

Title: Genetically encoded tools for in vivo GPCR agonist detection at cellular resolution

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Abstract

G-protein-coupled receptors (GPCRs) are the most abundant receptor type in the human body and are responsible for regulating many physiological processes, such as sensation, cognition, muscle contraction, and metabolism. Further, GPCRs are widely expressed in the brain where their agonists make up a large number of neurotransmitters and neuromodulators. Due to the importance of GPCRs in human physiology, genetically encoded sensors have been engineered to detect GPCR agonists at cellular resolution in vivo. These sensors can be placed into two main categories: those that offer real-time information on the signaling dynamics of GPCR agonists and those that integrate the GPCR agonist signal into a permanent, quantifiable mark that can be used to detect GPCR agonist localization in a large brain area. In this review, we discuss the various designs of real-time and integration sensors, their advantages and limitations, and some in vivo applications. We also discuss the potential of using real-time and integrator sensors together to identify neuronal circuits affected by endogenous GPCR agonists and perform detailed characterizations of the spatiotemporal dynamics of GPCR agonist release in those circuits. By using these sensors together, the overall knowledge of GPCR-mediated signaling can be expanded.

Introduction

This is the author manuscript accepted for publication and has undergone full peer review but has not been through the copyediting, typesetting, pagination and proofreading process, which may lead to differences between this version and the [Version of Record](#). Please cite this article as [doi: 10.1002/ctm2.1124](https://doi.org/10.1002/ctm2.1124).

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G-protein-coupled receptors (GPCRs) are seven-transmembrane domain receptors responsible for modulating an abundance of physiological processes, including sensation¹, cognition², muscle contraction³, and metabolism⁴. Consisting of more than 800 receptor types, GPCRs are the largest family of membrane receptors in the human body⁵. Once activated by external stimuli including small molecule neurotransmitters, peptides, lipids, ions, and light, the GPCR undergoes a conformational change which allows G-protein and/or β -arrestin binding⁶⁻⁸. G-protein binding catalyzes the activation of a downstream signaling cascade, where, depending on the G-protein subtype, can cause a change in the intracellular concentration of second messengers, such as cAMP, IP₃, Ca²⁺, and diacylglycerol, while β -arrestin binding can cause internalization of the GPCR^{9,10} (Figure 1).

GPCR signaling malfunction is involved in a large number of diseases, such as heart disease¹¹, depression¹², dementia¹³, Alzheimer's disease¹³, Parkinson's disease¹³, and Huntington's disease¹³. Approximately 35% of drugs approved by the U.S. Food and Drug Administration (FDA) target GPCRs¹⁴. Additionally, many neuromodulators and neurotransmitters are GPCR agonists, such as glutamate, GABA, acetylcholine, serotonin, dopamine, and various neuromodulating peptides. GPCRs, therefore, play critical roles in modulating the activity of the nervous system. GPCR activity is mainly regulated by the spatiotemporally-regulated release of endogenous agonists. Therefore, there is particular interest in studying the spatiotemporal dynamics of endogenous GPCR agonists to understand how the precise timing and spatial localization of GPCR activation affects the downstream signaling cascade and overall biological response.

Overview of genetically encoded sensors for detecting endogenous GPCR agonists

To gain information about the spatiotemporal dynamics of GPCR activation in biological models, a variety of genetically encoded sensors have been engineered to detect GPCR agonists. Genetically encoded GPCR sensors are advantageous over non-genetically encoded bioanalytical methods, such as microdialysis¹⁵ and fast scanning cyclic voltammetry¹⁶⁻¹⁹, for GPCR agonist detection, because genetic encoding offers cell-type specificity when imaging in the animal brain and can image at cellular or even subcellular resolution. Genetically encoded GPCR sensors, therefore, can be used to determine the precise localization and release dynamics of GPCR agonist release.

Genetically encoded sensor designs are mainly based on the agonist-induced conformational change of the third intracellular loop, G-protein or β -arrestin binding, or the change of concentration of second messengers (Figure 1). The readout of these sensors can be the activation of a fluorescent protein, an enzymatic reaction, or the transcription of a reporter gene.

Genetically encoded GPCR sensors can be divided into two main categories: those that offer real-time information about GPCR agonist-induced activation and those that

integrate the GPCR agonist signal, leaving a permanent quantifiable signal over a large region of the brain (Figure 2). Real-time sensors are advantageous because they offer important information about the signaling dynamics of GPCRs; however, they can only detect agonists in a small brain region due to imaging challenges. Integrators, due to their permanent signals, can be used to determine the localization of GPCR agonists in a large area of the brain at cellular resolution but cannot be used to study signaling dynamics. In this review, we'll discuss the recent advancement of genetically encoded real time sensors and integrators for detecting GPCR agonists. We will also discuss how these two categories of sensors can be used together to provide detailed information about the spatiotemporal release of GPCR agonists in a whole brain at cellular resolution.

Genetically encoded real-time sensors

A variety of genetically encoded real-time sensors for detecting GPCR agonists have been recently developed using five major strategies: 1) intra or intermolecular resonance energy transfers; 2) circularly permuted green fluorescent protein (cpGFP) insertion into periplasmic binding proteins; 3) cpGFP inserted into the third intracellular loop of a GPCR; 4) translocation assays; and 5) detection of second messengers following GPCR activation. Real-time sensors are beneficial for studying the signaling kinetics of neurotransmitters and neuromodulators; therefore, these sensors benefit from having fast on- and off- rates. In this review, we will focus on the various designs of real-time sensors but will not discuss in detail the kinetics of real-time sensors. Readers can see reference²⁰ for time constant values of many of the sensors discussed here.

FRET and BRET-based sensors

Fluorescence resonance energy transfer (FRET)-based GPCR agonist sensors use a donor and acceptor fluorescent protein, where the donor's emission spectrum overlaps with the acceptor's excitation spectrum. FRET efficiency is inversely proportional to r raised to the 6th power, where r is the distance between the donor and acceptor fluorophore. Therefore, FRET efficiency can be highly sensitive to GPCR activation-dependent conformational change. The FRET fluorescent protein pairs can either be placed on different locations on the same GPCR (intramolecular FRET) or one protein can be fused to the GPCR and the other can be fused to a protein interactor, such as a G-protein mimic or β -arrestin (intermolecular FRET). Bioluminescence resonance energy transfer (BRET) is similar to FRET, except that a bioluminescent protein acts as the energy donor to activate the fluorescent protein acceptor²¹.

Here, we will mainly discuss the FRET- and BRET-based sensors that can potentially be used for detection of GPCR agonists in vivo at cellular resolution. For a more comprehensive review of FRET- and BRET-based GPCR sensors, see reference²¹. We will divide FRET and BRET sensors into four main categories: 1) those that detect G-protein

activation; 2) those that detect β -arrestin recruitment; 3) those that detect intramolecular GPCR conformational changes; and 4) those that detect the conformational change of binding proteins.

BRET and FRET sensors based on G-protein binding and activation

To detect G-protein activation, a bioluminescent protein, such as *Renilla* luciferase (Rluc) or the smaller luciferase enzyme, Nano Luciferase (NLuc)²², fused to a GPCR can act as the BRET donor and a fluorescent protein, such as Venus or yellow fluorescent protein (YFP), fused to a $G\alpha$ -protein or a $G\alpha$ -mimic can act as the BRET acceptor. When the GPCR is activated by its agonist, the $G\alpha$ -protein/mimic binds to the intracellular portion of the GPCR, thereby bringing the two BRET pairs into proximity where energy transfer can take place (Figures 3A and 3B). BRET-based GPCR sensors have been used to detect GPCR activation at subcellular resolution in cell cultures to observe activation at the plasma membrane, Golgi apparatus, and endosomes²³.

G-protein-based BRET sensors have been developed for several GPCRs, including the β -adrenergic 2 receptor (β 2AR), α -adrenergic 2 receptor (α 2AR), vasopressin 2 receptor (V2R), sensor neuron specific receptor, and thromboxane A2 receptor. In these sensors, a BRET acceptor, GFP10, was fused to G-proteins and a BRET donor, RLuc was fused to the GPCR²⁴ (Table 1). These sensors can monitor agonist-induced desensitization in real-time²⁴.

Conformation-specific nanobodies engineered to bind to active GPCRs have been used as $G\alpha$ -mimics and are advantageous in their easy expression in cell culture and high affinities for active GPCRs. These nanobodies have been used in BRET-based sensors, where one BRET pair is attached to the nanobody and the other is attached to the corresponding GPCR to give an increase in BRET signal upon GPCR activation²⁵ (Figure 3B). Alternatively, a nanobody that binds to the inactive GPCR, such as Nb6 for the kappa opioid receptor (KOR), can see a decrease in BRET signal upon GPCR activation²⁵. Although this tool has not been used in any in vivo applications, it was able to observe the different conformational states of KOR induced by different ligands²⁵. A disadvantage of nanobody-based BRET sensors is that only a few GPCRs have conformation-specific nanobody binders. These nanobody-GPCR pairs include Nb39 and Nb6 for opioid receptors (ORs)^{26,27}, Nb80 for β 2AR²⁸, Nb.AT110 for angiotensin receptors²⁹, and Nb9-8 for the M2-muscarinic receptors³⁰ to name a few.

For GPCRs without conformation-specific nanobodies, miniG proteins can be used as G-protein mimics. MiniG proteins are derived from the $G\alpha$ subunit of the G-protein complex, but with the membrane anchor and $G\beta\gamma$ binding surface removed, thereby allowing easier protein expression than the wild-type $G\alpha$ protein³¹. MiniG-based BRET sensors have been developed by fusing the miniG protein to the BRET acceptor, such as Venus, and fusing the donor, such as Rluc or NLuc, to the GPCR (Figure 3B). In the presence of agonist, miniG can bind to the GPCR and cause an increase in BRET signal. MiniG-based BRET sensors are very versatile, having been designed for several $G\alpha$ -protein types: $G\alpha_i/o$, $G\alpha_s$, $G\alpha_{12/13}$, and $G\alpha_q/11$ ²³. Consequently, miniG protein-based BRET sensors can be

used for a wide variety of GPCRs in cell culture; however, to our knowledge, miniG protein-based BRET sensors have not been used in vivo.

Fusion of the GPCR to the BRET pair involves genetically altering the GPCR; consequently, these types of sensors cannot detect endogenous GPCR activation. Alternatively, BERKY is a BRET-based GPCR agonist sensor that does not alter the endogenous GPCR and can, therefore, detect endogenous GPCR agonists (Figure 3C). BERKY is a single protein chain where the BRET donor, NLuc, is bound to a membrane anchoring sequence and the acceptor, YFP, is bound to a synthetic peptide called KB-1753. The donor and acceptor BRET pairs are separated by an ER/K α -helix linker. Once the $G\alpha$ -protein is activated, KB-1753 interacts with the GTP-bound $G\alpha$ -protein, bringing YFP to the membrane and increasing BRET. BERKY sensors have also been developed for $G\alpha_q$, $G\alpha_{13}$, $G\beta\gamma$, and Rho to probe the activation of endogenous GPCRs³². Additionally, BRET and FRET pairs inserted into the $G\alpha\beta\gamma$ heterotrimer have also proven useful in measuring endogenous GPCR activation. The proximity of the $G\alpha$ and $G\beta\gamma$ subunits changes based on the activation-state of the GPCR, thereby changing the energy transfer efficiency and indicating GPCR activation^{33–39} (Figure 3D). Finally, NanoBRET was engineered by fusing NLuc to GRK3ct, a GPCR kinase, and Venus to $G\beta\gamma$ to observe G-protein activation⁴⁰. While the above two types of BRET sensors can monitor GPCR activation without direct fusion to the GPCR, they can result in false positives since the sensor activation is dependent on general GPCR activity and not the specific activity of the GPCR of interest. Consequently, control studies without the expression of the GPCR of interest is needed to ensure the sensor activation observed is due to the GPCR of interest.

The advantage of G-protein binding and activation-based BRET sensors is that these sensors have been illustrated for a wide variety of G-protein subtypes ($G\alpha_i$, $G\alpha_o$, $G\alpha_s$, $G\alpha_q$, $G\alpha_{12}$, and $G\alpha_{13}$), enabling the detection of a large number of GPCR agonists²¹. A disadvantage of BRET sensors is that they have poor sensitivity for some G-protein subtypes²¹. However, a higher sensitivity can be achieved by systematic optimization of the insertion points for the BRET pairs. This systematic optimization was performed in the BRET biosensors TRUPATH, where fourteen BRET-based GPCR sensors were developed⁴¹. Still, individual optimization for each BRET biosensor is strenuous, making it desirable to have a more universal sensor design platform.

In a similar design to the BRET sensors, FRET sensors have been developed with the fluorescent proteins CFP and YFP, where one FRET pair is attached to the GPCR and the other is attached to the G-protein or G-protein mimic. The activation of the GPCR causes YFP to come in proximity to CFP and energy transfer can occur (Figure 3B). This sensor type has been developed for the α_{2A} AR^{42,43}, the muscarinic-M4 receptor⁴³, the A_{2A} -adenosine receptor⁴⁴, and the β -adrenergic receptor¹⁴⁴ (Table 1). eCFP and eGFP were used as FRET pairs for the bradykinin receptor type 2⁴⁵. FRET has faster kinetics than BRET; however, FRET's need for excitation of the donor fluorescent proteins could lead to cell damage or photobleaching. BRET's use of a chemical substrate, therefore, can be less destructive for the cell.

BRET and FRET sensors based on GPCR- β -arrestin interaction

BRET sensors have been designed based on the intermolecular interaction between β -arrestin and the active GPCR's C-terminus. RLuc as the BRET donor has been fused to the C-terminus of a GPCR and YFP as the BRET acceptor has been fused to β -arrestin2 to detect the activation of β 2AR⁴⁶, the thyrotropin releasing hormone receptor^{47,48}, and the oxytocin receptor⁴⁹ (Figure 3E). Additionally, RLuc and GFP from *Renilla reniformis* have been attached to a GPCR and the transmembrane domain, respectively, where β -arrestin-induced internalization causes a change in energy transfer efficiency⁵⁰. This sensor type has been designed for the angiotensin II receptor type 1, melanocortin type 4 receptor, and the V2R and can be used to interrogate receptor activation as well as GPCR recycling events⁵⁰. The GFP and RLuc BRET pair fused to V2R and β -arrestin, respectively, was also used to interrogate how palmitoylation of the V2R carboxyl tail affects β -arrestin recruitment⁵¹.

A BRET sensor with improved efficiency was designed, in which the V2R is fused to hRLuc/RLuc8 and β -arrestin2 is fused to a modified form of GFP, called 2GFP⁵². Cells expressing the above BRET sensor were transplanted into a mouse kidney and a transparent, plastic window was fitted into the skin and body wall adjacent to the kidney. The window enables BRET to be measured in deep tissues⁵². The same BRET pair was used to design a BRET-based GPCR sensor by fusing β 2AR to RLuc and β -arrestin2 to 2GFP⁵³. Due to poor light penetration in deep tissues, sufficient BRET signal was only achieved in the testes of mice and not in other tissues⁵³. In future work, a red shifted BRET pair can improve BRET detection in deep tissues.

BRET sensors have also been developed to detect the specific conformational change found in β -arrestin upon recruitment to the GPCR. To design this sensor, RLuc and YFP have been attached to either terminus of β -arrestin^{54,55}, as well as NLuc and a red shifted fluorescent protein⁵⁶ (Figure 3F). Different from the above intermolecular BRET sensors, these BRET sensors detect the intramolecular BRET efficiency change due to arrestin's conformational change. Compared to intermolecular BRET and FRET, intramolecular BRET and FRET have a smaller change in the distance between the BRET/FRET pairs and, therefore, a lower signal dynamic range. However, these intramolecular β -arrestin-based BRET sensors can observe the different types of active conformations that β -arrestin adopts and have been used for a variety of GPCRs, such as the angiotensin 1 receptor, vasopressin receptors 1 and 2, β -adrenergic receptors 1 and 2, muscarinic 1 receptor, chemokine receptor 5, delta opioid receptor, serotonin receptor, platelet-activating factor receptor, chemokine receptor type 7, and the glucagon-like peptide-1 receptor⁵⁴⁻⁵⁶.

Similar to the G-protein-based FRET design, CFP and YFP have been used as FRET pairs. One FRET pair is attached to the GPCR and the other is attached to β -arrestin1 or 2 (Figure 3E). These sensors have been applied in cell cultures to determine the time it takes β -arrestin2 to be recruited to the parathyroid hormone 1 receptor⁵⁷ and β 2AR⁵⁸. They have also been applied to determine the differential recruitment of β -arrestin1 and 2 to P2Y2R⁵⁹.

Compared to the G-protein based sensors, β -arrestin-based BRET and FRET real time sensors can generate a stable on-signal, because the GPCR- β -arrestin interaction

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occurs on the order of minutes to tens of minutes, while the GPCR-G protein interaction is much more transient, on the order of seconds. Consequently, β -arrestin-based BRET and FRET real time sensors are also limited by their off kinetics. Additionally, not all GPCRs strongly recruit β -arrestin. Therefore, G-protein based and β -arrestin-based BRET and FRET sensors should be chosen depending on their different applications.

BRET and FRET sensors based on the conformational changes in the GPCR

G-protein-based BRET sensors require knowledge of the specific G-protein type that binds to the GPCR which is not always known. β -arrestin-based BRET sensors need to use GPCRs that couple strongly to β -arrestin which, as stated previously, is not true for all GPCRs. To address these limitations in sensor designs, BRET and FRET sensors have been developed that utilize the conformational change of the GPCR upon activation rather than utilizing the intermolecular interactions of the G-proteins and β -arrestin with the GPCR. Theoretically, these types of sensors should be more universal given that all GPCRs undergo a conformational change upon activation. CFP can be inserted into the third intracellular loop of the GPCR where the major conformational change takes place upon GPCR activation, and YFP can be fused to the C-terminus of the GPCR, or the other way around (Figure 3G). This strategy has been used for the PTHR and α_{2A} AR to observe the different conformational changes of GPCRs induced by different types of ligands^{57,60}. These sensors have been developed for other types of GPCRs as well⁶¹⁻⁶³ (Table 1). These intramolecular sensors based on the conformational changes in the GPCR have a lower signal dynamic range than the intermolecular BRET and FRET sensors.

BRET and FRET for detecting the conformational change of binding proteins

Although the majority of FRET/BRET sensors for GPCR agonists involve GPCRs or G-proteins in their design, the glycine FRET sensor (GlyFS) and the glutamate sensor (FLIPE) use soluble protein domains that can bind to glycine and glutamate in their sensor designs. In GlyFS, two FRET pairs, enhanced cyan fluorescent protein (ECFP) and Venus, were both attached to a rationally-designed soluble glycine-binding protein⁶⁴. Glycine binding to the sensor reduces the FRET between the two fluorescent proteins (Figure 4A). GlyFs was used in acute hippocampal slices to reveal changes of extracellular glycine concentrations during development, enrichment of glycine outside of the synapse, and the increase of extracellular glycine from neuroplasticity⁶⁴. The glutamate sensor, FLIPE, has a similar design to GlyFs, where ECFP and Venus were inserted into a soluble glutamate binding protein, ybeJ⁶⁵. FLIPE was used to show that glutamate uptake in the cytosol of PC12 cells has a minimal effect on overall cytosolic glutamate levels⁶⁵.

Summary of advantages and limitations of BRET- and FRET-based GPCR sensors

BRET- and FRET-based sensors have been designed for a wide variety of GPCRs, mainly based on the activated GPCR's interaction with G-proteins or β -arrestin or the conformational change of the GPCR to indicate GPCR activation. BRET and FRET designs are highly versatile and can be generally applied to most GPCRs by using different types of $G\alpha$ -proteins or β -arrestin. However, BRET is limited by the low signal intensity of luminescence which requires longer BRET acquisition time. FRET is limited in its signal-to-noise ratio (SNR), which is most commonly <100%, due to potential spectral overlap or non-proximity-based energy transfer. Here, we define SNR for real-time sensors as either the agonist-induced change of fluorescence over the initial fluorescence or the signal over background ratio. Additionally, these sensors take extensive engineering to determine the correct placement of the donor and acceptor, so the specific conformational-induced change of the distance of the FRET donor-acceptor pair can be maximal. Lastly, BRET and FRET requires the use of two fluorescence channels, limiting the use of multiple fluorescence markers and sensors.

To overcome these limitations, single fluorescent-protein based systems have been developed by using cpGFP. Circular permutation involves the fusion of the original N and C terminus of a protein and the creation of a new N and C terminus in a different position on the protein. cpGFP was engineered to have its new N and C terminus near the fluorophore pocket, thereby making the fluorophore more exposed to the solvent environment. Environmental changes around the cpGFP fluorophore can cause a change in the fluorescence emission intensity. This resulting change in fluorescence is instantaneous and reversible, making cpGFP a good tool for real-time detection of neurotransmitters and neuromodulators. cpGFP as a real time sensor has been used in two ways: 1) cpGFP inserted into periplasmic binding proteins; and 2) cpGFP inserted into the third intracellular loop of a GPCR.

Single-color fluorescent sensors based on cpGFP insertion into periplasmic binding proteins

Fluorescent sensors for several neuromodulators and neurotransmitters have been engineered using periplasmic binding proteins, receptors found in bacteria that undergo a conformational change in the presence of a variety of small molecules. Specifically, these are Venus-fly trap-like proteins that close in the presence of their binding ligand. cpGFP was inserted into periplasmic binding proteins to create a series of neurotransmitter and neuromodulator sensors, named Sensing Fluorescent Reporters (SnFRs). For the SnFR design, ligand binding causes a conformational change in the periplasmic binding proteins which causes a change in the fluorophore environment of cpGFP, resulting in a fluorescence change (Figure 4B). SnFRs have been developed to detect several GPCR ligands, including glutamate, GABA, acetylcholine, and serotonin, with SNR from 1 and 25 (Table 1). The glutamate version of SnFR, called iGluSnFR, has been used to detect glutamate in *C. elegans*, zebrafish, mice, and ferrets⁶⁶. The original iGluSnFR used circularly permuted enhanced GFP (cpEGFP) but new versions have since been developed with circularly permuted super folder GFP (SF-iGluSnFR) which has higher expression level and

fluorescent signals⁶⁷. Additionally, different colored variants and variants with a variety of affinities for glutamate have been developed⁶⁷. More recently, iGluSnFR3 was developed through directed evolution and has a higher SNR, dynamic range, expression, and photostability than SF-iGluSnFR⁶⁸. Finally, faster versions of iGluSnFR have been developed to resolve individual glutamate release events in rat hippocampal slices⁶⁹. The SnFR engineering technique has also been used to engineer a sensor for an important inhibitory neurotransmitter, GABA. iGABASnFR has been used in vivo to track the concentration of GABA in the mitochondria in zebrafish models and track the GABA signaling dynamics during interictal spikes and seizures in mice⁷⁰. Recently, iAChSnFR was developed to detect acetylcholine release in mice, zebrafish, flies, and *C. elegans*⁷¹.

Naturally occurring periplasmic binding proteins do not exist for the majority of neurotransmitters and neuromodulators, limiting the generalizability of the SnFR design; however, engineering of the binding pocket of existing periplasmic binding proteins could lead to the development of new sensors. Using machine learning, researchers re-engineered the binding pocket of iAChSnFR's periplasmic binding protein to bind to serotonin, generating iSeroSnFR. iSeroSnFR was used to detect serotonin release in mice during different behavioral assays, such as fear conditioning, social interaction, and sleep/wake transitions⁷². Further, site saturated mutagenesis of a periplasmic binding protein was performed to select for a variant that can bind specifically to nicotinic acetylcholine receptor agonists, thereby creating iNicSnFR. iNicSnFR was used to determine the rate at which different nicotinic receptor agonists leave the endoplasmic reticulum⁷³. The iNicSnFR was subsequently mutated using site saturated mutagenesis and rational design to select for variants that specifically bind to S-methadone and not cholinergic ligands; this sensor is called iS-methadoneSnFR⁷⁴.

To further illustrate the versatility of the SnFR design platform, a new set of SnFRs were developed recently for a series of smoking-cessation drugs: dianicline, cytisine, 10-fluorocytisine, and 9-bromo-10-ethylcytisine. These sensors are called iDrugSnFRs and have been used to determine the differing rates of membrane crossing of smoking-cessation drugs, a property that affects the drug's pharmacokinetics⁷⁵.

The application of the SnFR sensors has been demonstrated in a variety of model organisms to interrogate the neurobiology underlying various neurotransmitter and neuromodulator-mediated signaling events. The wide applicability of SnFRs for in vivo experiments illustrates the usefulness and robustness of the sensor design.

Single-color fluorescent sensors based on cpGFP insertion into the third intracellular loop of GPCRs

GPCR activation-based (GRAB) and Light sensors were developed by inserting cpGFP into the third intracellular loop of a GPCR. Upon agonist binding, the third intracellular loop undergoes a conformational change that changes the fluorophore environment of cpGFP, resulting in a fluorescence change (Figure 5). These sensors have been developed to detect dopamine^{76,77}, endogenous opioid peptides^{76,78}, serotonin^{79,80}, noradrenaline^{76,81,82}, endocannabinoids⁸³, adenosine^{84,85}, and acetylcholine^{86,87} with most

SNRs ranging between 2 and 10 (Table 1). For some versions of these sensors, portions of the third intracellular loop have been truncated to allow a larger fluorescence change. Different linkers connecting cpGFP to the third intracellular loop have also been screened to change the sensitivity and SNR of the sensors. Both rational design and site-saturated mutagenesis have been used to engineer these sensors and improve their SNR and overall brightness⁸⁸. In addition, red-shifted versions have been developed for both GRAB and Light sensors to allow for multiplexed imaging of different neurotransmitters and neuromodulators at a single time^{89,90}. For a more comprehensive overview of the engineering used to design the GRAB/light sensors, see reference⁸⁸.

GRAB sensors have been used in a variety of in vivo models, including flies, zebrafish, and mice⁷⁷. Light sensors have been illustrated in vivo in mice and rats⁷⁶. These sensors were used in animal models to reveal important information about neurotransmitter and neuromodulator-mediated neurobiology. To highlight a few applications, the GRAB_{Ado} sensor was used to study sleep homeostasis and found that adenosine is at high concentration during rapid eye movement (REM) sleep but low outside of the REM sleep cycle⁸⁴. Using GRAB_{Ado}, researchers also found that the activation of glutamatergic neurons resulted in an increase of adenosine. Further, dLight helped researchers find that heroin activates dopaminergic neurons located in the medial area of the VTA⁹¹, dopamine is involved in relapse and alcohol-seeking behaviors⁹², and that the pattern of morphine exposure affects dopamine release in the nucleus accumbens⁹³. GRAB_{DA} was used to show that alcohol, but not fat, significantly increased dopamine in the nucleus accumbens and that fat, but not alcohol, increased dopamine in the dorsal striatum⁹⁴.

Translocation assays

Rather than genetically altering the GPCR, translocation assays can be performed to indicate GPCR activation. When a GPCR is activated, GPCR-interacting proteins can translocate from the cytoplasm to the cell membrane or from the membrane to the cytoplasm. Consequently, these interacting proteins, such as β -arrestin⁹⁵⁻⁹⁷, protein kinase C⁹⁵, and G-proteins/mimics^{98,99} have been tagged with fluorescent proteins and their agonist-induced translocation has been detected with high magnification imaging. Total internal reflection fluorescence microscopy has also been used to analyze the fluorescence ~100 nm from the cell surface, allowing precise measurements of membrane fluorescence⁹⁸. A major limitation of translocation assays is their low SNR.

Detection of second messengers

The activation of a GPCR can increase or decrease the concentration of second messengers, such as cAMP, IP₃, Ca²⁺, and diacylglycerol. Consequently, sensors that can detect the concentration of second messengers can be used to detect GPCR activation by an agonist. These sensors can be generalizable to many GPCRs, because the activation of most GPCRs will cause a change in the concentration of at least one of the second messengers previously mentioned. However, second messaging events can also be caused by non GPCR-mediated signaling, potentially giving false positive signals. It is, therefore,

better to use GPCR agonist sensors that detect more upstream signals, such as conformational changes in the GPCR or β -arrestin/G-protein binding, when available.

One of the most widely used second messenger sensors, GCaMP, was first developed in 2001 to detect intracellular calcium changes in living cells¹⁰⁰. GCaMP involves the fusion of a calcium-dependent protein binding pair, M13 and calmodulin, to either terminus of cpGFP. In the presence of calcium, these proteins bind, changing the fluorophore environment of cpGFP and resulting in a fluorescence change (Figure 6A). After 20 years of dedicated work in sensor optimization, improved versions of GCaMP have been published that have better SNR, heightened brightness, faster kinetics, and heightened calcium sensitivity than the original version^{101–104}. GCaMP has been used in a variety of model organisms in vivo, such as *C. elegans*^{102,104}, flies^{101–104}, mice^{101–104}, and zebrafish^{101,102}. Further, GCaMP has been used in vivo to detect calcium spiking in axons, illustrating its subcellular resolution¹⁰⁵. The impact of GCaMP on the field of neuroscience cannot be understated. Because this review focuses on neuromodulators and not calcium sensors, a more extensive review of GCaMP and its different optimized versions, as well as other genetically encoded calcium sensors, can be found in references^{106,107}.

A FRET sensor used to detect cAMP, called CAMYEL, was developed by inserting a protein that interacts with cAMP, Epac, in between two BRET pairs, YFP and RLuc¹⁰⁸. Upon the binding of cAMP to CAMYEL, there is a quantifiable decrease in BRET signal (Figure 6B). This tool was used to discover that sphingosine 1-phosphate can increase the amount of intracellular cAMP that is stimulated by isoproterenol and prostaglandin E₂¹⁰⁸. In addition to CAMYEL, many other FRET sensors for cAMP detection have been developed and are summarized in reference¹⁰⁹. Epac was also inserted into cpGFP, where cAMP binding to cpGFP causes a change in cpGFP fluorescence intensity¹¹⁰ (Figure 6C).

A fluorescent sensor for diacylglycerol was developed by fusing cpGFP to an isoform of protein kinase II that only responds to diacylglycerol¹¹¹ (Figure 6D). This diacylglycerol sensor was, then, improved by replacing cpGFP with the brighter mNeonGreen fluorescent protein¹¹⁰. Additionally, FRET was used to develop a sensor for protein kinase A activity, an important protein in downstream GPCR signal transduction¹¹².

GPCR agonist integration sensors

Integration-based GPCR agonist sensors integrate the GPCR agonist signal into a permanent and measurable readout for further analysis post mortem. Integration sensors have complementary strengths to real time sensors by enabling the examination of the localization of GPCR agonists across a large area of interest or the whole brain to interrogate a GPCR's effect on neuronal circuitry globally. Integration-based GPCR agonist sensors have been engineered for a wide variety of GPCRs and can be divided into three main categories based on their signal output: 1) transcriptional activation; 2) cpGFP fluorophore formation; and 3) split protein complementation sensors.

A transcriptional activation-based GPCR sensor uses unique transcription factors and promoters that are orthogonal with the cellular system to activate a reporter gene upon agonist binding. The reporter genes commonly code for fluorescent proteins, so that activation of the sensors by GPCR agonists will fluorescently label the cells. Transcription-based GPCR sensors were first developed in 2008, in which the C-terminus of the V2R was fused to the tTA transcription factor via the N1a tobacco etch virus (TEV) protease cleavage site (TEVcs), and β -arrestin2 was fused to TEV protease¹¹³ (Figure 7A). In the presence of the V2R agonist, the β -arrestin2 fused to TEV protease will bind to the C-terminus of V2R, bringing the protease in proximity to TEVcs for cleavage to take place, releasing tTA. No longer tethered to the membrane, tTA can then translocate to the nucleus and activate a tTA-dependent reporter gene. This sensor design, called Tango, was developed for 89 other GPCRs, illustrating the robustness of the technique¹¹³. It is important to note, however, that the V2R's C-terminal tail needs to be added to the other GPCRs to enhance β -arrestin2 recruitment and consequently increase the SNR. Here, we define SNR of integrators as the signal over background ratio.

To apply Tango to the majority of the druggable GPCRs within the human genome, PRESTO-Tango was developed¹¹⁴. PRESTO-Tango uses a "modular design strategy" to efficiently design Tango-based GPCR sensors for over 300 GPCRs. The researchers, then, demonstrated these 300+ Tango-based sensors can be used to test compounds against nearly the entire druggable human GPCR genome. They tested two compounds against 133 GPCR-Tango targets and showed that one compound, LSD, has activities against GPCRs that were not previously known¹¹⁴. Additionally, they screened 91 GPCR-Tango targets against FDA-approved drugs and found that the diabetes drug, nateglinide, has activity against MRGPRX4, a receptor thought to be involved in pain and itch¹¹⁴.

The PRESTO-Tango system represents one of the most widely applied high-throughput screening platform for GPCR ligands. However, its application in detecting GPCR ligands has so far been limited to cell cultures. In the PRESTO-Tango system, the TEV protease has basal cleavage activity that could accumulate over time in animal models, reducing the SNR. Additionally, Tango activation requires hours of GPCR agonist incubation due to the low catalytic efficiency of the TEV protease. To overcome these limitations, a light-activated Tango was developed in 2017 called inducible Tango (*i*Tango2)¹¹⁵.

Because integrators can start accumulating background signal immediately after the sensor is expressed, *i*Tango2's use of a light gating reduces the overall background of the system and allows signal to accumulate only in the light window. Additionally, light can be used as a temporal gating to record signaling events during a specific behavior. The *i*Tango2 design differs from Tango in two main ways (Figure 7B). First, the TEV protease is split into two halves, where one is tethered to β -arrestin2 and the other is attached to the C-terminus of the GPCR. Second, an engineered light sensing protein based on AsLOV2 is added to cage the TEVcs in the dark state. In light, the J α helix of the AsLOV2 protein undergoes a conformational change, uncaging the TEVcs to be accessible to the TEV protease. Agonist activation recruits β -arrestin2, bringing split TEV protease half TEV-C to the GPCR-TEV-N fusion protein, and allowing the split TEV protease to reconstitute. Therefore, only in the

presence of both light and agonist activation will the split TEV protease be reconstituted and can cleave the light-uncaged TEVcs, releasing tTA. iTango2 showed a SNR 20-fold higher than the original Tango system¹¹⁵. iTango2 has been developed for DRD2¹¹⁵, cannabinoid receptor type 1¹¹⁵, serotonin receptor 1A¹¹⁵, neuropeptide Y receptor type 1¹¹⁵, and the oxytocin receptor¹¹⁶ (Table 2). Additionally, iTango2 has been used to label dopamine-sensitive neuronal populations during reward-based learning in mice¹¹⁵ and was used to detect the change of CCR5 activity after learning in mice¹¹⁷.

Similar to iTango2, SPARK is a light-controlled, transcription-based GPCR agonist sensor that utilizes TEV protease cleavage to release a unique transcription factor that is tethered to the GPCR and can activate reporter gene expression^{118–120} (Figure 7C). SPARK differs from iTango2 in two main ways. First, it does not use split TEV protease and instead uses a C-terminal truncated TEV protease that has low affinity for the TEVcs. This allows protease cleavage only to occur when the TEV protease is brought to proximity to the TEVcs. Second, the TEVcs is caged by an evolved version of the AsLOV2 domain (eLOV). eLOV was evolved using directed evolution and was shown to have a 10x better light-to-dark signal ratio than the original AsLOV2¹²¹. In the initial testing of the SPARK design for detecting GPCR agonists, β 2AR was fused to the eLOV domain-caged TEVcs and a GAL4 transcription factor and β -arrestin was fused to the truncated TEV protease. To illustrate the generalizability of SPARK in GPCR agonist detection, SPARK was tested with eight different GPCRs. Six of these GPCRs had robust light- and ligand-dependent gene activation with five having ligand-dependent SNR above 15¹¹⁸ (Table 2).

One main advantage of the transcriptional system is the versatility of the signal output. To illustrate this, researchers used β 2AR-SPARK to drive luciferase expression, illustrating how SPARK can be used for high-throughput screening assays that need an easily quantifiable readout¹¹⁸. SPARK and iTango were compared using the β 2AR versions of both tools and transfection in HEK293T cells. SPARK was found to have a 16.4-fold higher ligand-dependent SNR¹¹⁸ and a higher sensitivity than iTango2¹¹⁸.

Recently, SPARK2 was developed using an evolved TEV protease, uTEV1 Δ , with improved catalytic efficiency¹²². SPARK's activation efficiency is limited to the number of cleavage events performed by the TEV protease. Due to the relatively slow catalytic rate of the TEV protease, SPARK requires 10 to 15 minutes of light and agonist stimulation to see a sufficient signal, limiting its application in situations where the GPCR agonist-induced protein-protein interaction (PPI) occurs on the order of seconds to a few minutes. SPARK2 uses uTEV1 Δ which has a faster cleavage rate than the original TEV protease, and consequently only 1 minute of light and agonist stimulation is needed to see a robust signal for β 2AR-SPARK2¹²².

cpGFP fluorophore formation-based GPCR agonist-integration sensors

SPARK and TANGO are multi-component systems, where the SNR is highly protein expression level-dependent and multiple viruses need to be delivered in animal model studies. Single protein chain-based integration systems could overcome some of the

limitations of the multi-component systems for interrogating GPCR signaling in a whole brain at cellular resolution. Recently, cpGFP-based sensors with a fluorophore formation-based mechanism have been developed to detect opioids for the mu, kappa, and delta opioid receptors (MOR, KOR, and DOR, respectively)¹²³ (Table 2). Named SPOTIT1, these tools take advantage of the ability of Nb39 to both bind to the active MOR and inhibit cpGFP fluorophore formation, thereby providing a sufficient protein switch for opioid detection. As shown in Figure 7D, in the absence of opioids, Nb39 remains bound to cpGFP and cpGFP fluorophore formation is inhibited. In the presence of opioids, Nb39 dissociates from cpGFP to bind to the active OR. This allows the cpGFP fluorophore to form and, therefore, a subsequent fluorescence increase. SPOTIT1 showed a SNR up to 12.5 in HEK293T cell culture and was shown to be effective in detecting a variety of opioid agonists, such as peptide agonists, synthetic agonists, and partial agonists. SPOTIT1 was also shown to be functional in neuronal cell culture, illustrating its potential applicability in animal models. However, SPOTIT1 had reduced brightness in neuronal cell culture¹²³. To improve the brightness of the SPOTIT sensors, SPOTIT2 was developed by adding four amino acids to the N-terminus of cpGFP to better encapsulate the fluorophore and increase its quantum yield¹²⁴. In HEK293T cell culture, SPOTIT2 is 11x brighter than SPOTIT1 and 2.7x brighter in neuronal culture. Our lab recently tested SPOTIT2 in mice and a significant difference in signal was detected between the morphine- and saline-treated mice (unpublished data), illustrating SPOTIT2's potential applications in animal models.

As an integration reporter, SPOTIT accumulates opioid signal over time until reaching a saturation point. Similarly, the background signal, which arises from the basal activity of the ORs, could also accumulate over time. To minimize the background signal accumulation, a chemical-gated version of SPOTIT, called SPOTon, was designed¹²⁵. In the SPOTon sensor design, SPOTIT is split into two components which are fused to a heterodimerizing PPI pair, FKBP and FRB; the OR was fused to FRB and cpGFP-Nb39 was fused to FKBP. In the presence of the small molecule rapamycin, FKBP and FRB heterodimerize, bringing cpGFP-Nb39 to the OR and creating a functional opioid sensor. Then, the opioid activated OR can recruit Nb39, allowing the cpGFP fluorophore to form. This sensor had a comparable opioid-dependent SNR as the SPOTIT sensors and strong rapamycin-dependence¹²⁵. Due to rapamycin's low blood brain barrier (BBB) penetrability, the engineering of a chemical-gated SPOTon that is activated with a BBB penetrable molecule would be needed for animal studies. Further, even though SPOTIT has only been demonstrated to detect opioid receptor agonists, the SPOTIT design could potentially be extended to other GPCR agonists by using other conformational specific binding nanobodies.

Split fluorescent protein complementation

Termed bimolecular fluorescence complementation (BiFC), split fluorescent protein (FP) components have low fluorescence when the split FP halves are apart, due to the inability of the fluorophore to form, and a high fluorescence when the components re-associate. Split FP can be designed to be proximity dependent by tuning the binding affinity of the split components, such that the complementation of the components only occurs when

they are brought into proximity. These proximity-dependent split FP can then be fused to PPI pairs to detect PPIs (Figure 7F).

Split GFP has been used extensively in BiFC assays to detect GPCR agonists. GFP has a 11-stranded β -barrel structure and, due to its very stable structure, can be split in a variety of positions. Tripartite GFP, where GFP is split at β -strand 10 and 11 to give three fragments: β 10, β 11, and β 1-9, has better SNR than a bipartite system due to reduced reassociation of the split fragments in the absence of PPI¹²⁶. Tripartite GFP was used to detect GPCR activation by fusing β 11 to the C-terminus of a GPCR, β 10 to β -arrestin, and expressing β 1-9¹²⁷. Agonist-induced GPCR activation recruits β -arrestin to the GPCR's C-terminus, allowing all three split fragments to re-associate and fluorophore formation to occur (Figure 7F). This assay, named Trio, was generated for the protease-activated receptor 1, β 2AR, neurokinin receptor, and MOR (Table 2). However, similar to Tango and SPARK, this system is limited to GPCRs that recruit β -arrestin and is a multiple component system.

Summary

Table 1 and 2 list various real-time sensors and integration sensors for a range of GPCR agonists discussed in this review. When choosing which existing sensor to use, one must consider both the SNR and affinity of the agonist for the sensor. Higher SNRs are generally preferred to provide a more robust and sensitive application in animal models. Sensors with a range of affinities for the agonist need to be evaluated in the specific animal model used for testing to determine which sensor can detect agonist concentrations with the highest dynamic range. While some sensors have been validated in animal models, many sensors have only been tested in cell cultures. For the latter, careful evaluations in animal models are needed before the sensors can be applied to study endogenous agonist release. In some cases, further engineering of the sensor SNR or affinity for ligand might be needed to achieve robust application in animal models.

Generally, real-time sensors have lower SNRs than integration sensors, partially due to the mechanism of the sensor design and also their transient detection, while integration sensors can accumulate signals over time to provide an end point readout. However, real time sensors can record the agonist-induced fluctuation of signal from the same cells; therefore, even a low percentage of signal change could potentially be detected if the background of the system is low. Integration sensors require higher SNRs than real time sensors, because their agonist-induced signal changes are not compared within the same cell but, instead, between different animals treated under different conditions. Typically, a minimum SNR of 5 or higher is required for integration reporters. Additionally, due to the protein level-dependence for integration sensors, single-component integration reporters are less variable and more advantageous than multi-component integration reporters.

Due to their complementary strengths and limitations, both real time and integration sensors are useful for studying endogenous GPCR agonist release and their effects on neuromodulation and other physiological processes. Genetically encoded real-time sensors of GPCR agonists are useful for interrogating the spatiotemporal dynamics of GPCR

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signaling. However, real-time imaging is limited to a small field of view. Cellular resolution integrators are useful for mapping GPCR agonist release in a large area, being able to interrogate whole-brain neuronal circuitry and label neurons exposed to the agonists for further interrogation. Integration sensors can be used first for unbiased search of the neuronal circuits affected by endogenous GPCR agonists across the brain. Real-time sensors can then be used to perform detailed characterizations of the spatiotemporal dynamics of endogenous GPCR agonist release in those circuits. The combined use of these sensors, therefore, can help expand our overall knowledge of GPCR-mediated neuronal signaling.

ACKNOWLEDGEMENTS AND FUNDING

This work is supported by the University of Michigan. W.W. is supported by NIH R21AG07200901, DP2MH132939, and Scialog grant 28431. K.K. is supported by NIH F31MH12915001.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

AUTHOR CONTRIBUTIONS

All authors contributed equally.

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Table 1: Real time sensors

Sensor Type	NT/NM detection	SNR	Affinity for ligand (EC50/IC50/Kd)	Model	References
FRET based	Acetylcholine	100%	600 nM carbachol	CHO, HEK	36,43
	Epinephrine/ norepinephrine	32%	12 nM-17 μ M	HEK, CHO, PC12	37,38,42– 44,57,58,6 0,62,63
	Bradykinin	60%	NA	HEK	45,61
	Adenosine	16%	NA	HEK	44
	Histamine	20%	~1 μ M	HeLa, MEF	33
	Lysophosphatidic acid	40%	NA	HeLa	34
	Parathyroid hormone	20%	16 nM	HEK	57
	Adenine and uridine nucleotides	4%	NA	HEK	59
	Glycine (GlyFS)	~20%	20 μ M	HEK, brain tissue	64
	Glutamate (FLIPE)	27%	0.6 μ M-1 mM	PC12, neuron culture	65
	Epinephrine/norepinephrine	300%	9.1 pM-120 nM isoproterenol	HEK, HeLa, neuron culture, and mice	23,24,32,3 5,39– 41,46,53– 56
	Acetylcholine	450%	0.77 μ M carbachol ~1 μ M	HEK	23,32,35,3 9,40,56

BRET based			acetylcholine		
	Angiotensin II	130%	1 nM to 4.2 nM	HEK	35,50,54-56
	Chemokine ligands 19 and 21	130%	NA	HEK	56
	Chemokine ligands 3, 4, 3-like 1	~18%	NA	HEK	39,55
	C-X-C motif chemokine 12	~38%	NA	HEK	39
	Calcitonin gene-related peptide	~20%	NA	HEK	39
	Prostaglandin E2	~55%	NA	HEK	39
	Vasoactive intestinal peptide	~25%	NA	HEK	39
	Secretin	~17%	NA	HEK	39
	Glucagon-like peptide-1 and glucagon	120%	NA	HEK	56
	Platelet-activating factor	3.6%	NA	HEK	55
	Serotonin	130%	NA	HEK	56
	Endogenous opioids	2.5%	0.09 nM salvinorin A 5.9 nM-580 nM DAMGO	HEK	25,39-41,55
	endocannabinoids	NA	49 nM-65 nM WIN 55,212-2	HEK	41
	Neurotensin	NA	34 pM-390pM	HEK	41
Adenosine	NA	~ 0.5 μM	HEK	23	

	Thyrotropin-releasing hormone	NA	8.8 nM	HEK COS-1	47,48
	Oxytocin	NA	NA	COS-7	49
	Vasopressin	130%	3 nM AVP	HEK and mice	24,39,50– 52,55,56
	Parathyroid hormone	6.8%	10 nM	HEK	54
	Dopamine	NA	~0.1 μ M	HEK	23,39,40
	Thromboxane	50%	9.6 nM U46619	HEK	24,35
	Sensory neuron-specific receptor	NA	NA	HEK	24
	Endothelin-1	NA	~1 nM	HEK	23
	Bradykinin	NA	NA	HEK	40
	Melanocortin	NA	NA	HEK	50
SnFRs	Glutamate	~25	0.6 μ M-600 μ M	HEK, neuron culture, C. elegans, zebrafish, mice, ferrets	66–69
	GABA	~1.7	30-110 μ M	HEK, neuron culture, mice, and zebrafish	70
	Acetylcholine	14	0.4 μ M -35 μ M	HEK, neuron culture, C. elegans, fly, zebrafish, mouse	71,73
	Serotonin	17	390 μ M	HEK, neuron culture, mice	72
	Dopamine	3.4	4 nM-130 nM	HEK, neuron culture, Fly, zebrafish, mice	77,89

GRAB	Epinephrine/ norepinephrine	2.3	83 nM-930 nM	HEK, neuron culture, Zebrafish, mice	⁸¹
	Serotonin	2.8	14 nM	HEK, neuron culture, flies, mice	⁷⁹
	Endocannabinoids	9.5	9 μ M 2-AG 0.8 μ M AEA	HEK, neuron culture, mice	⁸³
	Adenosine	~2.0	60 nM-3.6 μ M	HEK, neuron culture, mice	^{84,85}
	Acetylcholine	2.8	0.7 μ M	HEK, neuron culture, flies, mice	^{86,87}
	Lights	Dopamine	9.3	4.1 nM-1.6 μ M	HEK, neuron culture, mice, rats
Dynorphin		0.6	NA	HEK, mice	^{76,78}
Serotonin		0.8	26nM	HEK, neuron culture, mice	⁸⁰
Epinephrine/ norepinephrine		2.1	760nM	HEK, neuron culture, mice	^{76,82}

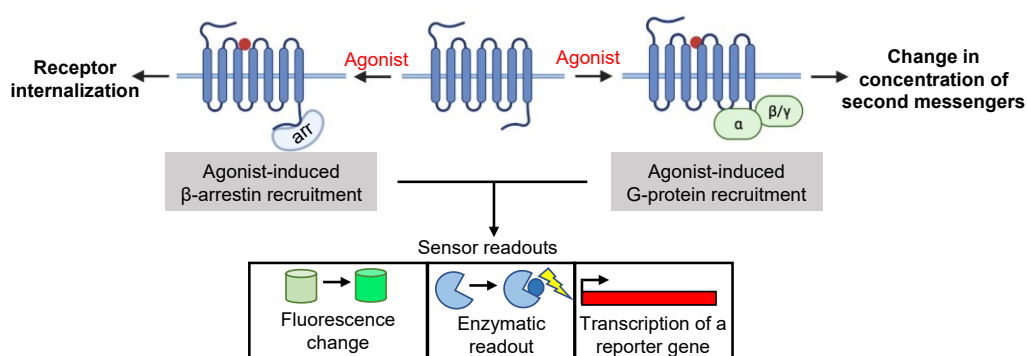
Table 2: Integrator sensors

Sensor Type	NT/NM detection	SNR	EC50	Model	References
TANGO	Has been developed for	1.3 to	NA	HEK, neuron	^{113,114}

	over 300 GPCRs	180		culture, Mouse	
iTANGO	Dopamine	8.9	~50 nM	HEK, neuron culture, mice	115,117
	Cannabinoid	~3.6	NA	Neuron culture	115
	Serotonin	~4.5	NA	Neuron culture	115
	Neuropeptide Y	~6.3	NA	Neuron culture	115
	Oxytocin	6.6	~2.5 μ M	HEK, neuron culture, mice	116
	Epinephrine/norepinephrine	24	52 nM iso	HEK	118,122
SPARK	Motilin	18	NA	HEK	118
	dopamine	19	NA	HEK	118
	Bombesin	37	NA	HEK	118
	Vasopressin	4.2	NA	HEK	118
	Angiotensin II	1.4	NA	HEK	118
	Opioid peptides	MOR: 13 KOR: 38 DOR: 2.7	15 nM fentanyl 0.77 μ M salvinorin A	HEK and neuron culture	123-125
SPOTIT					

Trio	Epinephrine/norepinephrine	NA	NA	HEK	127
	Neurokinin	NA	NA	HEK	127
	Opioid peptides	NA	NA	HEK	127

Table details: ~Estimated values, because raw values were not available in the publication. All values were rounded to 1-2 significant digits. For all SNR, the highest values were taken from the publications referenced. For the EC50/Kd values, a range of values were recorded from the publications referenced. EC50/Kd values for different versions of a sensor are included in the range. For those tested in multiple cell types, the SNR and EC50/Kd range are chosen for the cell type that gave the best SNR and lowest EC50/Kd. For FRET and BRET sensors, the agonist-induced percent change in BRET/FRET efficiency were recorded as either the agonist-induced change of fluorescence over the original fluorescence or the signal in the presence of agonist over the background signal without agonist. For the integrator sensors, the SNR was calculated as the signal in the presence of agonist over the background signal without agonist. For the light-gated integrators, the agonist-dependent SNR in the presence of light was recorded. The SNR of the intermolecular BRET sensors were not recorded in the table, because the majority of these sensors were reported to have “0” BRET without agonist, making it impossible to calculate a percent change. MEF, mouse embryonic fibroblast. HEK, human embryonic kidney cells. CHO, Chinese hamster ovary cells.



Graphical abstract: G-protein-coupled receptor (GPCR) agonist sensors utilize downstream GPCR signaling events, such as GPCR conformational changes and protein binding, to give a measurable sensor readout.

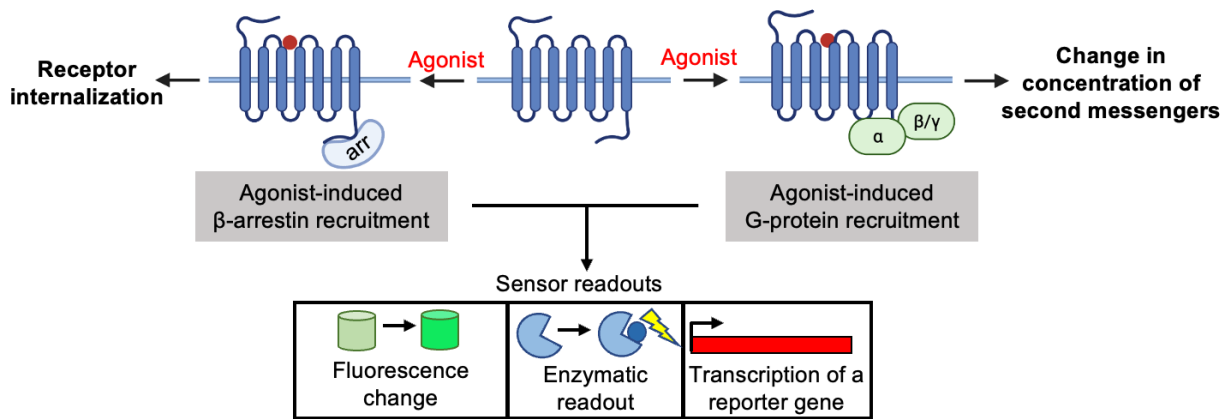


FIGURE 1 Schematic of GPCR signaling cascade. Agonist binding causes a conformational change in the GPCR which recruits G-proteins and/or β -arrestin to the GPCR. G-protein binding leads to a change in concentration of second messengers and β -arrestin binding causes internalization of the GPCR. These agonist-induced changes can be utilized in sensors, where they can cause the change of fluorescence, an enzymatic readout, or the transcription of a reporter gene.

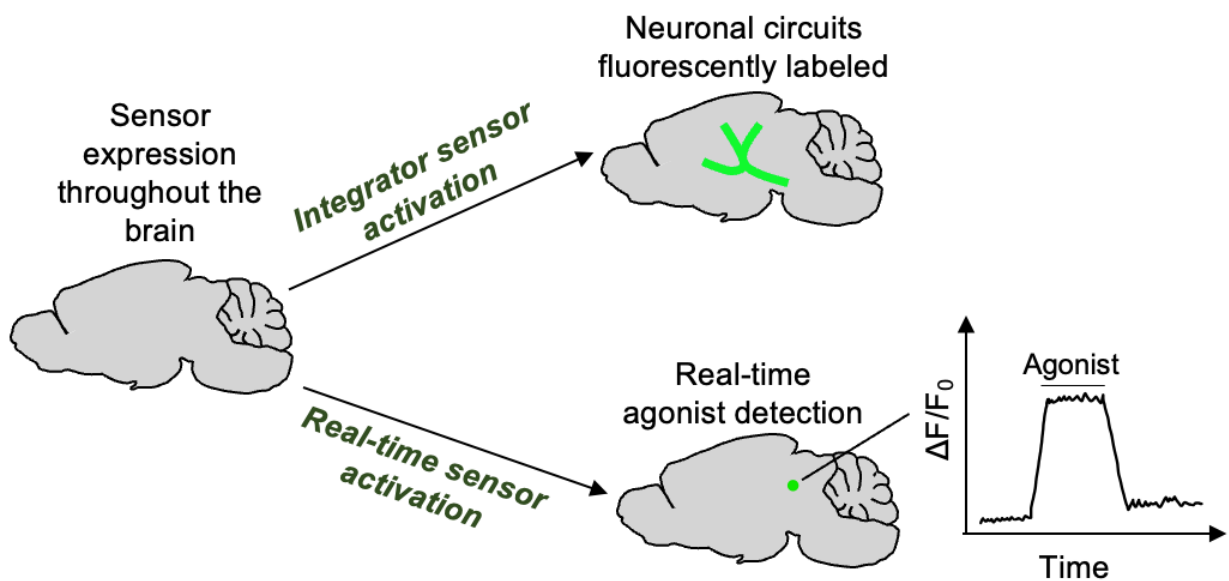


FIGURE 2 Integrators versus real time sensors. Integrators leave a permeant mark in the cells exposed to GPCR agonists, so large brain regions can be analyzed at cellular resolution. Real-time sensors can be used to observe the real-time agonist-induced neuronal activity in a small field of view.

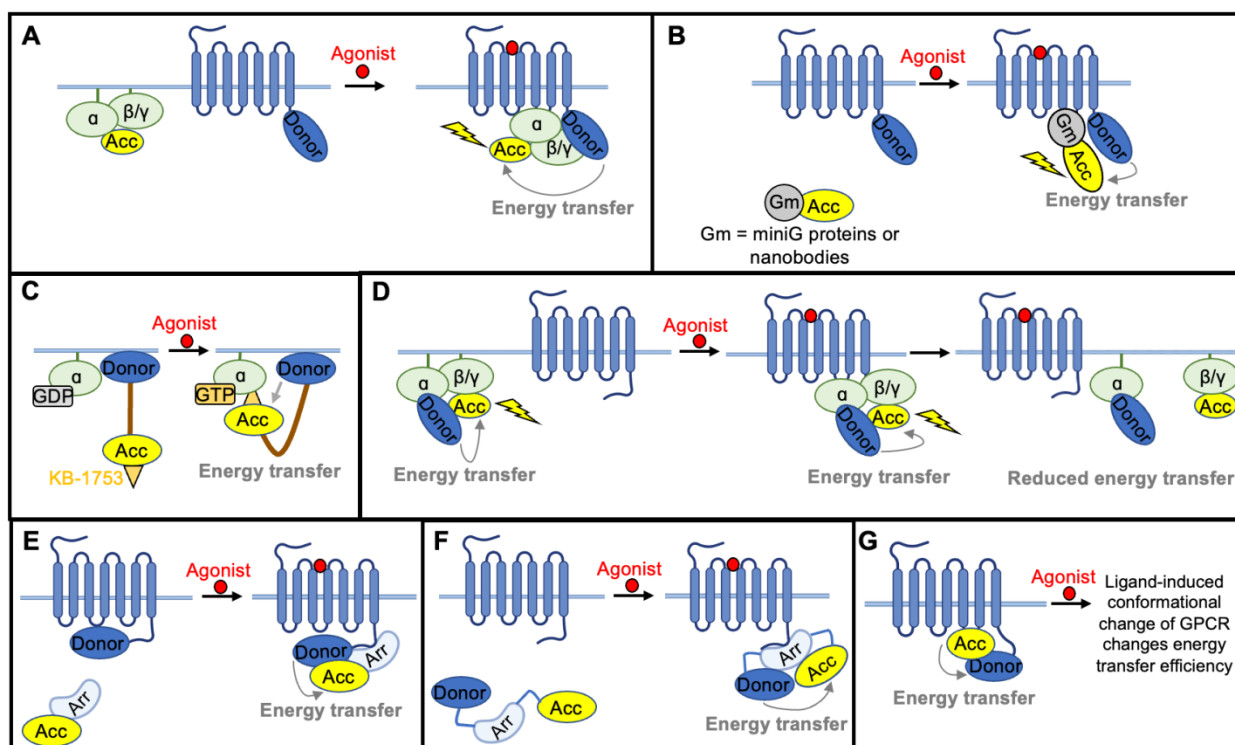


FIGURE 3 FRET and BRET-based sensors. (A) Resonance energy transfer (RET)-based sensor, where one RET pair is attached to the $G\alpha$ -protein and the other is attached to the GPCR. GPCR activation, recruits the G proteins, increasing the energy transfer efficiency. Acc, RET acceptor. (B) RET-based sensor, where one RET pair is attached to the $G\alpha$ -protein mimic, such as a miniG protein or a nanobody, and the other is attached to the GPCR. GPCR activation, recruits the G-protein mimic, increasing the energy transfer efficiency. (C) Schematic of BERKY. BRET pairs are separated by an ER/K α -helix linker, where the acceptor is at the end of the linker and the donor is at the start of the linker, fused to the membrane. KB-1753, a peptide that binds to the active $G\alpha$ -protein, is attached to the acceptor pair. Agonist results in G-protein activation which results in KB-1753 binding and an increase in energy transfer efficiency. (D) RET-based sensor, where one RET pair is attached to the $G\alpha$ -protein and the other is attached to the $G\beta$ - and γ -proteins. GPCR activation causes the distance to increase between the $G\alpha$ and $G\beta\gamma$ proteins, decreasing the energy transfer efficiency. (E) RET-based sensor, where one RET pair is attached to β -arrestin and the other is attached to the GPCR. GPCR activation recruits β -arrestin, increasing the energy transfer efficiency. (F) RET-based sensor, where RET pairs are attached to either terminus of β -arrestin. Agonist activation of the GPCR causes a conformational change in β -arrestin, where there is an increase in energy transfer efficiency. (G) RET-based sensor, where one RET pair is attached to the third intracellular loop of the GPCR and the other is attached to the GPCR's C-terminal tail. Agonist-induced activation causes a conformational change in the GPCR, thereby changing the energy transfer efficiency.

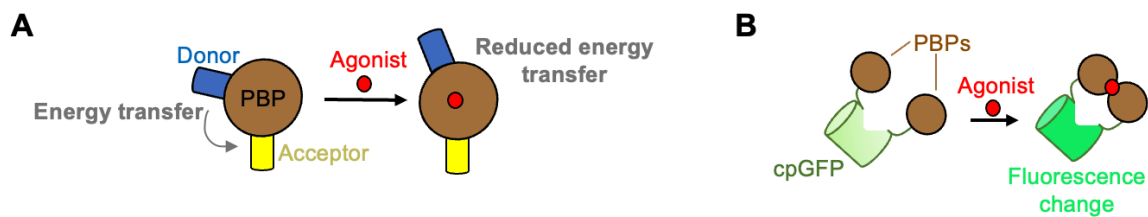


FIGURE 4 PBP-based tools. (A) Resonance energy transfer pairs are attached to periplasmic binding proteins (PBPs). Ligand binding induces a conformational change in the PBP which results in a change in energy transfer efficiency. (B) PBPs are attached to either terminus of cpGFP, where ligand binding to the PBP causes a conformational change in cpGFP, resulting in a fluorescence change.

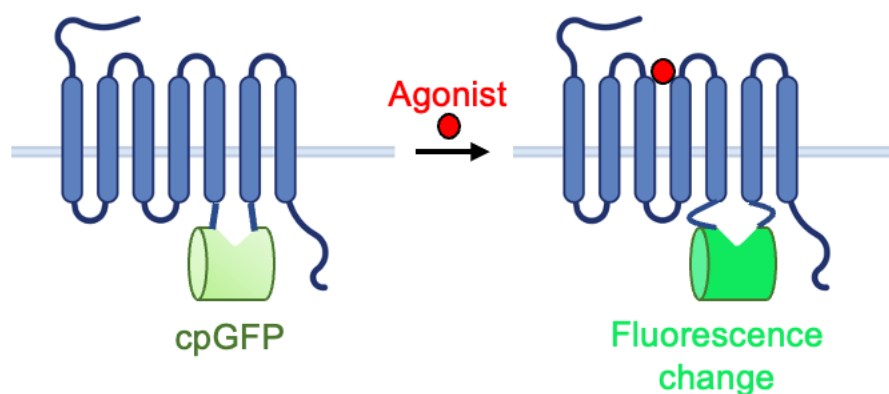


FIGURE 5 cpGFP inserted into the third intracellular loop of a GPCR. Upon agonist-induced GPCR activation, the third intracellular loop of the GPCR undergoes a conformational change, changing the fluorophore environment of cpGFP and resulting in a fluorescence change.

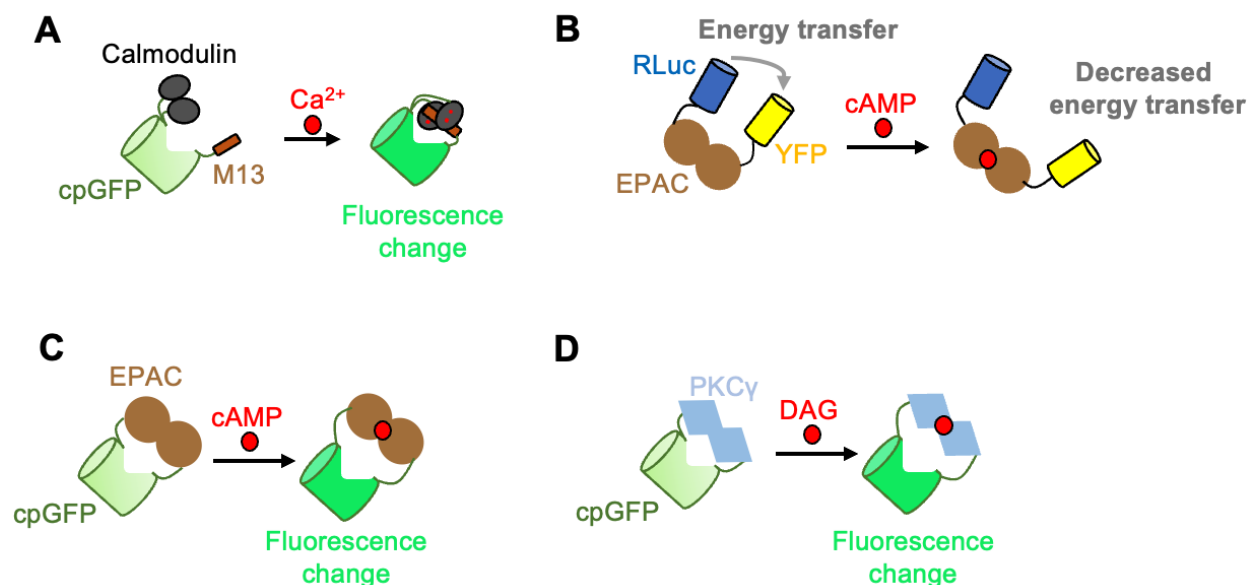


FIGURE 6 Sensors that detect secondary messengers. (A) Calcium-sensing proteins, calmodulin and M13, are attached to either terminus of cpGFP. In the presence of calcium, calmodulin and M13 interact, resulting in a change in the fluorophore environment of cpGFP and a fluorescence increase. (B) Schematic of CAMYEL. BRET pairs RLuc and YFP are tethered to EPAC. cAMP binding causes a

conformational change in EPAC, where the energy transfer efficiency is decreased. (C) EPAC was attached to either terminus of cpGFP. cAMP-induced EPAC conformational change results in a change in cpGFP fluorescence. PKC γ , which only binds to diacylglycerol (DAG), is tethered to either terminus of cpGFP. DAG induces a conformational change in PKC γ , changing the cpGFP fluorescence.

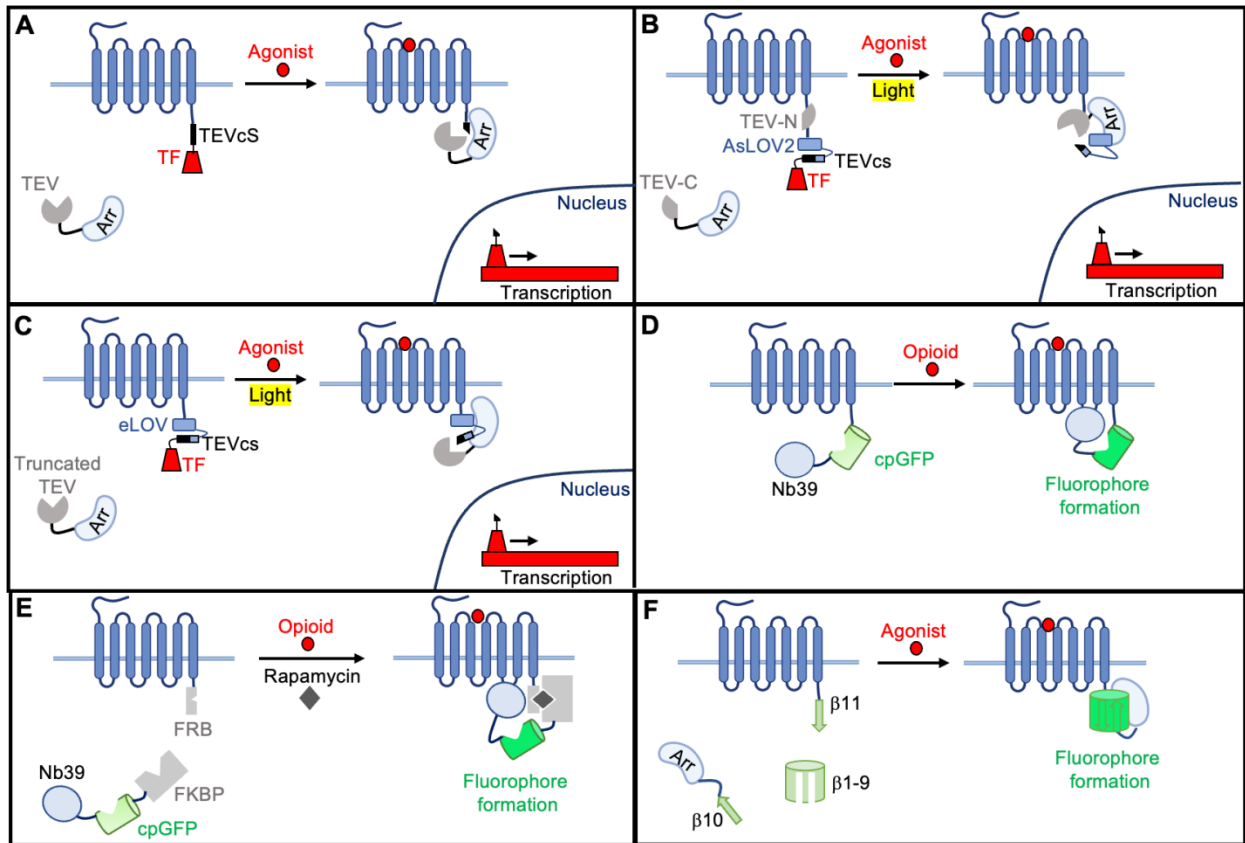


FIGURE 7 Integrators. (A) Schematic of Tango. The C-terminus of the GPCR is fused with a TEVcs and TF. β - arrestin is tethered to a TEV protease. Agonist binding to the GPCR recruits β -arrestin, where the TEV protease can then cut at the TEVcs, releasing the TF so it can translocate to the nucleus and activate a reporter gene. (B) Schematic of iTango2. Same basic mechanism as Tango, except the TEV protease is split into two components (TEV-N and TEV-C) that are fused to β -arrestin and the GPCR. Additionally, the LOV domain cages the TEVcs. Agonist recruits β -arrestin fused split TEV to the GPCR, where the split protease components can reassociate and light uncages TEVcs, allowing the protease to cut and release the TF. (C) Schematic of SPARK. Same basic mechanism as iTango2, except the TEV protease is not split, a truncated protease is used instead. (D) Schematic of SPOTIT. cpGFP and Nb39 are tethered to the C-terminus of the GPCR. Nb39 inhibits cpGFP fluorophore formation. Agonist activates the OR, recruiting Nb39 to the IL3, releasing cpGFP and allowing the fluorophore to form. (E) Schematic of SPOTon. Same basic mechanism of SPOTIT, except the OR is fused to FRB and cpGFP-Nb39 are fused to FKBP. Rapamycin induces heterodimerization of FKBP and FRB, bringing cpGFP-Nb39 to the OR. Opioid activates the OR, recruiting Nb39 to the third intracellular loop and allowing the cpGFP fluorophore to mature. (F) Schematic of Trio. GFP is split into three component: β 1-9, β 10, and β 11. β 10 is attached to β -arrestin, β 11 is attached to the GPCR, and β 1-9 is expressed in the cytosol. Agonist-induced GPCR activation recruits β -arrestin to the Cterminus of the GPCR, allowing the three split components of GFP to re-associate and a fluorescence increase.