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Optogenetic approaches to control Ca2+-modulated physiological processes

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Abstract

As a versatile intracellular second messenger, calcium ion (Ca^{2+}) regulates a plethora of physiological processes. To achieve precise control over Ca^{2+} signals in living cells and organisms, a set of optogenetic tools have recently been crafted by engineering photosensitive domains into intracellular signaling proteins, G-protein coupled receptors (GPCRs), receptor tyrosine kinases (RTKs), and Ca^{2+} channels. We highlight herein the optogenetic engineering strategies, kinetic properties, advantages and limitations of these genetically-encoded Ca^{2+} channel actuators (GECAs) and modulators. In parallel, we present exemplary applications in both excitable and non-excitable cells and tissues. Furthermore, we briefly discuss potential solutions for wireless optogenetics to accelerate the *in vivo* applications of GECAs under physiological conditions, with an emphasis on integrating near-infrared (NIR) light-excitable upconversion nanoparticles (UCNPs) and bioluminescence with optogenetics.

Graphical Abstract

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The authors have no competing interests or conflict of interest to declare.

Keywords

optogenetics; Calcium signaling; protein design and engineering; LOV2; CRY2; Near-infrared light; upconversion nanoparticles; genetically-encoded calcium channel actuators; immune response; neuromodulation; CRISPRa; NFAT; synthetic biology; Ion channel

1. Introduction

 $Ca²⁺$, as a signal for life and death and secondary messenger, controls a myriad of cellular events, including muscle contraction, neurotransmitter release, lymphocyte activation, gene expression and metabolism¹⁻³. In mammalian cells, the intracellular Ca^{2+} concentration is tightly regulated through coordinated actions of a repertoire of Ca^{2+} -signaling components, including G-protein coupled receptors (GPCRs), the receptor tyrosine kinases (RTKs), various Ca^{2+} channels and transporters/exchangers, and intracellular Ca^{2+} sensing and buffering proteins^{1–3}. Upon engagement of membrane receptors by ligands or antigens, phospholipase C gamma (PLC γ) is phosphorylated and activated to catalyze the hydrolysis of phosphatidylinositol 4,5-biphosphate $(PI(4,5)P_2)$, yielding two important secondary messengers, diacylglycerol (DAG) and inositol 1,4,5-triphosphate (InsP₃). InsP₃ further binds to ER-resident InsP₃ receptors to induce Ca^{2+} release from the ER Ca^{2+} store, followed by the activation of store-operated Ca^{2+} entry (SOCE)². SOCE is best exemplified by the Ca^{2+} release-activated Ca^{2+} (CRAC) channel composed of ORAI1 and the stromal interaction molecule 1 (STIM1), and constitutes the primary route of Ca^{2+} entry in nonexcitable tissues^{4–6}. By taking advantage of naturally-existing photoreceptors derived from microbes, plants or mammals, these signalling events can be recapitulated through optogenetic engineering to achieve tailored function. Compared to pharmacological, chemogenetic and genetic approaches, these genetically-encoded Ca^{2+} channel actuators (GECAs) have the advantages of rapid reversibility, superior spatiotemporal resolution, and non-invasiveness⁴. In this review, we present the latest progress in this endeavor, with a focus on the design principles, kinetic features and caveats to be taken into consideration during the application of these tools. When used in combination with genetically-encoded Ca^{2+} indicators, GECAs permit the simultaneous monitoring and interrogation of Ca^{2+} modulated physiological processes both in vitro and in vivo.

2. Engineered photoresponsive GPCRs

As the founding member of the optogenetic field, channelrhodopsin-2 (ChR2) derived from the green alga Chlamydomonas reinhardtii has transformed the neuroscience field by enabling precise control of neural circuits with light both *in vitro* and *in vivo*^{$7-9$}. As a cation permeant channel, ChR2 permits the non-selective entry of Na^+ , H^+ , and Ca^{2+} into the cytosol upon blue-green light stimulation (450–510 nm), thereby depolarizing the plasma membrane (PM) to trigger neuronal firing¹⁰ (Fig. 1a). However, given the over 75-fold higher concentrations of extracellular Na⁺ over Ca²⁺ (136–151 mM *versus* 1.4–1.8 mM), the Ca^{2+} permeability of ChR2 under physiological conditions is very low $(P_{Ca}/P_{Na}: 0.15)^{11}$, thus greatly hampering its application in non-excitable cells and tissues. One ChR2 variant, designated CatCh for Calcium translocating Channelrhodopsin, harbors a point mutation L132C with the Ca²⁺ permeability enhanced by 1.6-fold $(P_{Ca}/P_{NA}: 0.24)^{11, 12}$, but still falls far behind voltage- or store-operated Ca^{2+} channels $(P_{Ca}/P_{Na}: > 1000)^6$. More Ca^{2+} -specific ChR2 variants are yet to be developed.

Instead of site-directed mutagenesis, Airan et al. explored an alternative approach by generating a series of rhodopsin-GPCR chimeras, designated "OptoXRs"13, to photomanipulate intracellular second messengers, including cAMP and Ca^{2+} . In a typical OptoXR, the intracellular loops of rhodopsin were replaced with the intracellular regions of Gq-coupled adrenergic receptors (Fig. 1b). Upon photostimulation at 504 nm, OptoXR evoked cytosolic Ca^{2+} elevation via the activation of the PLC pathway in HEK cells, and was further applied to photo-control behaviors in awake mice¹³. A limitation of ChR2 variants and OptoXRs is that the broad rhodopsin absorption peak (440–590 nm) may potentially conflict with biosensors used to report the functional consequences of optogenetic tools^{14, 15}. In addition, owing to the rapid *trans*-to-*cis* isomerization of the retinal co-factor (tens of ms), sustained Ca^{2+} entry requires constant photostimulation and tends to cause undesired photo-toxicity. The latter concern is partially mitigated by the use of naturally-existing melanopsin (encoded by OPN4), a photopigment protein found in human retinal ganglion cells and the iris of mice and primates¹⁶ (Fig. 1c). Melanopsin can generate both sustained and oscillatory Ca²⁺ signals within seconds ($t_{1/2, ON}$: < 10 s) to activate a downstream Ca^{2+} -responsive master transcription factor, the nuclear factor of activated-T cells $(NFAT)^{17}$. Melanopsin has been introduced into mouse embryonic stem cell-derived embryoid bodies (EBs) to generate pacemaker-like activity in cardiomyocytes within EBs following photostimulation¹⁸. When expressed in non-excitable primary T cells, melanopsin could photo-boost tumor killing in a mouse model of hepatocarcinoma¹⁹. Similar to the design of OptoXRs, a chimeric receptor, Opto-mGluR6, composed of melanopsin and metabotropic glutamate receptor 6 (mGluR6), has been devised and expressed in retina to restore the vision of blind mice with retinal degeneration²⁰. One potential caveat associated with these GPCR-based optogenetic actuators is the paralleled stimulation of the DAG/PKC pathway following PLC activation, which could cause non- $Ca²⁺$ -related functional crosstalk to complicate data interpretation.

3. Photoactivatable RTKs

Similar to GPCR, the engagement of RTKs by their cognate ligands often leads to receptor multimerization to activate the PLC γ pathway for Ca²⁺ mobilization. To faithfully recapitulate this process, optical multimerizers, including Arabidopsis cryptochrome 2 (CRY2), the light-oxygen-voltage domains (LOV) of aureochorme, and Deinococcus phytochrome BphP (DrBphP), have been fused to the catalytic intracellular kinase domain of RTKs to generate Opto-RTKs^{21–24} (Fig. 1d). Most engineering efforts have been devoted to two RTKs, the fibroblast growth factor receptor 1 (FGFR1)^{21, 22} and the tropomyosinrelated kinase $(\text{Trk})^{24}$. Upon light stimulation, the photosensitive moieties undergo dimerization or oligomerization to drive the activation of the intracellular kinase domain of RTKs. Repetitive Ca^{2+} signals can be generated by switching on and off blue light for CRY2 or LOV-based tools^{21–24}, or toggling between near-infrared (780 nm) and far-red light (660 nm) for the case of DrBphP²⁵. Opto-RTKs can thus be applied to control intracellular Ca^{2+} signals using both visible and non-visible light. The use of NIR light further enables multiplexed experiments using fluorescent probes and biosensors in the blue-green range without activation of the engineered receptors. Opto-RTKs, however, are fraught with the lack of Ca^{2+} signaling fidelity because of simultaneous activation of multiple downstream pathways. For example, aside from the Ca^{2+} -associated PLC pathway, photostimulation of Opto-FGFR1 could co-activate proliferation-and survival-related RAS-MAPK, PI3K-AKT and STAT-dependent signaling cascades^{21–24}. Extra cautions, therefore, have to be taken when using Opto-RTKs to interrogate Ca^{2+} -regulated signaling events.

4. Photoswitchable Ca2+ releaser

Optogenetic engineering has also been extended to intracellular Ca^{2+} sensor proteins, such as the four EF-hand-containing calmodulin. A photoactivatable Ca^{2+} releaser (PACR) has been generated via optimized insertion of LOV2 into an engineered calmodulin-M13 fusion protein (CaM-M13) (Fig. 1e). In the dark, PACR interacts with Ca^{2+} with a dissociation constant (K_d) of approximately 16 nM. Upon blue light stimulation, structural changes within LOV2 disrupt the CaM-M13 interaction, which in turn causes Ca^{2+} release from PACR because of a pronounced reduction in the affinity for $Ca^{2+} (K_d = 3.75 \,\mu\text{M})^{26}$, thereby leading to an elevation of $[Ca^{2+}]_{\text{cyto}}$. Even though highly specific for Ca^{2+} , the application of PACR faces three drawbacks: (i) the limited Ca^{2+} releasing capability (<90 nM increase of $[Ca^{2+}]_{cvto}$ and (ii) cell-to-cell variations because of varying levels of PACR expression; and (iii) the potential perturbation to resting cytosolic Ca^{2+} level and complications from other CaM-regulated targets.

5. Optogenetic actuators engineered from the CRAC channel

During SOCE activation, STIM1 dynamically couples to ORAI1 via two major molecular steps (Fig. 1f–g): (i) the initiation of STIM1 activation via Ca^{2+} -depletion induced oligomerization of the luminal domain containing the EF-SAM domain; (ii) inside-out signal propagation toward the STIM1 cytoplasmic domain (STIM1ct) to overcome an intramolecular inhibition primarily mediated by the interaction between the coiled-coil 1 region (CC1) and the STIM-Orai activating domain $(SOAR)^{2, 5, 6, 27}$. Two major optogenetic

engineering approaches, using either CRY2 or LOV2 photosensory domains derived from plants, have been applied to reconstruct these key molecular steps to enable optical control of Ca^{2+} influx through CRAC channels (Fig. 1g).

5.1 CRY2-based OptoSTIM1 and variants

To recapitulate EF-SAM oligomerization during initiation of SOCE, the luminal and transmembrane domains of STIM1 have been replaced by the photolyase-homology region (PHR) of CRY2 to yield OptoSTIM1²⁸, which has an activation kinetics of ≤ 1 min in response to blue light stimulation and a deactivation kinetics of around 4–6 min upon the withdrawal of light (Fig 1g and Table 1). Very recently, two further improved versions, monster-OptoSTIM1 (monSTIM1) and enhanced OptoSTIM1 (eOS1), have been generated to enhance the light sensitivity and dynamic range of Ca^{2+} response by 55-fold and 5~10fold, respectively. The deactivation half-life of the OptoSTIM1 toolset generally ranges from 4 to 7 min^{28, 29}. For eOS1, it takes about 10 min to return to the basal level³⁰. In addition to CRY2-mediated homo-oligomerization, the fusion of optical dimerizers, made of iLID/SspB or CRY2/CIBN, with STIM1ct can likewise photo-trigger Ca^{2+} influx in mammalian cells, with the activation and deactivation half-lives falling in the range of \sim 30 sec and 40–190 sec, respectively³¹. Furthermore, ER-tethered versions were developed to manipulate membrane contact site (MCS) assembly at ER-PM junctions and induce localized Ca²⁺ influx ($t_{1/2, \text{on}}$ = 37.8 \pm 5.3 s and $t_{1/2, \text{off}}$ = 97.5 \pm 16.5 s) in response to blue light stimulation³¹. These tools set the stage for more physiologically-relevant mimicry of native STIM1 puncta formation and Ca^{2+} microdomain generation.

5.2 Opto-CRAC series and other LOV2-based GECAs

To mimic the intramolecular autoinhibition of STIM1 mediated by the juxtamembrane CC1 and SOAR, CC1 was replaced by LOV2 and various STIM1ct fragments were optimized to generate LOV2-STIM1ct chimeras (Opto-CRAC^{31, 32}, BACCS³³, and LOVS1K³⁴). These LOV2-based GECAs enable more rapid and readily reversible control of Ca^{2+} signals, with the activation kinetics ranging from 10 sec to 30 sec and the deactivation half-lives in the range of 30–50 sec (Table 1). In these chimeric constructs, LOV2 cages the fused STIM1ct fragment in a largely inactive conformation presumably through steric hindrance. Upon photostimulation, the photochemical reaction within LOV2 leads to the unfolding of the Cterminal Jα helix to unleash the caged STIM1ct fragments, thereby exposing the SOAR/CAD region to engage and gate ORAI Ca^{2+} channels in the PM (Fig. 1g). Opto- $CRAC³²$ or BACCS variants³³, either anchored to PM or using STIM1ct fragments derived from species other than human and rodent (fruit fly³³, chicken³⁵, and zebrafish³⁶) further diversify the kinetic properties of the LOV2-based GECAs. In addition, these LOV2-based GECAs can be directly fused to ORAI1 channels, in the form of a single or two copies $32, 35$, to enable optogenetic control of Ca^{2+} signaling in cells or tissues with no or little expression of endogenous ORAI proteins.

Collectively, compared to Opto-GPCRs and Opto-RTKs, STIM1-based GECAs directly gate one of the most Ca^{2+} -selective channels, the ORAI protein family, and therefore, retain high Ca^{2+} selectivity over other cations³². Nonetheless, one has to bear in mind the caveat that STIM1, and presumably STIM1-derived GECAs too, might cross-talk with other types of

 $Ca²⁺$ channels (e.g., certain types of transient receptor potential (TRP) channels and voltage gated Ca^{2+} channels) in selected cell types.

6. Genetically-encoded photoswitchable inhibitors of Ca2+ signaling

Complementary to the GECA toolkit, two optogenetic tools capable of suppressing Ca^{2+} entry and/or ER Ca^{2+} release have been generated. To develop a light-inducible OFF-switch for voltage-gated Ca^{2+} (Ca_V) channel, Ma *et al*. developed an optogenetic construct, termed OptoRGK 37 (Fig. 2a), by using a light-sensitive heterodimerization system (iLID/SspB) to control the subcellular localization of a small G protein, Rem_{1-266} . Rem belongs to the RGK family of G proteins and functions as negative regulator of Ca^{2+} entry mediated by Ca_V channels in excitable cells. OptoRGK has been successfully used to perturb Ca^{2+} oscillations in HL-1 cardiomyocytes, and shows promise to rescue cardiac arrhythmia³⁷. In a second case, Hannanta-anan et al. utilized a similar strategy using the CRY2/CIBN optical dimerizer to recruit regulators of G-protein signaling (RGS) toward PM, thereby reducing the lifespan of GTP-bound α-subunits to terminate agonist-evoked calcium oscillations via inhibition of Ga_q -mediated downstream signaling (including $Ca²⁺$ mobilization from internal stores) in response to blue light stimulation³⁸ (Fig. 2b). Complementary to the genetic approach, photopharmacological tools building upon azopyrazoles have also been invented to selectively suppress CRAC channel activity in a light-dependent manner³⁹.

7. Applications of GECAs and wireless optogenetics

Given the strict Ca^{2+} selectivity and high signalling fidelity, STIM1-derived GECAs have been used *ex vivo* to control $Ca^{2+}/NFAT$ -dependent gene expression (Fig. 3a) and cytokine production in cells of the immune system (Fig. 3b). When coupled with the CRISPR activation (CRISPRa) technique⁴⁰, Opto-CRAC can further be repurposed to control the expression of endogenous genes. The calcium-responsive transcriptional reprogramming tool (CaRROT)³⁶, made of NFAT1–460-dCas9-VP64, has been devised to serve this purpose, which allows precise control of transcriptional outputs at user-defined genomic loci, thus expanding the Ca^{2+} -responsive effectors to almost any endogenous genes (Fig. 3c). Moving beyond in vitro applications, recent studies have reported encouraging progress in the use of non-opsin-based optogenetics in living animals.

7.1 Neuromodulation in awake mice

A bottleneck with the existing optogenetic approach is a situation where the depth of tissue penetration of visible light is shallow $(<2-3$ mm)⁴¹, requiring optogenetic protocols to use invasive fiber optic probes or microLED to deliver visible light (in the range of 400–500 nm) into tissues (Fig. 3d). These invasive procedures are technically demanding, tend to elicit inflammatory responses, and remain incompatible with chronic experiments. To overcome these issues, monSTIM1 was generated by using a hyper-sensitive CRY2 variant with one mutation and nine additional amino acids in the C-terminus of $CRY2_{PHR} (CRY2_{E281A}-A9)$ to boost its clustering capability. The new construct displayed 55-fold increase in lightsensitivity compared to the parental OptoSTIM1²⁹ (Fig 1g). Hence, monSTIM1 was utilized to manipulate Ca^{2+} signaling and activate Ca^{2+} -dependent gene expression in deep-brain regions (hippocampal and thalamic regions), and to boost the social learning capability of

awake mice without invasive surgical procedures²⁹ (Fig. 3d). However, upon non-invasive light stimulation, how deep the tissues can be reached to effectively photoactivate monSTIM1 remains elusive. Furthermore, blue light emitting at 450–470 nm may generate local heat and oxidative stress while STIM1 is also sensitive to both heat and $ROS²⁷$. It remains to be rigorously validated if the observed behavior changes are indeed caused by $Ca²⁺$ signals per se rather than undesired side effects of photostimulation. Another attractive strategy to overcome this bottleneck is to exploit optogenetic modules that can be switched on and off by NIR that has a greater depth of tissue penetration (up to centimeters)⁴².

7.2 Immunomodulation in living animals

STIM1-based GECAs have been successfully applied in vivo to control the motility and function of cells of the immune system. eOS1 has been employed to manipulate Ca^{2+} signals in single T cells and photo-tune T cell migration, adhesion and chemokine release in vivo via two-photon-based photoactivation30. Although two-photon optogenetics provide a better spatial specificity and deep penetration in scattering tissue⁴³, potential heat generation during photostimulation and special equipment requirements make it non-ideal for daily use in the laboratory setting, let alone further clinical applications. A wireless nano-optogenetic approach based on NIR-excitable upconversion nanoparticles (UCNPs; Fig. 3e) has been developed to accelerate in vivo application using immune cells. The most significant advantage of UCNPs is their unusual inverse excitation and emission profiles: UCNPs are excited using low power (mW) and deep tissue-penetrating $(1-2 \text{ cm})$ NIR light. This low energy input is efficiently converted to a higher energy output emission at diverse shorter wavelengths, including blue light emission at 470 nm that can activate most optogenetic constructs (Fig. 3e). Since UCNPs can be illuminated from NIR sources located outside the body, they can act as "nano-illuminator relays" to activate optogenetic constructs in a wireless manner. The nano-optogenetic concept has been validated in a dendritic cell-based immunomodulatory therapy against tumor in a mouse mode of melanoma³² (Fig. 3e).

8. Conclusions

In summary, a variety of optogenetic tools, based on both opsin or non-opsin photoreceptors, have been crafted to enable remote control of intracellular Ca^{2+} signaling both *in cellulo* and in living organisms (summarized in Table 1). Most tools are engineered from membrane receptors (GPCRS and RTKs) and signaling proteins (CaM, G proteins and STIM1) that are directly or indirectly involved in Ca^{2+} signaling. Among these tools, GECAs inspired by STIM1 are regarded as the most Ca^{2+} selective and have been most widely applied to modulate the central nervous system and the immune system. The *in vivo* application of these tools will likely be accelerated by developing NIR-responsive optogenetic modules and combining existing tools with nanotechnologies or bioluminescence to enable in vivo optogenetics. The feasibility of bioluminescence-aided optogenetic stimulation (Fig. 3f), at the cost of partially sacrificing the spatial resolution, has been validated in several recent studies^{44–46}. These innovative platforms might represent alternative solutions for future wireless optogenetics.

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Highlights

- **•** Overview of optogenetic tools to control Ca2+-modulated physiological processes
- Engineering strategies for genetically-encoded Ca^{2+} channel actuators (GECAs)
- **•** Guidelines for selecting appropriate GECAs to achieve tailored function
- **•** Insights into potential solutions for wireless optogenetics in vivo

Figure 1. Optogenetic tools for remote activation of intracellular Ca2+ signaling. a. Channelrhodopsin-2 (ChR2) as a cation-permeant channel is non-selective toward $H^+, K^+,$ $Na⁺$, and $Ca²⁺$.

b. OptoXRs as light-sensing rhodopsin-GPCR chimeras. The extracellular and TM regions are derived from a light-sensitive rhodopsin, while intracellular regions from different adrenergic receptors (β_2AR or α_1AR).

c. Blue light-mediated isomerization of the light-absorbing retinal cofactor triggers the conformation change of melanopsin (OPN4) and subsequent activation of Ga_q phospholipase C (PLC) to induce Ca^{2+} influx via TRPC channel.

d. Opto-RTKs as photoactivatable membrane-anchored receptor tyrosine kinases. The intracellular domain of RTKs (FGFR or TrK), is fused to CRY2 or other optical dimerizers, thereby enabling light-inducible clustering to activate downstream signaling, including the PLC pathway and subsequent Ca^{2+} mobilization.

e. LOV2-CaM-M13 hybrid protein as a photoactivatable Ca^{2+} releaser (PACR). Photostimulation results in the release of bound Ca^{2+} into the surrounding environment. f. Domain architecture of STIM1. SP, signal peptide; EF, EF-hand Ca^{2+} -binding motif; SAM, sterile alpha motif; TM, transmembrane domain; CC1, coiled-coil domain 1; SOAR, STIM-Orai activating region; PS, proline-serine rich domain; PB, polybasic domain. g. Genetically-encoded Ca^{2+} channel actuators (GECAs) derived from STIM1. (Left) schematic illustrating functional STIM1-ORAI1 coupling in response to store depletion. Ca^{2+} depletion within the ER lumen induces oligomerization of the luminal EF-SAM domain to initiate STIM1 activation and triggers conformational switch to expose SOAR/CAD and the C terminal PB domain. Next, STIM1 undergoes further oligomerization and migration toward the PM, a process facilitated by the association between SOAR/CAD and ORAI1, as well as the interaction between positively-charged PB and negativelycharged phosphoinositides embedded in PM. SOAR/CAD is responsible for direct engagement and activation of ORAI channels to mediate Ca^{2+} influx. (*right*) Optogenetic engineering to convert store-operated Ca^{2+} entry (SOCE) into lightoperated Ca^{2+} entry (LOCE). (i) A protein oligomerization-based engineering strategy using various optical dimerizers (Opto-CRAC2) and CRY2-STIM1 fusion proteins (OptoSTIM1, monSTIM1, and eOS1). Upon light illumination, CRY2 or optical dimerizers undergoes oligomerization, mimicking Ca^{2+} -depletion induced EF-SAM multimerization in ER lumen, to trigger STIM1 activation and the subsequent opening of ORAI Ca^{2+} channels in the PM. (ii) The design of GECAs based on LOV2 conformational switch (Opto-CRAC1, LOVS1K or BACCS), in which the CC1-SOAR interaction-mediated intramolecular autoinhibition is mimicked by LOV2-SOAR fusion in the dark. Upon photostimulation, LOV2 undergoes conformational changes to expose SOAR/CAD, which engages PMresident ORAI Ca²⁺ channel to induce Ca²⁺ influx.

Figure 2. Genetically-encoded photoswitchable inhibitors of intracellular Ca2+ signaling. a. Cartoon illustration of OptoRGK to photo-control Cay channel activity in cardiomyocytes. An optical dimerizer pair made of LOV2-SsrA (iLID) and SspB is utilized to induce the translocation of Rem1 core domain (a member of the RGK family of GTPbinding proteins) from the cytosol to PM, where it engages the auxiliary beta subunits of CaV channel to suppress its activity upon photostimulation.

b. Design of optogenetic regulators of G-protein signaling (Opto-RGS2). In this design, RGS2 is split into two components, including the catalytic box (R2-box) which is critical for inhibition of Ga_q signaling and the N-terminal amphipathic helix (R2-helix). The two split parts are fused with either CRY2_{PHR} or its binding partner CIBN. In the dark, R2-box is distributed in the cytoplasm. Upon blue light illumination, the forced dimerization between CIBN-R2-helix and CRY2-R2-box triggers reconstitution of the active protein and

subsequently inhibit Ga_q signaling and terminates Ca^{2+} release originated from GPCR signaling.

Figure 3. Applications of GECAs and potential solutions for wireless optogenetics.

a. GECAs (e.g., Opto-CRAC) utilized to control Ca^{2+} -dependent gene expression using synthetic response elements (RE) derived from various Ca^{2+} -responsive transcription factors. NFAT, nuclear factor of activated-T cells; CREB: cAMP response element-binding domain; SRE, serum response element.

b. GECAs used to control $Ca^{2+}/NFAT$ -driven cytokine production in T cells in the presence of co-stimulatory signals to activate activator protein 1 (AP-1).

c. Coupling GECAs with CRISPRa to enable light-tunable transcriptional programming. Upon blue light illumination, Opto-CRAC1 induces cytosolic Ca^{2+} influx, which in turn activates CaM/calcineurin to dephosphorylate the N-terminal non-DNA binding fragment of NFAT (NFAT_{1–460}) and eventually drive the nuclear entry of the fusion protein (NFAT_{1–460}dCas9-VP64) – a calcium-responsive transcriptional reprogramming tool (designated as CaRROT). In the presence of small guide RNAs (sgRNAs), CaRROT can be precisely

targeted to the promoter regions of any desired genomic loci to photo-tune the expression of endogenous genes.

d. Optogenetic neuromodulation with GECAs (e.g., OptoSTIM1 or monSTIM1). Blue light can be delivered into a specific brain region via implantation of optical fiber (i) or μLED (ii). Photoactivatable Ca²⁺ influx can lead to induction of Ca²⁺-responsive gene expression (e.g., c-Fos) in both neurons and astrocytes in deep-brain regions such as hippocampal and thalamic regions (iii), thereby modulating the behavior (e.g., social fear learning) in awake mice (iv).

e. A nano-optogenetic platform for wireless optogenetic immunomodulation in a mouse model of melanoma. (i) schematic illustration of the core/shell structure of lanthanide doped upconversion nanoparticles (UCNPs), which act as nanoilluminator to emit visible light upon near-infrared irradiation (NIR); (ii-iii) NIR light-controllable activation of Ca^{2+} influx in dendritic cells (DCs) expressing Opto-CRAC1 could boost antigen presentation and DC maturation to more efficiently prime T cells in the draining lymph nodes, thereby facilitating dendritic cell-based immunomodulatory therapy against melanoma.

f. Bioluminescence-aided optogenetic stimulation (BLOGS). (i) Cartoon illustrating the photoactivation of a chimeric luminopsin (LMO) receptor comprising a microbial opsin and Gaussia luciferase (GLuc) or NanoLuc luciferase (NLuc) in the presence of their cognate substrates coelenterazine (CTZ) or furimazine (FZ). (ii) Photochemical reactions catalyzed by GLuc or NLuc. (iii) Schematic depiction of wireless optogenetics using BLOGS in mice administered with luciferase substrates.

Table 1.

Summary of optogenetic tools for noninvasive and precise control Ca^{2+} signals in mammals. Summary of optogenetic tools for noninvasive and precise control Ca^{2+} signals in mammals.

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* cytosol-plasma membrane translocation cytosol-plasma membrane translocation

endothelial cells; iLID, improved light-induced dimer (LOV2-SsrA); LOV2. light-oxygen-voltage sensing) domain; LOVS1K, LOV2 fused STIM1 fragments (233-450); monSITM1, monster-OptoSTIM1; endothelial cells; iLID, improved light-induced dimer (LOV2-SsrA); LOV2, light-oxygen-voltage sensing) domain; LOVS1K, LOV2 fused STIM1 fragments (233–450); monSTIM1, monster-OptoSTIM1; terminal domain of CIB1; CRY2, cryptochrome2; eOS1, enhanced OptoSTIM1; FGFR1, fibroblast growth factor receptor 1; GECA, genetically encoded Ca²⁺ actuator; HUVECs, human umbilical vein terminal domain of CIB1; CRY2, cryptochrome2; eOS1, enhanced OptoSTIM1; FGFR1, fibroblast growth factor receptor 1; GECA, genetically encoded Ca²⁺ actuator; HUVECs, human umbilical vein OptoXRs, optogenetic control of intracellular signal transduction; S1, somatosensory cortex; S2, Drosophila S2 cells; SP, Signal peptide of STIM1; NIR, near-infrared; TM, transmembrane domain; Trk, PACR, photoactivatable Ca²⁺ releaser; Opto-a1AR, Optogenetic G_q-coupled a1-adrenergic receptor; Opto-CRAC, Optogenetic Ca²⁺ release-activated Ca²⁺ channel; OptoRGK, Optogenetic Ras-like GTPase Rad/Rem/Gem/Kir, Opto-RGS2, Optogenetic regulators of G-protein signaling; Opto-RTKs, Optogenetic receptor tyrosine kinases; OptoSTIM1, Optogenetic stromal interaction molecule 1;
OptoXRs, optogenetic control of int PACR, photoactivatable Ca²⁺ releaser; Opto-α1AR, Optogenetic Gq-coupled α1-adrenergic receptor; Opto-CRAC, Optogenetic Ca²⁺ release-activated Ca²⁺ channel; OptoRGK, Optogenetic Ras-like Abbreviations: BACCS, blue light-activated Ca²⁺ channel switch; BMDCs, bone marrow-derived DCs; CaM-M13, calmodulin fused with the M13 peptide; ChR2, Channelrhodopsin-2; CIBN, the N-Abbreviations: BACCS, blue light-activated Ca²⁺ channel switch; BMDCs, bone marrow-derived DCs; CaM-M13, calmodulin fused with the M13 peptide; ChR2, Channelrhodopsin-2; CIBN, the N-GTPase Rad/Rem/Gem/Kir; Opto-RGS2, Optogenetic regulators of G-protein signaling; Opto-RTKs, Optogenetic receptor tyrosine kinases; OptoSTIM1, Optogenetic stromal interaction molecule 1; tropomyosin-related kinase; tropomyosin-related kinase;