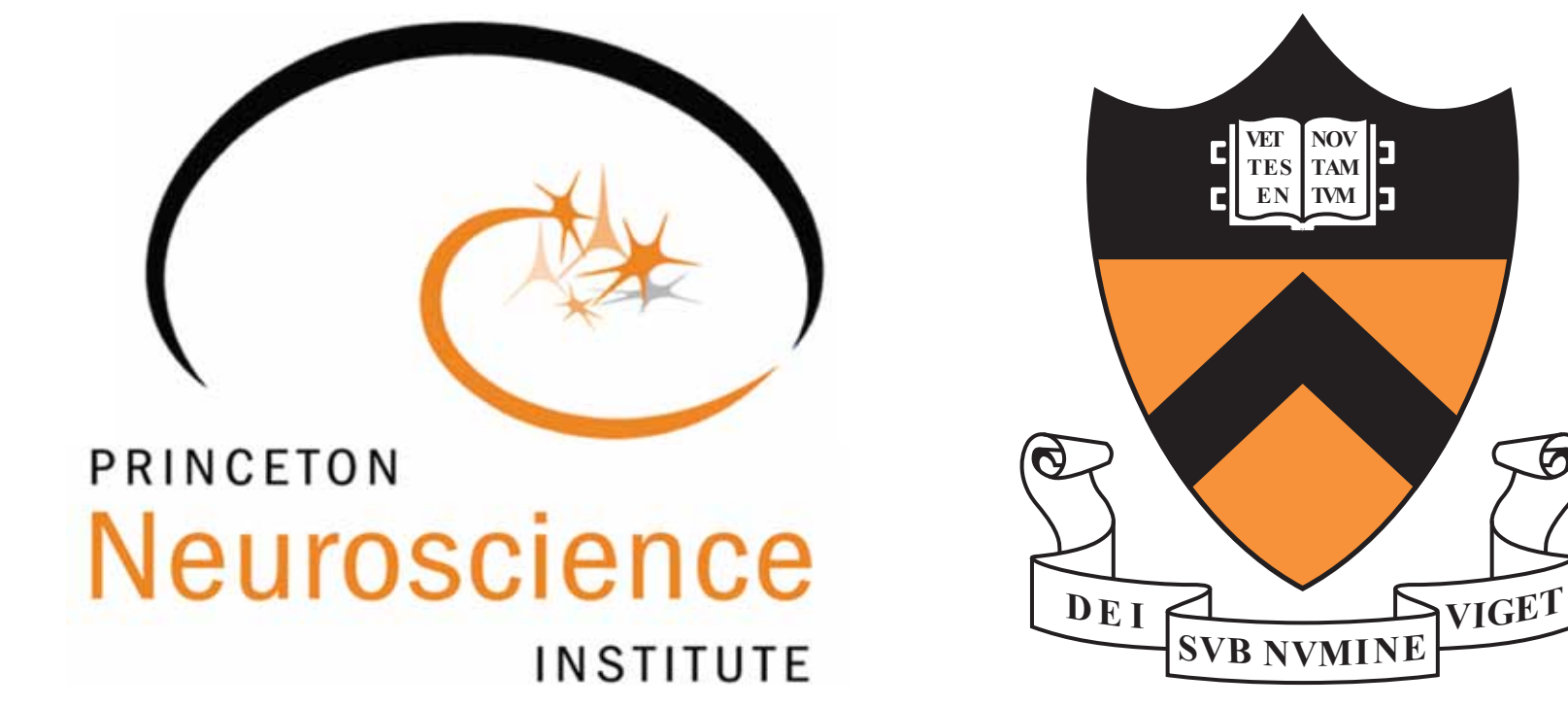


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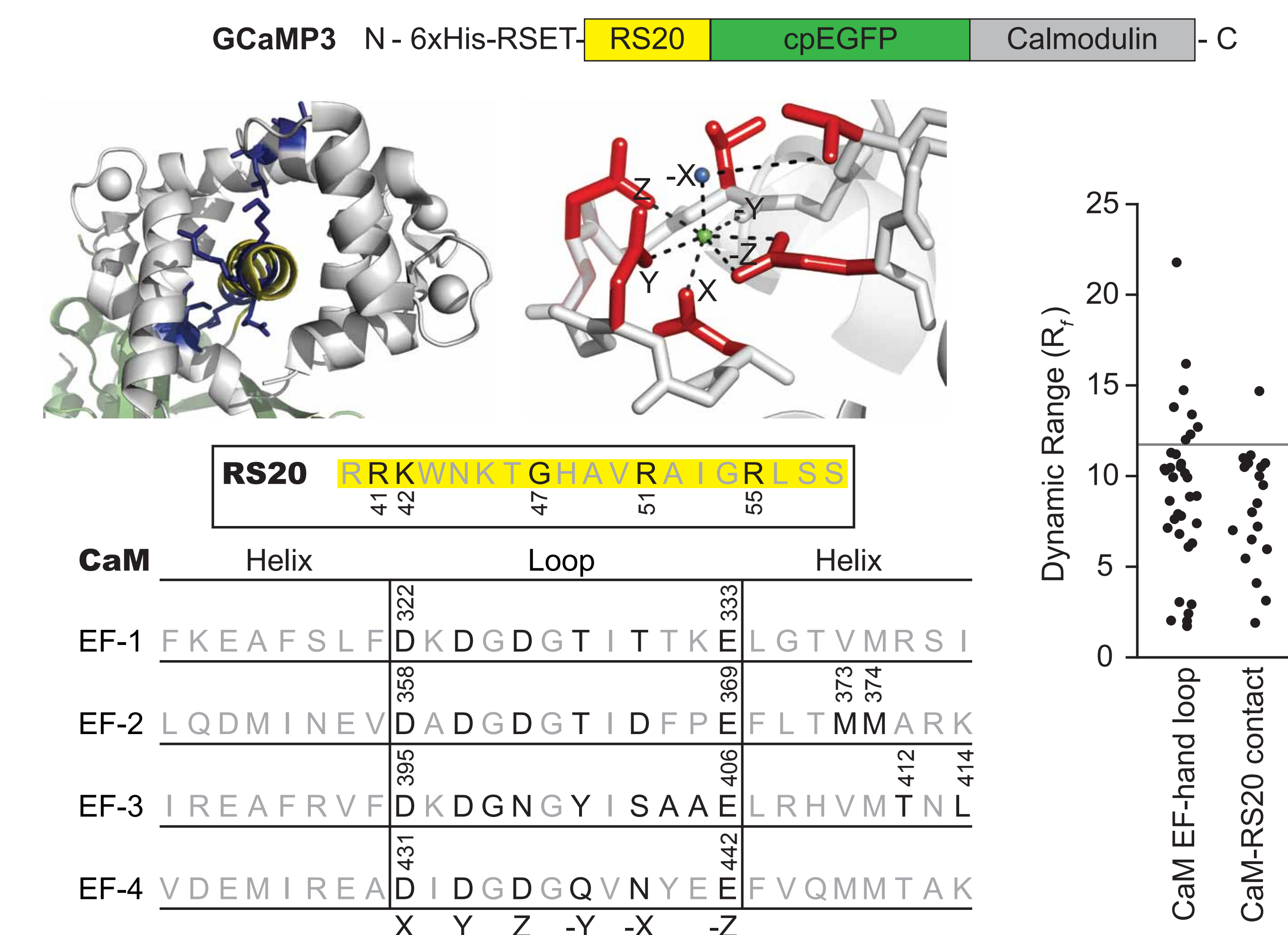
ABSTRACT

Although fluorescent genetically encoded calcium indicator (GECI) proteins provide a powerful tool for monitoring neuronal activity, they still have significant performance limitations compared with synthetic indicators such as Oregon Green BAPTA-1 (OGB-1). Because of high cooperativity originating from an EF hand protein-based detection mechanism, a given GECI is only sensitive to a small part of a neuron's likely calcium concentration range, which can exceed 1 μM during intense activity. GECIs also have up to 100-fold slower response kinetics than OGB-1, whose τ_{on} is ~ 1 ms to a step increase in calcium and $\tau_{\text{off}} \sim 7$ ms to a step decrease. Overcoming limitations in range and kinetics is a key step toward monitoring spike times and firing rates in cell-type-specific brain circuits.

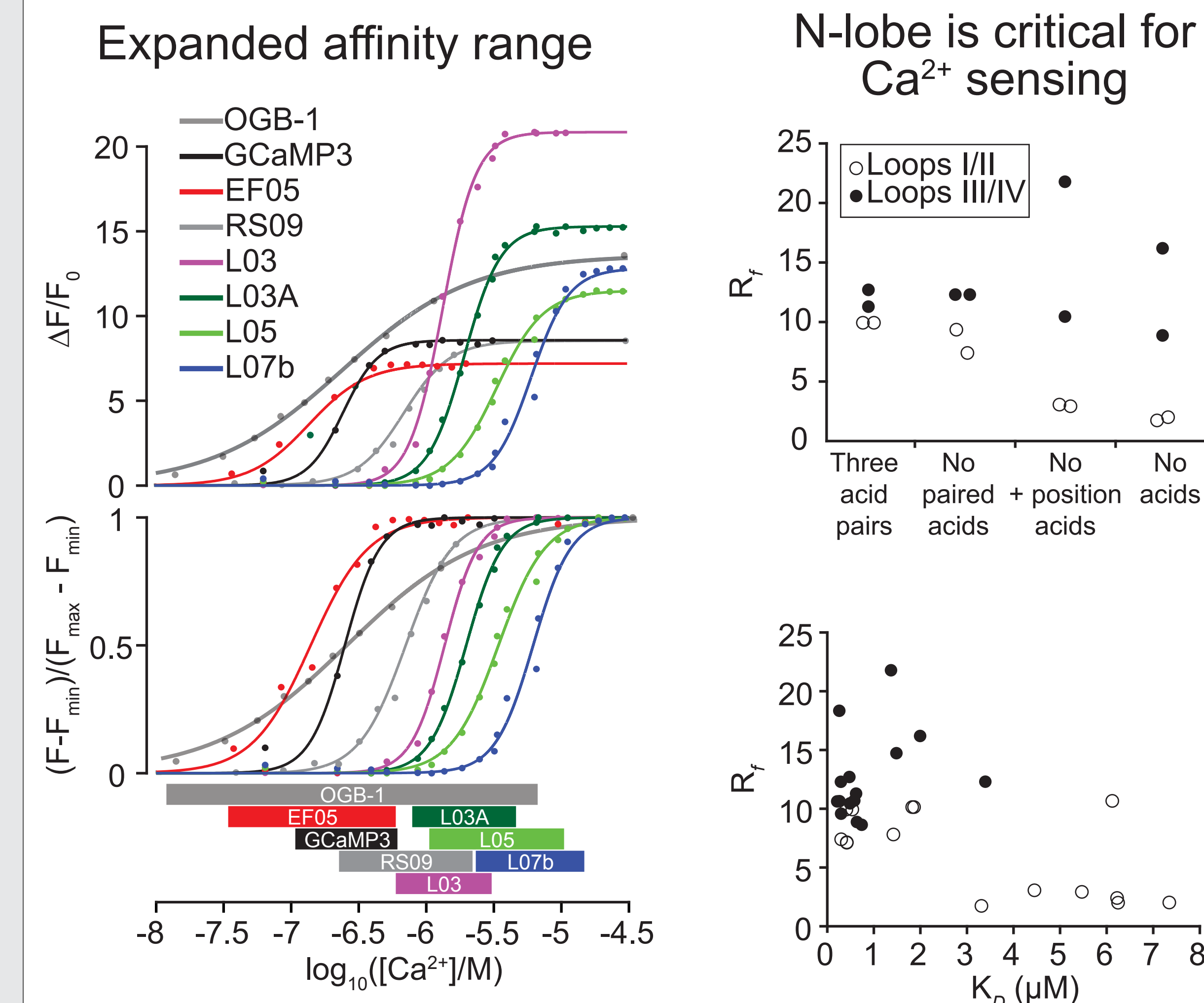
We are engaged in structure-based design to vary the affinity and accelerate the response kinetics of a popular GECI, GCaMP3 (Nakai et al. 2001 Nat. Biotechnol. 19:137; Tian et al. 2009 Nat. Methods 6:875). We have designed more than 50 variants by targeted mutation of GCaMP3's calmodulin domain and its intraprobe binding partner, RS20. In our cuvet characterizations of purified protein, we have (1) attained a nearly 40-fold (0.16-6 μM) range of K_D without impairing per-molecule brightness, and (2) made stopped-flow biochemical measurements in which off-responses (i.e. to decreasing steps in calcium) are more than 10 times faster than any other published GECI. Most of the gap in off-response speed between G-CaMP3 and OGB-1 could be closed without perturbing K_D (see Figure panel A).

We are using two-photon laser scanning microscopy to characterize our most promising variants in cultured sympathetic neurons by pseudorabies virus (PRV) infection, *Drosophila* neuromuscular junctions by transgenesis, and cerebellar neurons in brain slices by adeno-associated virus (AAV) infection (Figure panels B/C/D). Our long-term goal is to generate and identify variants and combinations of variants that detect single action potentials and/or firing rate modulations in the 1-100 Hz range.

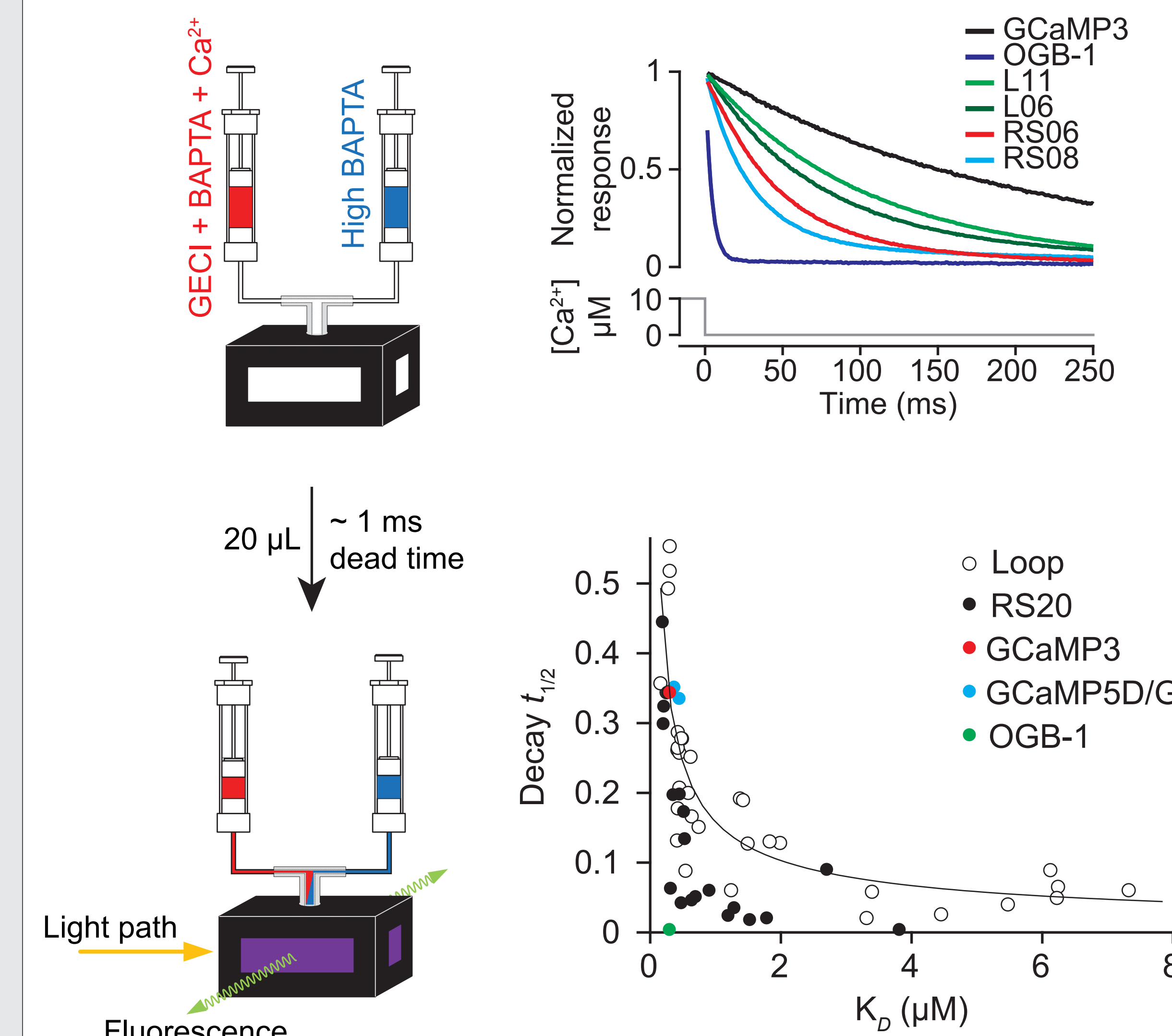
1 STRUCTURE-ORIENTED OPTIMIZATION



2 CONTROLLING AFFINITY



3 FASTER OFF-RESPONSES TO DECREASING CALCIUM

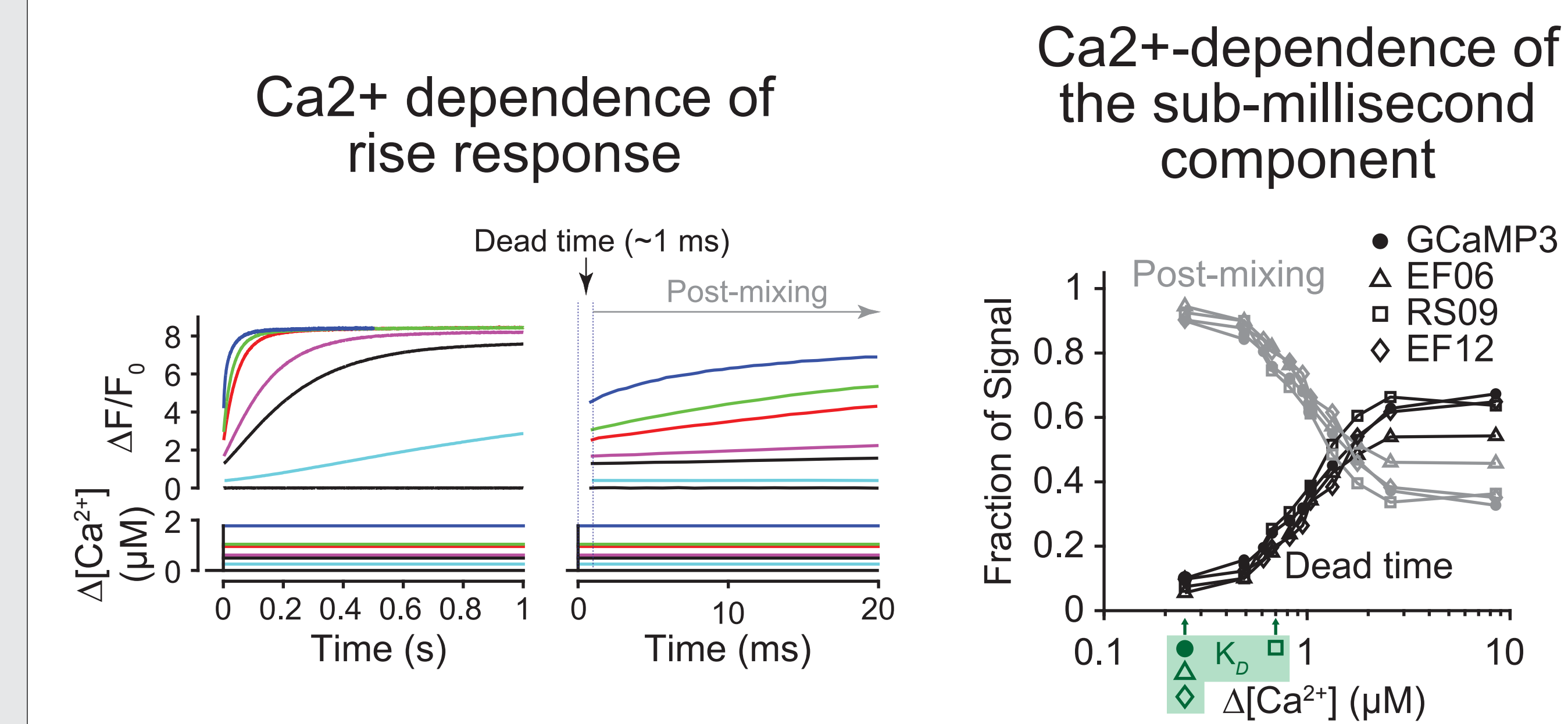


4 A RANGE OF AFFINITY AND OFF-RESPONSE TIMES

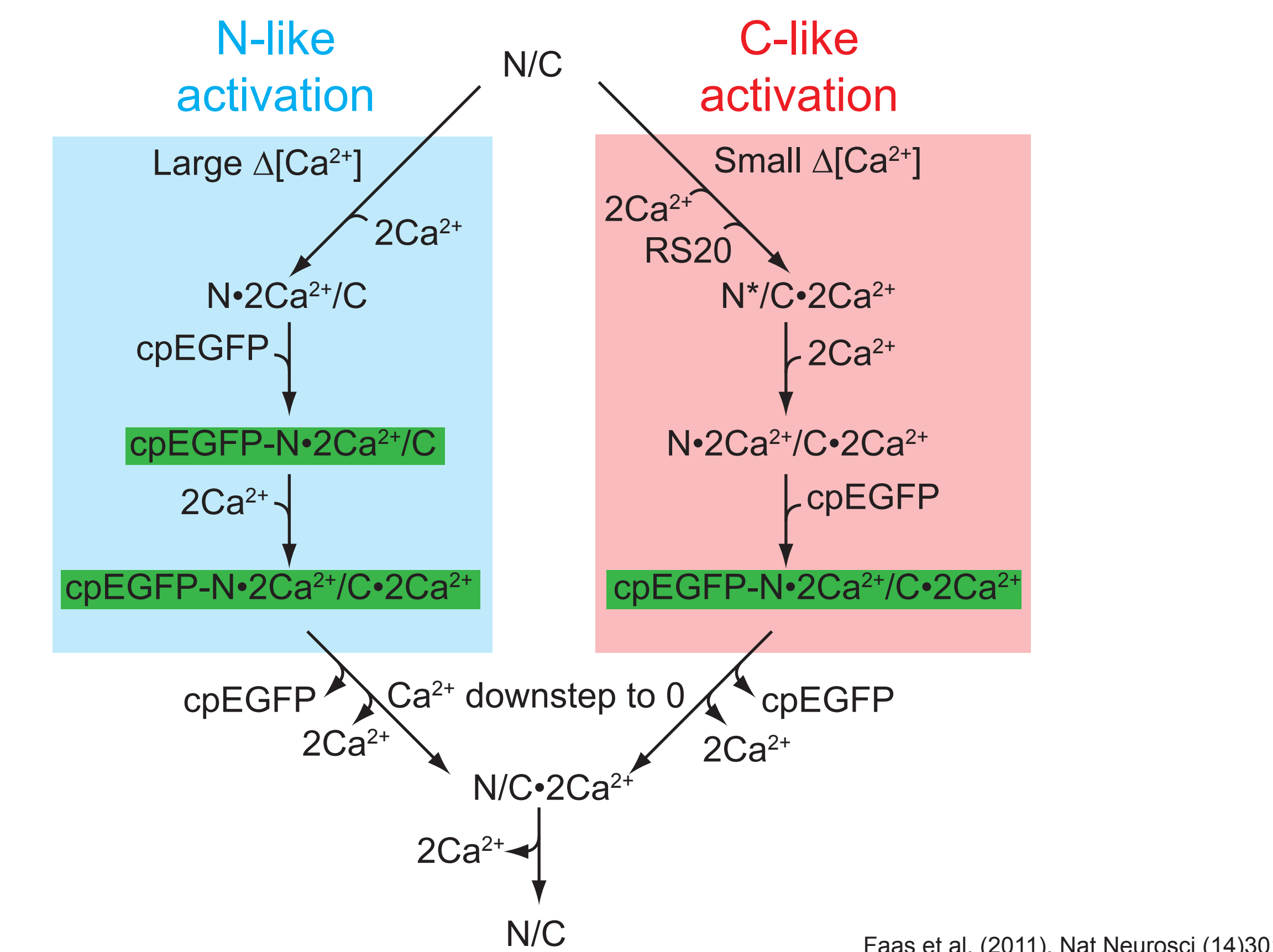
Variant	Altered domain	K_D (μM) ¹	R_f	Decay $t_{1/2}$ (ms) ²
OGB-1	-	0.24	14.0	5
GCaMP3	-	0.25	12.0	150
EF05	Loop	0.16	6.8	183
L03	Loop	1.37	21.8	80
L03A	Loop	1.99	16.2	52
L05	Loop	3.39	12.3	22
L07b	Loop	6.12	13.8	35
RS06	CaM-Helix	0.31	6.5	34
RS05	CaM-Helix	0.47	8.5	27
RS08	CaM-Helix	0.63	10.0	23
RS09	CaM-Helix	0.69	9.5	25
RSd03	RS20	0.90	11.1	22
RSd05	RS20	1.19	10.7	10
RSd06	RS20	1.78	10.6	7

¹23°C; ²37°C

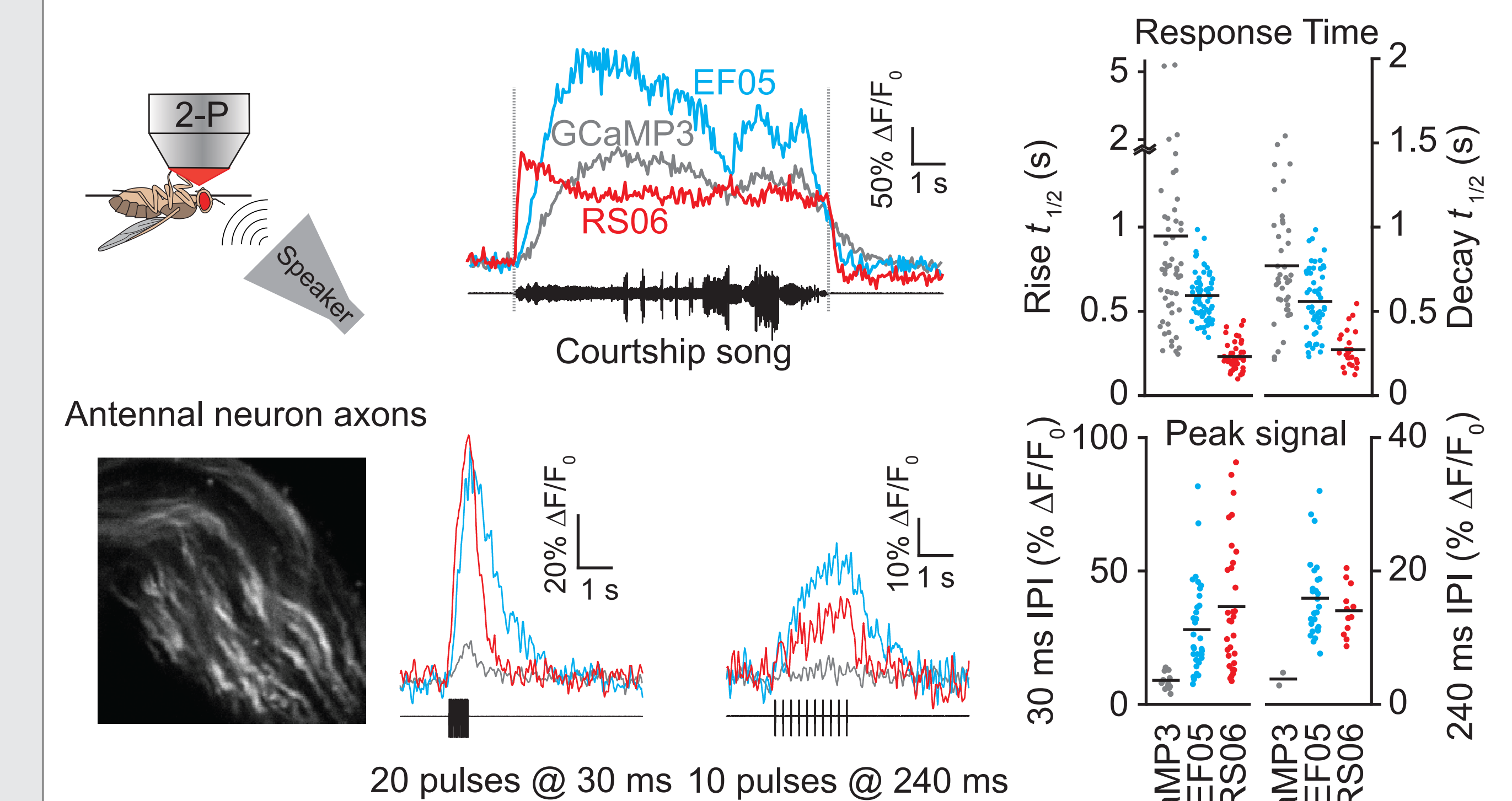
5 A LOW-AFFINITY MODE FOR RAPID ON-RESPONSES



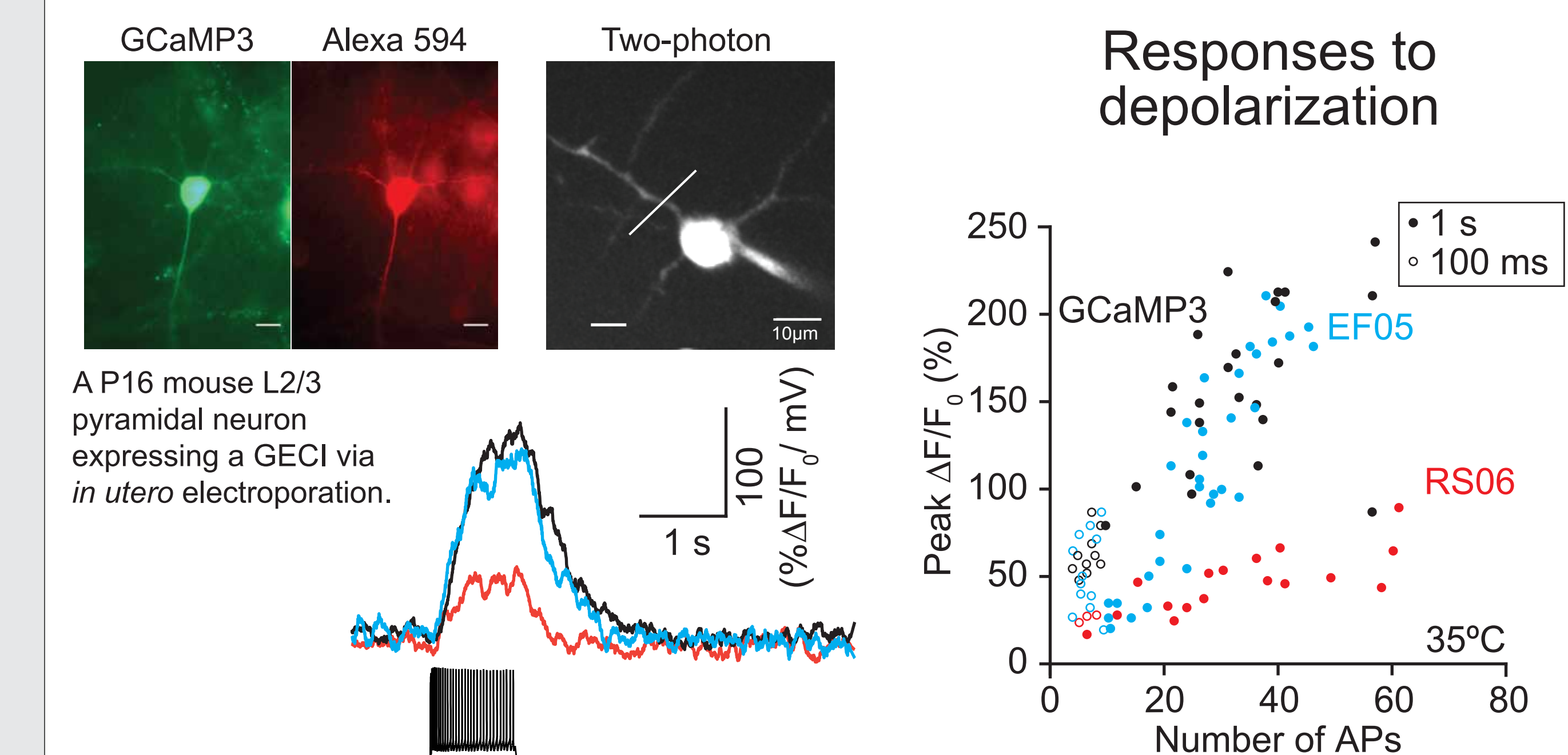
A TWO-PATH MODEL FOR GCaMP ACTIVATION



6 AXON RESPONSES IN DROSOPHILA



7 RESPONSES IN L2/3 NEURONS



SUMMARY

- Of over 50 variants made, 32 have dynamic ranges within 0.7-1.8 times that of GCaMP3.
- Variants achieved nearly 40-fold (0.16-6 μM) ranges of K_D .
- We achieved a 4- to 50-fold improvement off-response times.
- Fast, low affinity component to the rise response in all variants tested.
- In *Drosophila*, variants performed as expected from K_D , with fast variants showing a ~ 3 -fold faster rise and decay of signals.

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