

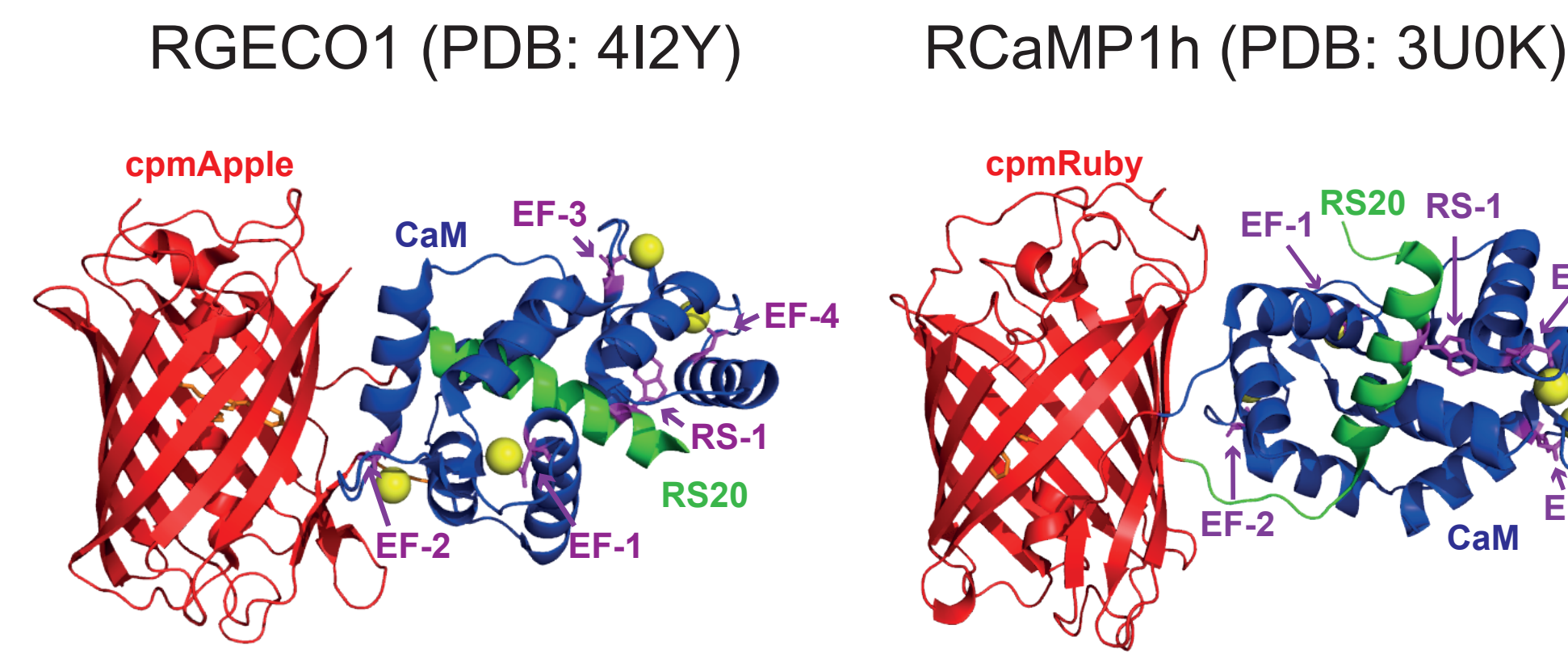
Fast-decay variants of red-fluorescent genetically-encoded Ca^{2+} sensors

S. Kerruth, C. Coates, K. Török

St. George's, University of London, Molecular and Clinical Sciences Research Institute, Cranmer Terrace, SW17 0RE, London, UK

Why make fast red probes?

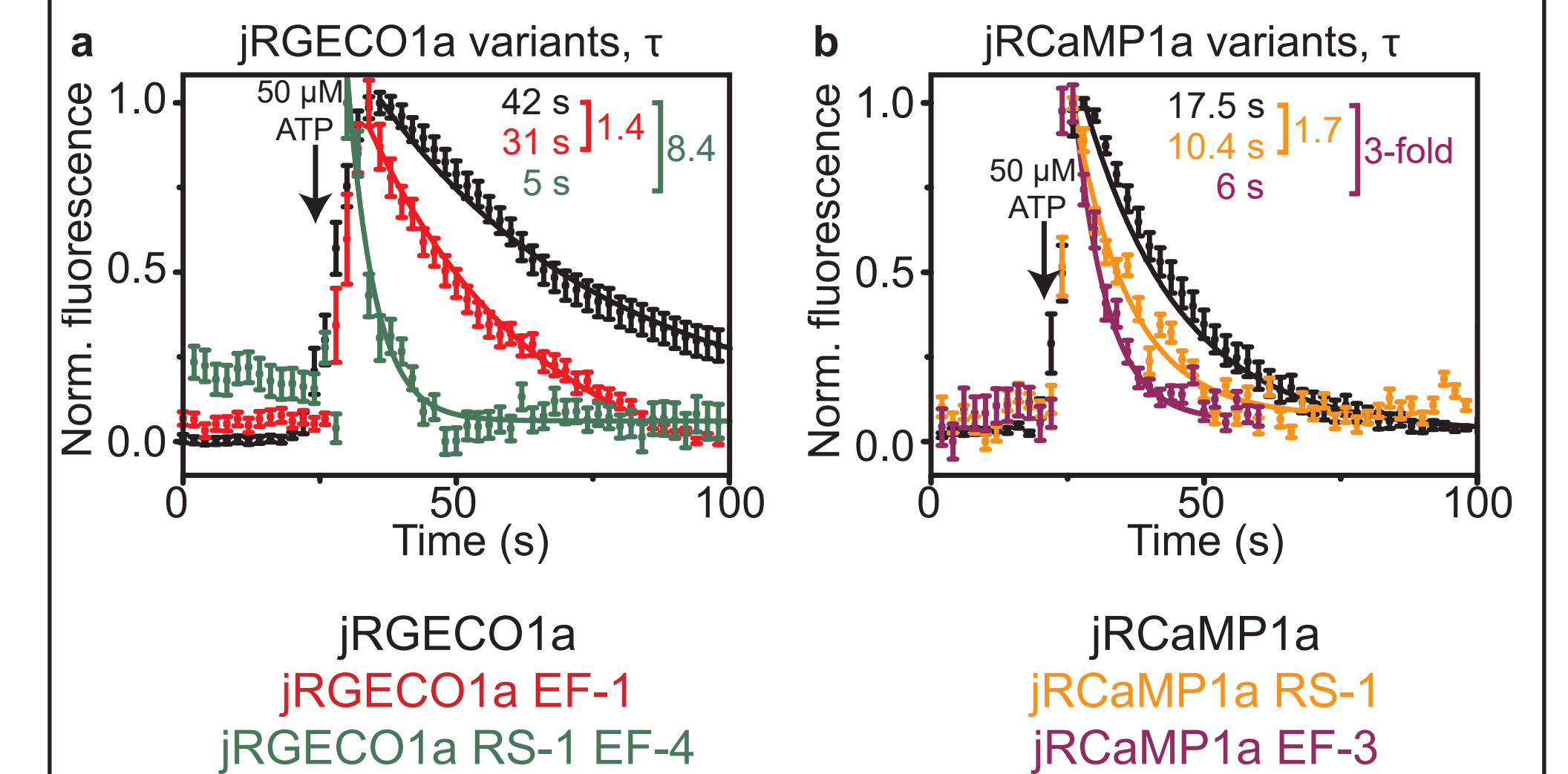
- Why red?** Red-fluorescent genetically-encoded Ca^{2+} indicators (RGECIs) monitor Ca^{2+} transients while cells can be simultaneously excited with blue-light for optogenetics.
- Challenge:** Current RGECIs^[1] have slow response, faster variants needed to monitor action potential firing
- Strategy:** To weaken the CaM and RS20 peptide interaction by mutating the RS20 peptide and disabling the CaM EF hands of jRGECO1a and jRCaMP1a^[2,3].



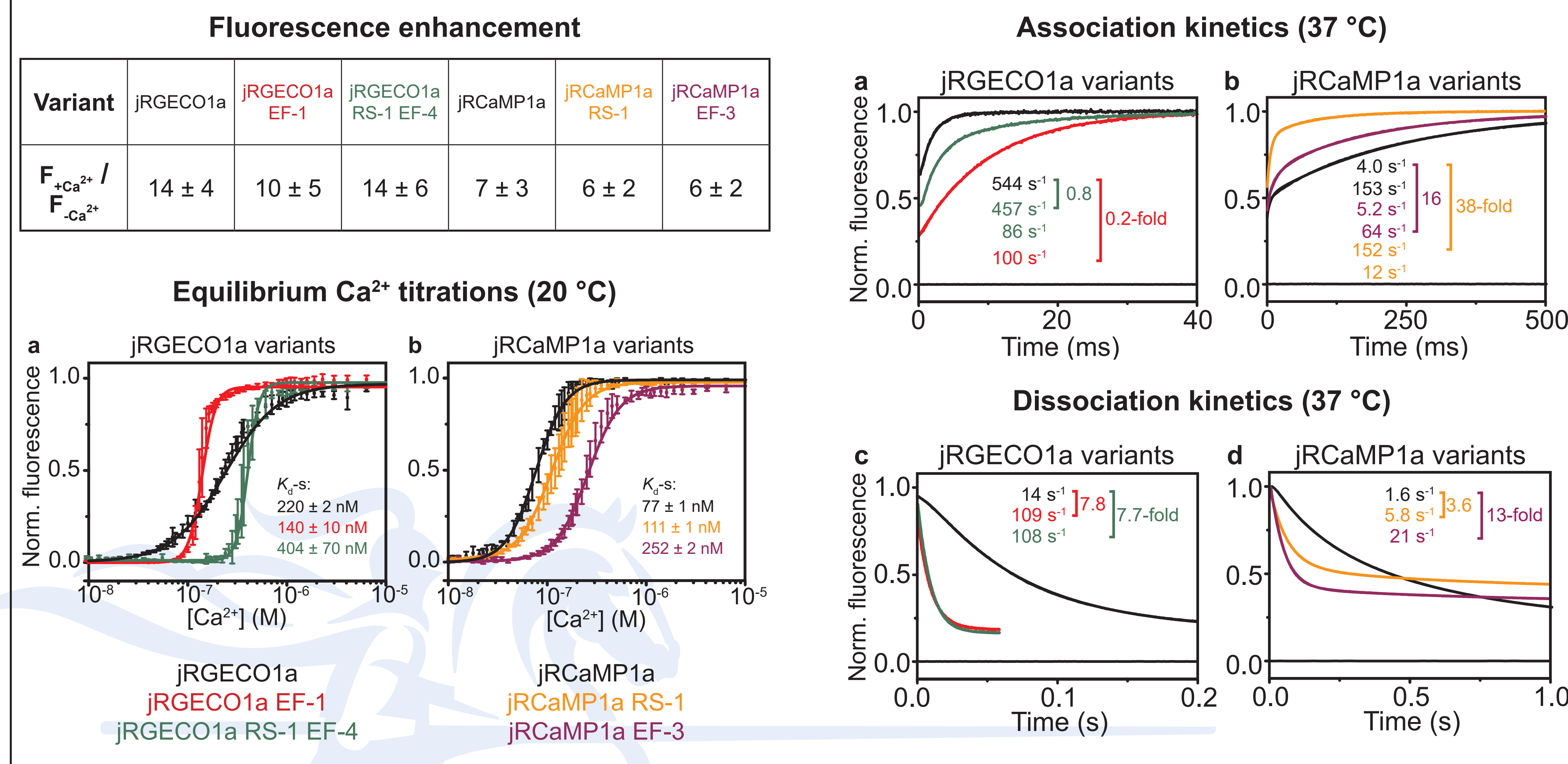
Summary

- Faster RGECI variants are developed without significant loss of brightness and fluorescence dynamic range
- jRCaMP1a RS-1** compared to jRCaMP1a: Limiting association rate 38-fold faster
- jRGECO1a RS-1 EF-4** and **jRCaMP1a RS-1** compared to jRGECO1a and jRCaMP1a:
 - Dissociation kinetics 8-fold and 13-fold faster, respectively
 - ATP-induced cell Ca^{2+} transient decay time courses are 8-fold and 3-fold faster, respectively
- Rate constants are fitted to two-step model for probes which show rapid binding followed by limiting isomerisation

Cellular Ca^{2+} dynamics



Biophysical characterization and kinetics



Ca^{2+} dependence of association kinetics

Observed rates increase with increasing $[\text{Ca}^{2+}]$ and showing sigmoidal dependence.

The k_{obs} were fitted to a model, with an equilibrium Ca^{2+} binding followed by an isomerization to form a fluorescent state.

(d) Irreversible isomerisation:

$$k_{\text{obs}} = \frac{k_2 K_1 [\text{Ca}^{2+}]^n}{1 + K_1 [\text{Ca}^{2+}]^n}$$

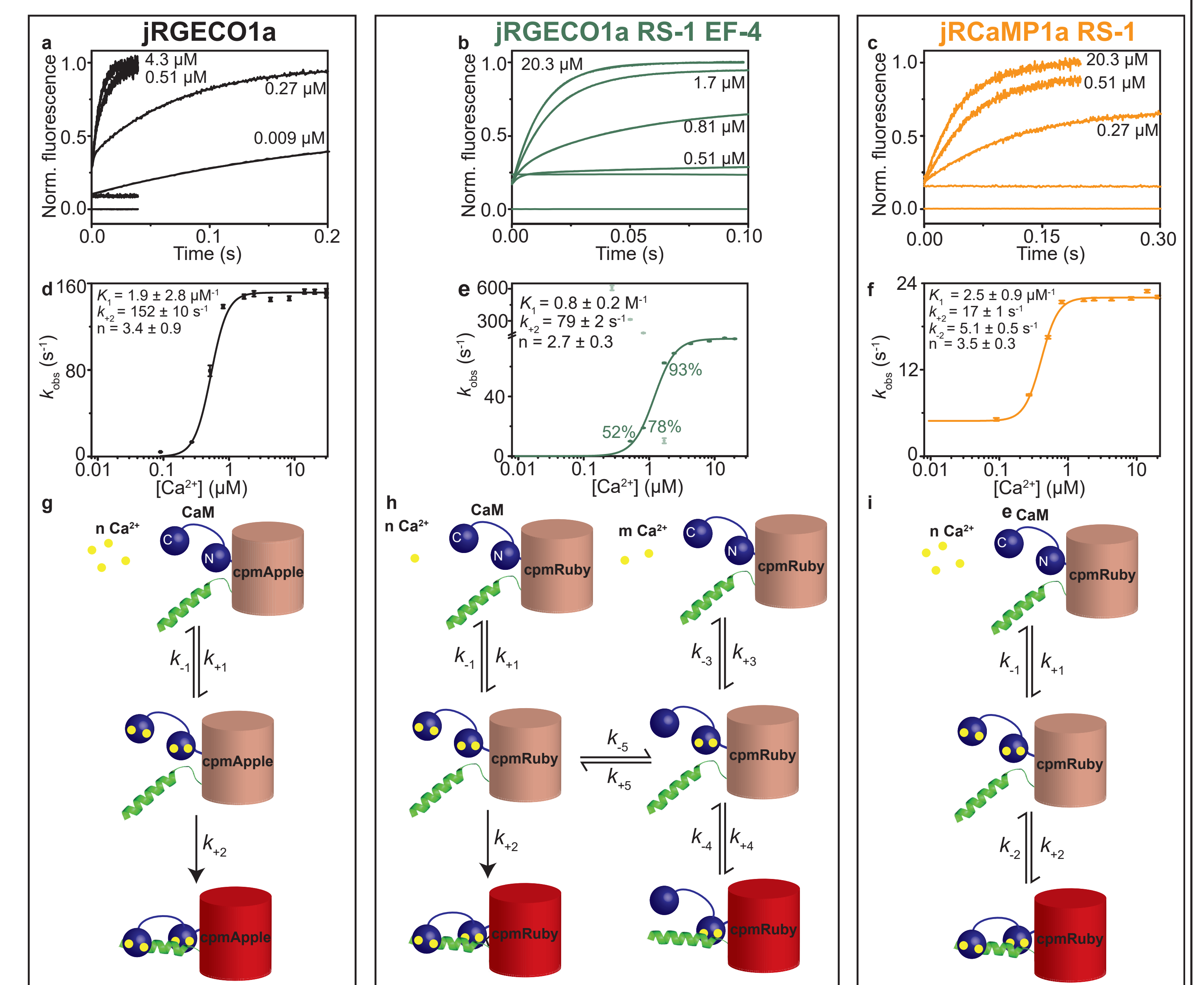
(e) Complex reaction with two fluorescent paths

$$k_{\text{obs}} = \frac{k_2 K_1 [\text{Ca}^{2+}]^n}{1 + K_1 [\text{Ca}^{2+}]^n}$$

$$k_{\text{obs}} = \frac{k_4 K_3 [\text{Ca}^{2+}]^m}{1 + K_3 [\text{Ca}^{2+}]^m} + k_4$$

(f) Reversible isomerisation:

$$k_{\text{obs}} = \frac{k_2 K_1 [\text{Ca}^{2+}]^n}{1 + K_1 [\text{Ca}^{2+}]^n} + k_2$$



Experimental Details

Ca^{2+} equilibrium titrations:

- RGECIs in assay buffer (50 mM K⁺-HEPES, 100 mM KCl, 2 mM MgCl₂, pH 7.5) + 10 mM EGTA
- continuous titration with 325 mM CaCl₂

Ca^{2+} transients in HEK293T cells:

- RGECIs expressed in HEK293T cells
- after 16-48 h imaged on confocal Zeiss AxioObserver Z1 at 37 °C (CMOS 2 s interval, λ_{ex} = 568 nm)
- Ca^{2+} transients triggered with 50 μM ATP

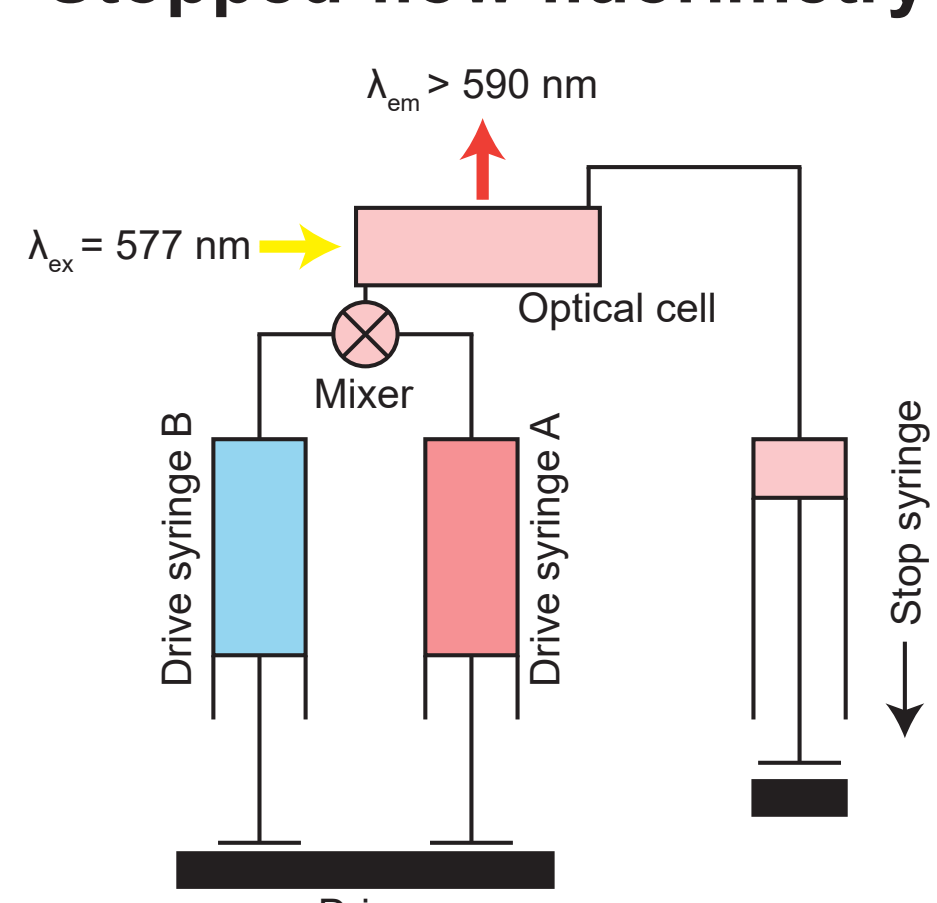
Association kinetics:

RGECIs in assay buffer + 10 mM EGTA were rapidly mixed with increasing CaCl₂ in assay buffer

Dissociation kinetics:

RGECIs in assay buffer + 1 mM CaCl₂ were rapidly mixed with 12.5 mM EGTA (conc. in mixing chamber)
Free $[\text{Ca}^{2+}]$ calculated by Maxchelator.

Stopped-flow fluorimetry



References

- [1] Dana et al., eLife, 2016.
- [2] Helassa et al., Sci. Rep., 2015
- [3] Helassa et al., Sci. Rep., 2016

Funding
BB/M02556X/1