

CHAPTER 2

CNIFERS: CELL-BASED BIOSENSORS WITH NANOMOLAR SENSITIVITY TO *IN VIVO* CHANGES IN NEUROMODULATION

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2.1 Introduction

Neuromodulators, such as dopamine, norepinephrine, serotonin, endocannabinoids and neuropeptides, all produce profound effects on circuits that control animal behavior. Perturbation of neuromodulation is implicated in several different brain disorders that include substance abuse and dependence, depression, and severe anxiety. A major obstacle to studying the effect of neuromodulators in the brain has been the inability to detect their release *in vivo* with sufficient sensitivity, chemical specificity, and spatial and temporal resolution. Here, we describe the development of new tools for detecting neuropeptides and other neuromodulators.

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2.2 Neuropeptides

Neuropeptides are small proteins that signal like neurotransmitters in the nervous system. Except for a few exceptions, such as FMRFamide-dependent opening of Na⁺-channels, neuropeptides typically signal through G protein-coupled receptors (GPCRs) (Lingueglia *et al.*, 1995), have a slower time course than ionotropic receptors, and produce persistent signaling. Peptide signaling through the GPCRs is amplified by G proteins, which stimulate/inhibit enzymes that generate second messengers like cAMP, IP₃, and Ca²⁺ that in turn modulate downstream kinases, e.g., protein kinase A (PKA), protein kinase C (PKC) (Sakurai *et al.*, 1998; Thibonnier *et al.*, 1998). These signaling pathways can also modulate the response of neurons through activation or inhibition of multiple ion channels, including voltage-gated calcium channels and potassium channels (Borgland *et al.*, 2008; Li and van den Pol, 2008).

Neuropeptides are synthesized as an inactive precursor on the rough endoplasmic reticulum in the soma, transferred to the Golgi apparatus for packaging, and then transported down the axon (Sossin *et al.*, 1989) (see Figure 2.1). The active peptide is generated by proteolytic cleavage of the precursor in large dense core

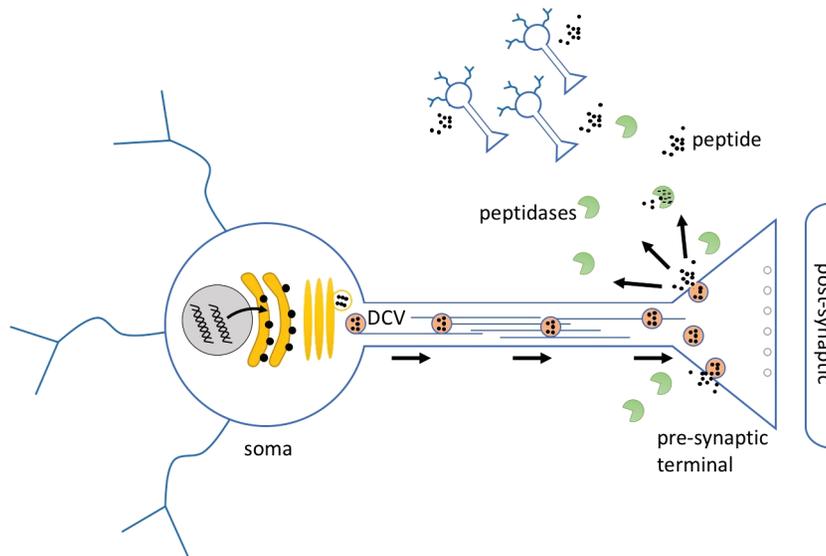


Figure 2.1 Cartoon of peptide synthesis and release in neurons. Neuropeptides are synthesized from mRNA in the soma on the rough ER as an inactive precursor, transferred to the Golgi apparatus for sorting, and transported down the axon in large dense core vesicles (DCVs). Active neuropeptide is generated by the proteolytic cleavage of the precursor peptide in DCVs. Neuropeptides act on receptors located on neurons several hundred micrometers away from release sites. Peptide half-life and diffusion are controlled, in part, by extrasynaptic peptidases, which catabolize the released peptides. In contrast to primary neurotransmitters, peptides are not recycled for re-use.

vesicles (DCVs). In general, DCVs contain multiple peptides and coexist with small vesicles housing small molecule transmitters (Chou *et al.*, 2001; Goodman *et al.*, 2007). In contrast to primary neurotransmitters, peptide-containing DCVs are mostly excluded from the active zone and found at greater distances from the synapse (Kreiner *et al.*, 1986; Fisher *et al.*, 1988). As most voltage-gated Ca^{2+} channels are localized to the active zone, peptide release is thought to require longer and higher frequency firing compared to small vesicle release (Sudhof, 2012).

The extrasynaptic localization of DCVs raises questions on the nature of neuropeptide signaling, particularly how far neuropeptides travel through the extracellular space. Minute-long half-lives of the neuropeptides and high-affinity of the neuropeptide receptors to the ligands ($K_d \sim \text{nM}$) have led researchers to postulate that peptides signal through volume transmission and act on extra-synaptic targets located several hundred micrometers away (Mens *et al.*, 1983; Solano *et al.*, 1996; Schloos *et al.*, 1997; Ludwig and Leng, 2006).

Perturbations in neuropeptide signaling have been implicated in several brain disorders. For example, alterations in neuropeptides appear to play a role in schizophrenia (Caceda *et al.*, 2007; LaCrosse and Olive, 2013). The hypothalamic neuropeptides oxytocin and vasopressin appear to be critically important for the regulation of social and cognitive processes that are abnormal in schizophrenia. The genes that encode oxytocin and vasopressin are associated with schizophrenia, with reduced pituitary and brain levels for both peptides found in individuals with the disease (Frank *et al.*, 2005). In contrast, oxytocin and vasopressin in the hypothalamus appear to be upregulated in individuals who suffer from major depressive disorder (Purba *et al.*, 1996).

Both of these neuropeptides can potentiate the effects of the corticotropin releasing factor (CRF) peptide and thus the activation of the hypothalamic pituitary adrenal (HPA) axis (Raadsheer *et al.*, 1994). Disruption in the stress related peptide CRF has also been implicated in drug addiction (Koob, 1999; Logrip *et al.*, 2011). Many drugs with abuse potential produce potentiation of the CRF system and likely drive atypical activity in response to stressors and contribute to relapse of drug abuse (Kreek and Koob, 1998; Roberto *et al.*, 2017).

2.3 Challenges of studying neuromodulators

Contemporary techniques for studying neuromodulators, such as microdialysis, have provided information on the extracellular concentration of neuromodulators *in vivo* (Day *et al.*, 2001). Microdialysis has been used to measure dopamine release in the medial prefrontal cortex (mPFC) in response to alcohol administration (Schier *et al.*, 2013), where alcohol administration produced significant peak increases in dopamine in the dialysate from mPFC. Microdialysis has also been

used to study other neuromodulators such as the neuropeptide oxytocin. Oxytocin is released in response to social stimuli in the dorso-lateral septum, and is blocked by social fear conditioning (Zoicas *et al.*, 2014). Although microdialysis is suitable for the accurate identification of different neuromodulators (Ji *et al.*, 2008; Greco *et al.*, 2013), there are some disadvantages. First, the collection of large CSF samples and the insertion of the microdialysis probe can disrupt neuromodulator activity near the probe track (Wang and Michael, 2012). Second, the temporal resolution is poor, with the sample collection typically occurring at 10 min intervals or longer, and not in real-time (Mingote *et al.*, 2004).

An alternative method for measuring the extracellular concentration of neuromodulators is fast-scan cyclic voltammetry (FSCV) (Robinson and Wightman, 2007). Compared to microdialysis, FSCV improves the spatial and temporal resolution as well as having nanomolar sensitivity. However, the number of neuromodulators that have been successfully examined with FSCV is limited (e.g., DA, serotonin, and NE). This technique is further limited by the inability to distinguish chemically similar compounds. For example, DA and NE differ by only one hydroxyl, and produce similar cyclic voltammetry signatures (Robinson *et al.*, 2003), making examinations of neuromodulator concentrations in brain regions where both are expressed a challenge.

Measuring changes in neuronal activity with single-cell electrophysiology is another technique for studying the effect of neuropeptides in the brain. Exposure to neuropeptides induces changes in excitatory or inhibitory post synaptic currents (amplitude or frequency), action potential firing, cell excitability, or current/voltage relationships (Radhakrishnan and Henry, 1995; Raggenbass, 2001; Jobst *et al.*, 2004; Gilpin and Roberto, 2012). However, electrophysiological measurements of neuropeptide release are hampered by the small amount of neuropeptide released by a single axonal bouton in the CNS (Dreifuss, 1975; Nordmann and Morris, 1984), which is likely due to a requirement for elevations in cytoplasmic Ca^{2+} that are higher than needed for fast, synaptic release (Tallent, 2007). In addition, the response of receptive cells is relatively slow (Rosenbaum *et al.*, 2009; Hazell *et al.*, 2012), taking many seconds to minutes, making it a challenge to correlate peptide release with a post synaptic response, or to use quantal analysis of single or multiple exocytotic events (Hazell *et al.*, 2012). Unlike fast neurotransmitters, such as glutamate or GABA, the rise time for neuromodulators is much slower making it harder to segregate individual events (van den Pol, 2012).

2.4 Cell-based neurotransmitter fluorescent engineered reporters

To overcome limitations with detecting release of neuromodulators *in vivo*, we have developed a new optical approach for detecting the release of neuromodulators. This

new technology is termed **cell-based neurotransmitter fluorescent engineered reporters** (CNiFERS).

CNiFERS leverage fluorescence resonance energy transfer (FRET) to translate transmitter binding into an optical signal. CNiFER cells are engineered from HEK293 cells by stably expressing a specific GPCR for the neuromodulator of interest and a FRET-based calcium sensor, e.g., TN-XXL (Nguyen *et al.*, 2010; Muller *et al.*, 2014; Lacin *et al.*, 2016). For Gq-coupled GPCRs, binding of the neuromodulator of interest to the GPCR expressed on the surface of HEK293 cells leads to an increase in intracellular Ca^{2+} which is detected by the FRET-based Ca^{2+} sensor, i.e. decrease in CFP (FRET donor) emission and an increase of YFP (FRET acceptor) emission (Figure 2.2). CNiFERS can be implanted directly into target brain regions where they can detect release of physiological concentrations of modulators with a temporal resolution of seconds and spatial resolution of less than 100 μm (Nguyen *et al.*, 2010; Muller *et al.*, 2014) (Figures 2.3 B–D). The dual wavelength measurement of the

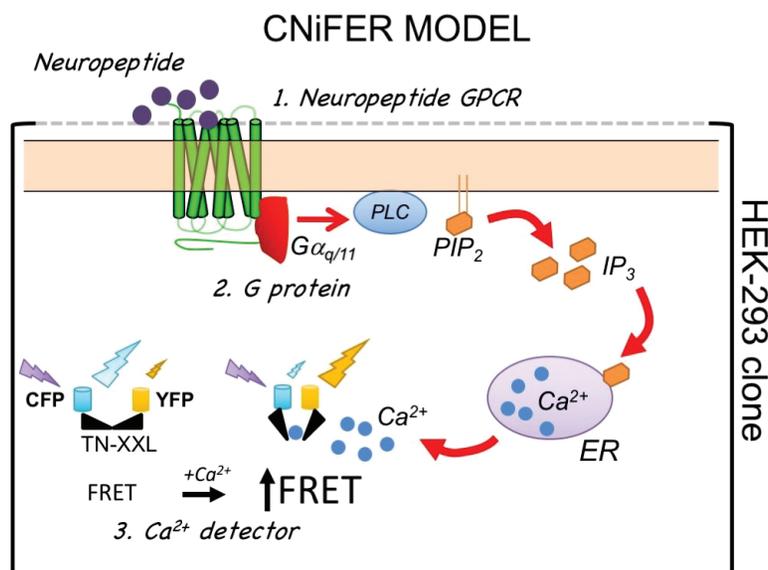


Figure 2.2 Design of peptide CNiFERS. (1) The GPCR for the neuropeptide of interest is expressed on the plasma membrane of HEK-293 cells. (2) Neuropeptide binding produces a conformational change and activation of the G_q G protein, which in turn activates phospholipase C (PLC). A chimeric protein (G_{q15} or G_{qs5}) can also be expressed to redirect G_{i/o} or G_s-coupled receptors, respectively, to the PLC pathway. Cleavage of PIP₂ into diacyl glycerol and IP₃ by PLC leads to the release of intracellular Ca²⁺ from the endoplasmic reticulum. (3) Ca²⁺ binds to the genetically encoded Ca²⁺ sensor which undergoes FRET. Two-photon excitation of the CNiFER in the absence of the neuropeptide agonist exhibits high CFP and low YFP fluorescence. Neuropeptide binding increases intracellular Ca²⁺ and produces an increase in FRET, detected as a decrease in CFP emission and increase in the YFP emission with CFP excitation.

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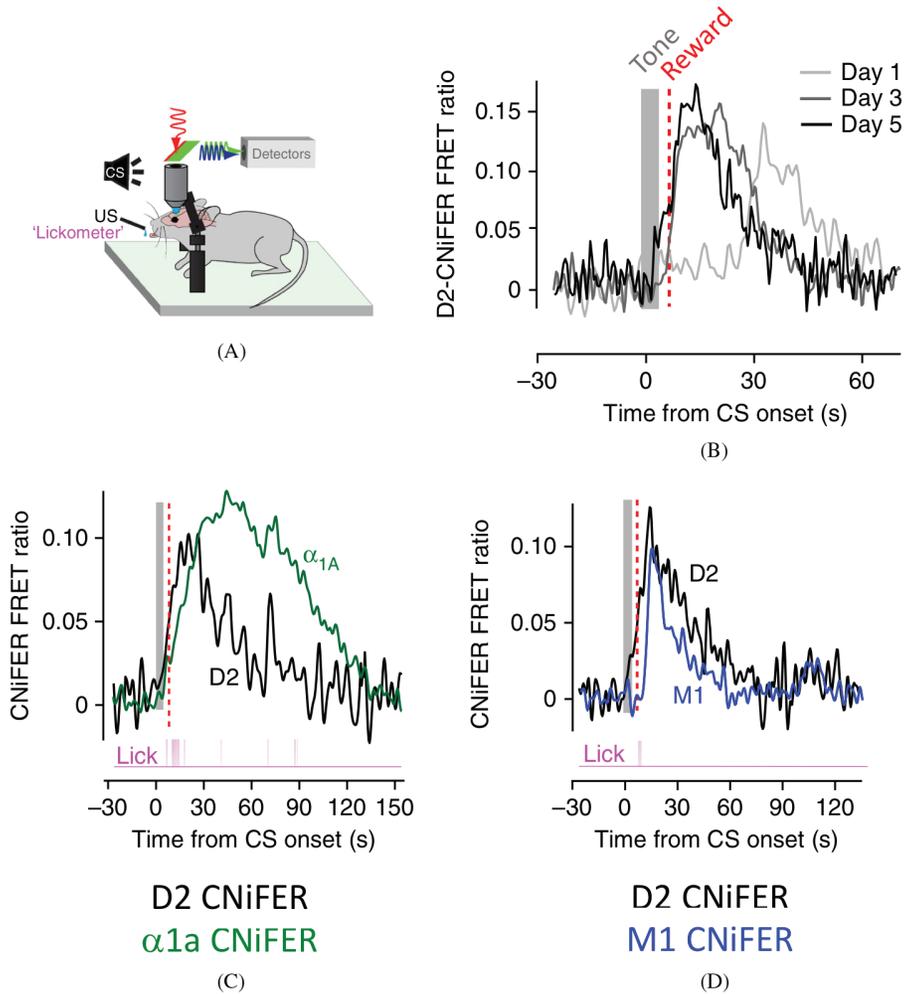


Figure 2.3 Example of CNiFER detection of DA, NE and ACh *in vivo*. (A) Procedure to measure licking behavior and CNiFER fluorescence in head-restrained mice during classical conditioning. (B) Single trial examples of the D2-CNiFER response in the same animal at days 1, 3 and 5 of training. (C) Simultaneous measurement of D2- and α_{1A} -CNiFER FRET, or D2- and M1-CNiFER FRET, in conjunction with bouts of licking, during a single conditioning trial (modified from Muller *et al.*, 2014).

The FRET-based Ca^{2+} sensor provides an intrinsic signal for correcting possible motion-induced artifacts.

2.4.1 Creation of cell-based neurotransmitter fluorescent engineered reporters (CNiFERs)

For each neuromodulator, an appropriate GPCR is stably expressed in a clonal HEK293 cell with a Ca^{2+} sensor, i.e., TN-XXL or equivalent Ca^{2+} sensor

(Figure 2.2). For $G_{i/o}$ - or G_s -coupled receptors, a chimeric G protein G_{qi5} or G_{qs5} , respectively, is also expressed in the HEK293 cell to redirect GPCR signaling to the G_q - Ca^{2+} pathway. The calculated fractional change in fluorescence, $\Delta F/F$, for each signal is used to obtain the FRET ratio, denoted $\Delta R/R$. Each clonal CNiFER is assessed for potency (e.g. EC_{50}), efficacy (i.e., largest increase in FRET ratio), and lowest background responses to other neurotransmitter/neuromodulators (Nguyen *et al.*, 2010; Muller *et al.*, 2014; Lacin *et al.*, 2016). The control CNiFER contains the same Ca^{2+} detection components but lacks the specific GPCR. The CNiFER approach is easily adapted for any neurotransmitter, neuromodulator or neuropeptide that signals through a specific GPCR. Thus, a library of CNiFERS can be created for assessing the release of a large number of different neuromodulators. In addition, implanting two different CNiFER types in the brain affords the opportunity for multiplexing, and simultaneous detection of the release of two different neuromodulators (Figures 2.3C and 2.3D) (Nguyen *et al.*, 2010; Muller *et al.*, 2014; Lacin *et al.*, 2016).

The creation of dopamine and norepinephrine CNiFERS provides a good example of simultaneous detection of chemically similar neurotransmitters (Muller *et al.*, 2014). The dopamine CNiFER was created with the D2 dopamine receptor (D2) and norepinephrine CNiFER with the α_{1A} receptor. The D2 and α_{1A} CNiFERS have an EC_{50} of 2.5 nM and 19 nM, respectively, suitable for detection of endogenously released transmitter (Muller *et al.*, 2014). We injected the D2 CNiFER and α_{1A} CNiFER into the cortex, approximately 200 μ m apart, and measured the release of DA and NE in awake mice undergoing Pavlovian conditioning, i.e. repeated pairing of a tone (CS) with a sugar reward (Figures 2.3A and 2.3B). We also compared the release of DA with ACh, using a muscarinic M1 CNiFER (Nguyen *et al.*, 2010) and D2 CNiFER for detecting DA (Figure 2.3C). This multiplexing of two different CNiFERS in the brain allows for monitoring the release of two different neuromodulators simultaneously. As predicted, the timing of DA release shifts from the reward to the CS with training. Interestingly, the timing of NE release is highly variable whereas ACh release is tightly correlated with licking (Muller *et al.*, 2014).

Multiplexing of CNiFERS could also be used to compare a single transmitter interacting with two different receptors. For example, dopamine signals through both D1 and D2 receptors, which have different affinities for DA (Lobo and Nestler, 2011). Endogenous dopamine release in a particular region may preferentially activate one CNiFER (D2 *versus* D1) over the other. Another example of multiplexing CNiFERS is to co-implant one CNiFER for a traditional neuromodulator, such as DA, with another CNiFER that expresses a GPCR for a neuropeptide, such as for the kappa opioid receptor to measure the neuropeptide dynorphin (Dyn). DA and Dyn CNiFERS could be implanted into the striatum

of a mouse to measure release dynamics in response to exposure to a drug of abuse such as alcohol (Resendez *et al.*, 2016).

The CNiFER design allows for measuring neuromodulator release on the time scale of seconds, mimicking the activation of natively expressed GPCRs. This temporal response of CNiFERs is a significant improvement over the response time of minutes to hours for microdialysis. The fast response time, high affinity, and the ability to generate a library of sensors capable of measuring any modulator that signals through a GPCR, makes CNiFERs well suited for examining the temporal dynamics of local neuropeptide release. A limitation with CNiFERs, on the other hand, is the inability to directly capture the activity of neuromodulators released within the synapse.

2.5 Future for CNiFER-based research

CNiFERs were first created with HEK293 cells because HEK293 cells typically have low expression of endogenous GPCRs and are easily transduced to create clonal lines. Although HEK293 cells are human cells, implantation into mice does not appear to produce a significant inflammatory response (Muller *et al.*, 2014) when mice are injected with an immunosuppressant (e.g., cyclosporine). Under these conditions, CNiFERs can remain functional for up to a week. However, monitoring neuromodulator release over longer periods might require repeated CNiFER injections. An alternative approach would be to create CNiFERs using rodent cells lines, such as mouse embryonic fibroblasts or chinese hamster ovary (CHO) cells. We have tested CHO cells and measured FRET responses comparable to HEK293 cells (unpublished data).

Currently, CNiFERs are implanted into cortex and monitored *in vivo* using two-photon laser scanning microscopy (TPLSM). However, neuropeptide-producing cells and terminals are widely distributed in the brain, including sub-cortical regions (Lein *et al.*, 2007; Sakurai, 2014; Acevedo-Rodriguez *et al.*, 2015). Standard TPLSM allows for imaging up to 1 mm in depth and cannot easily image areas below the cortex. In addition, animals are head-fixed for most TPLSM setups, limiting the range of behaviors that can be studied during imaging. Recent advances in imaging technologies have made it possible to monitor subcortical activity in freely moving animals (Gosh *et al.*, 2011; Cai *et al.*, 2016). For example, miniature microscopes (e.g. endoscopes or Miniscopes) contain all the electronic and optical components necessary for imaging and are directly mounted to a mouse skull and tethered to a cable. The head-mounted scopes are usually coupled with a gradient-refractive-index (GRIN) lens that is implanted near the target and relays the fluorescent signal to the scope (Girven and Sparta,

2017). Although currently available Miniscopes are designed for single-wavelength measurements, it should be possible to create a two-channel Miniscope without adding too much weight. Another approach for subcortical measurements of CNiFERS is to use fiber photometry. Fiber photometry is commonly used to measure Ca^{2+} transients using genetically-encoded Ca^{2+} sensors (e.g., GCaMPs) and involves insertion of an optical fiber and coupling to photomultiplier tubes (PMTs) (Girven and Sparta, 2017). Fibers have the advantage of producing less tissue damage because of the small size, ~ 0.2 mm, compared to GRIN lenses which typically are 0.5–1 mm in diameter (Girven and Sparta, 2017). Fiber photometry measurements lack spatial information, due to summation of the light signal from all cells. However, the average FRET activity is measured in multiple CNiFER cells within an implant, making fiber photometry well suited for CNiFER imaging.

An alternate approach to cell-based CNiFERS is the development of genetically encoded neuromodulator sensors that are capable of capturing the activity of neuromodulators within synapses. The first generation of genetically encoded neurotransmitter detectors were based on mutations of periplasmic binding proteins and detected glutamate release (iGluSnFR (Marvin *et al.*, 2013), EOS (Namiki *et al.*, 2007), Snifit (Brun *et al.*, 2011), FLIPE (Okumoto *et al.*, 2005), albeit with microMolar sensitivity. Recently, two groups have developed sensors for detecting DA based on coupling steric changes in a GPCR for DA, which occur upon binding of DA, into producing changes in the optical properties of a closely coupled fluorescent protein (FP). Patriarchi *et al.* (2018) created a fusion protein that directly couples dopamine binding induced conformational changes in an inert human dopamine receptor to changes in the fluorescence intensity of a circularly permuted green fluorescent protein (cpGFP) (termed dLight1). Sun *et al.* (2018) used a similar strategy but inserted cpEGFP in the intracellular loop of the human dopamine receptor (termed GRAB_{DA}). One advantage of these sensors is the ability to express the sensor in a particular neuronal population of interest and capturing the temporal dynamics of neuromodulator activity within the synapse (Patriarchi *et al.*, 2018; Sun *et al.*, 2018). Additionally, the wide expression enables detection of neuromodulator concentrations across the cortical mantle.

An advantage of using CNiFERS is the ability to multiplex and study the simultaneous release of multiple neuromodulators. Currently, the simultaneous imaging of two different transmitters is accomplished by injecting two types of CNiFERS within a 100 μm of each other. Because the two sensors are spatially distinct, it is possible to measure FRET in two implants simultaneously. Another approach is to mix CNiFERS of different colors in the same implant. For example, choosing a calcium sensor that emits at a longer wavelength than the CFP and

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YFP of TN-XXL, such as with a red Ca^{2+} sensor (Akerboom *et al.*, 2013). A CNiFER with a longer wavelength calcium sensor also has the advantage of imaging deeper in the brain (Akerboom *et al.*, 2013; Tischbirek *et al.*, 2015) and the longer wavelength excitation two-photon light necessary to excite a red or near-infrared fluorophore scatters less, produces less phototoxic damage, and penetrates deeper (Yaroslavsky *et al.*, 2002; Tischbirek *et al.*, 2015). Moreover, when CNiFER imaging is coupled to optogenetics, it enables continuous imaging of CNiFERs during blue light stimulation of channel-rhodopsin.

In summary, the methods for optically detecting the release of neurotransmitters or neuromodulators *in vivo* in awake animals are rapidly advancing, providing neuroscientists with state-of-the-art tools for probing the function of neuromodulators on brain circuits.

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