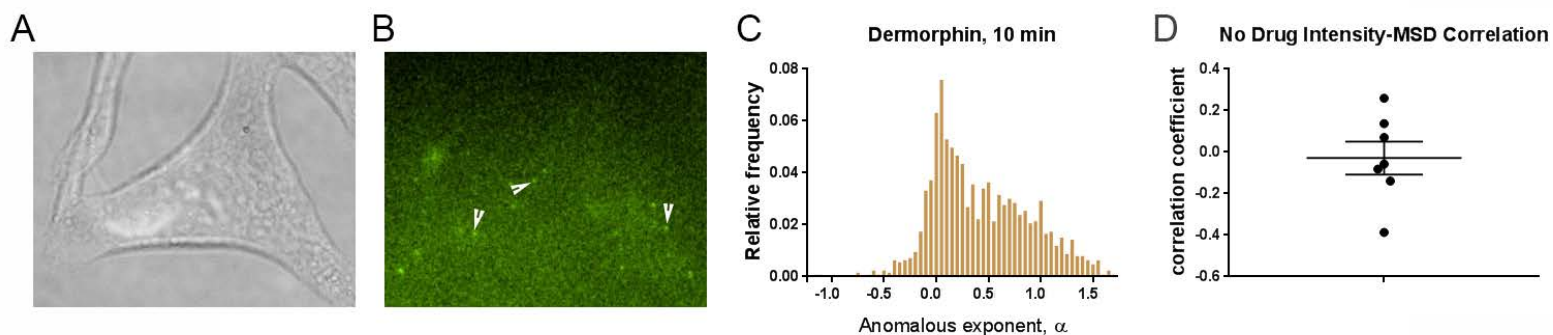


Figure S2. Single particle tracking after 10 min of Dermorphin-488 (60 pM) application reveals that the immobile population of MORs is not due to antibody-mediated crosslinking or Qdot hindering of mobility. A.) DIC image of an AtT20 cell labeled with Dermorphin-488. **B.)** The same cell is shown under fluorescence, and arrows indicate MORs labeled with a single Dermorphin-488 conjugate. **C.)** Distribution of α values after tracking of MOR-Dermorphin-488 conjugates ($n = 5$ cells, 1266 tracks) incubated for 10 min in the presence of 60 pM Dermorphin-488. The distribution of α values is similar to those observed with Qdot tracking, and the fraction of $\alpha < 0.27$ is 0.44 ± 0.08 , similar to MOR-Qdots in the 10 min DAMGO condition (0.45 ± 0.12). Values below 0 are likely due to errors made during tracking caused by the low signal to noise ratio with this labeling approach. **D.)** Distribution of MSD vs. fluorescence intensity correlation coefficients for individual cells in the no drug condition, with a mean correlation coefficient of -0.03 ± 0.08 .

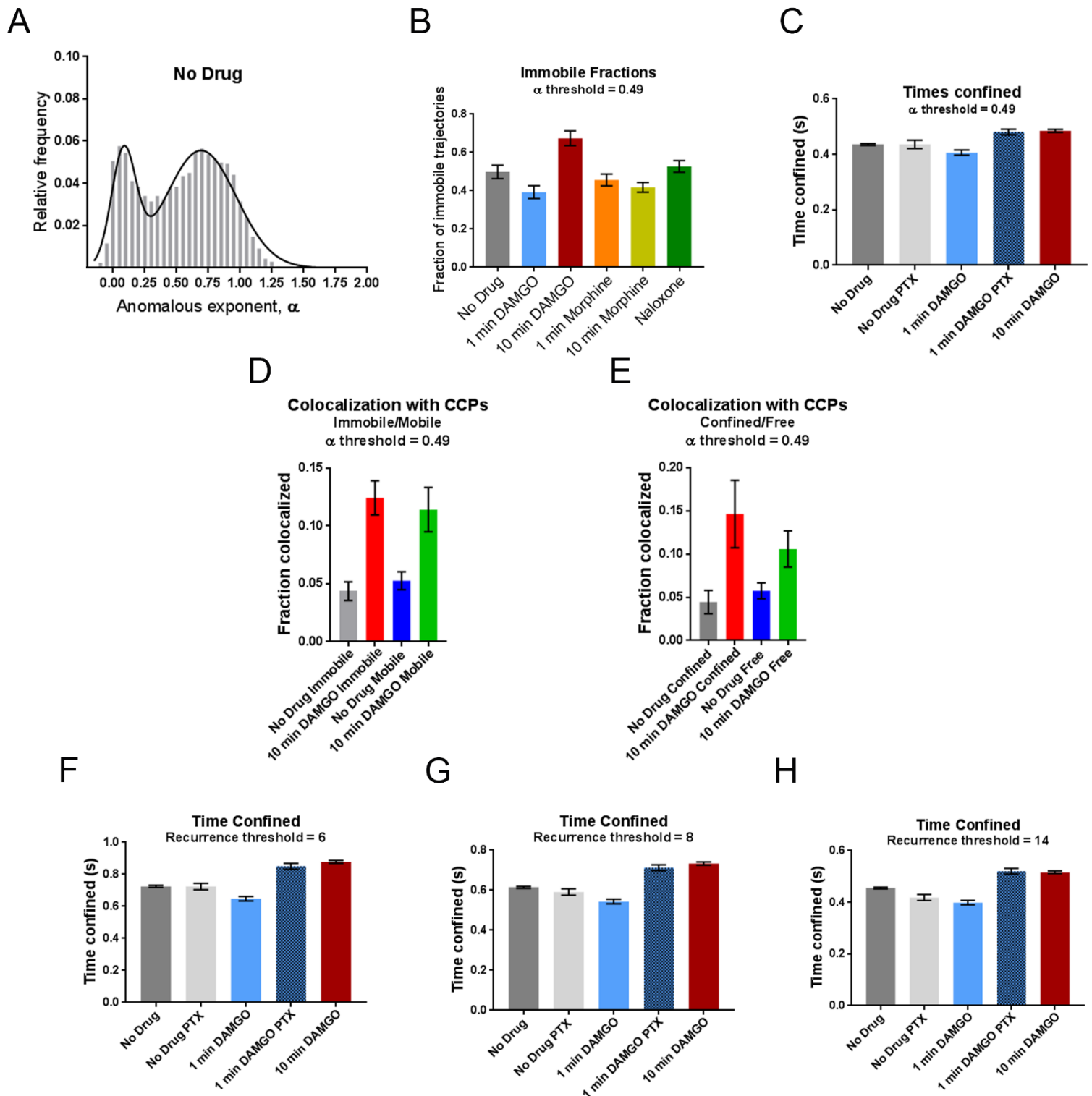


Figure S3. Reported results are robust to changes in mobile/immobile and confined/free thresholds.

A) Two Gaussian fitting results in a threshold near our chosen threshold of $\alpha = 0.27$. **B)** The pattern for fraction of immobile trajectories does not change when using the threshold of $\alpha = 0.49$ determined

using k-means for most experimental conditions, with the exception of 10 min Morphine. **C)** The pattern of confined times does not change at an $\alpha = 0.49$ threshold. **D)** The pattern of colocalization with CCPs does not change when comparing immobile and mobile trajectories or **E)** confined and free trajectories. **F)** Changing the recurrence threshold during recurrence analysis does not change the time confined when the recurrence threshold is 6, **G)** 8, **H)** or 14.

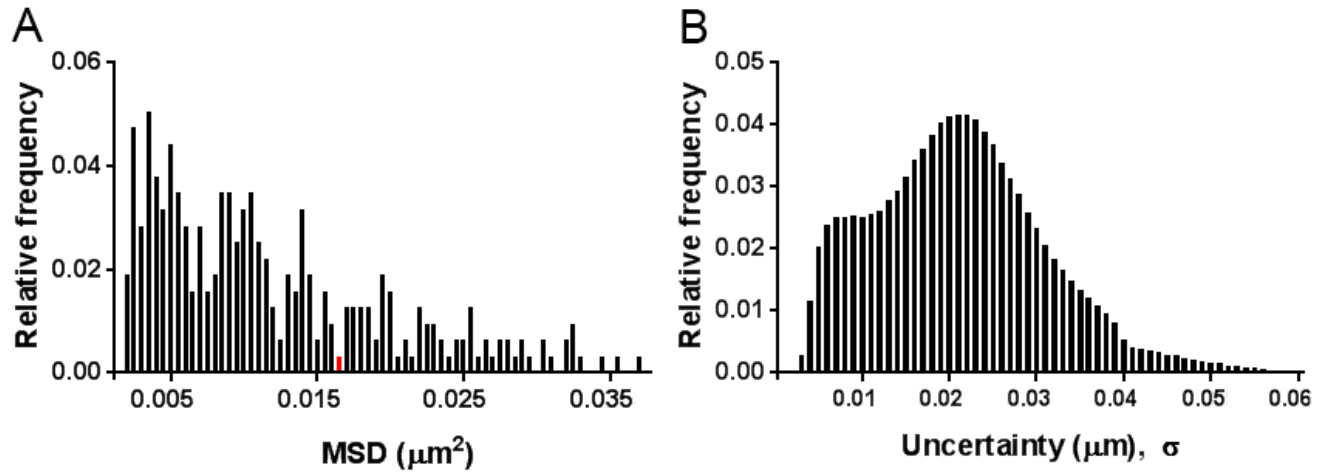


Figure S4. Data for trajectory corrections. A) Frequency histogram of MSD for glass-stuck Qdot-655 tracked without cells on the coverslip. The cutoff of $0.0165 \mu\text{m}^2$ is highlighted in red. Most MSDs are less than this cutoff. **B)** Histogram of localization uncertainties, σ , for each detected particle of a representative subset in the no drug condition. An average σ of $0.02 \mu\text{m}$ was used for uncertainty correction of tracks.