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Development of green fluorescent protein-based cAMP indicators for covering a wide range of cAMP concentrations†

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There is a wide range in the concentration of intracellular cyclic adenosine 3',5'-monophosphate (cAMP), which mediates specific effects as a second messenger in pathways affecting many physiological processes. Here, we developed green fluorescent cAMP indicators, named Green Falcan (Green fluorescent protein-based indicator visualizing cAMP dynamics) with various EC₅₀ values (0.3, 1, 3, 10 μM) for covering the wide range of intracellular cAMP concentrations. The fluorescence intensity of Green Falcans increased in a cAMP dose-dependent manner, with a dynamic range of over 3-fold. Green Falcans showed a high specificity for cAMP over its structural analogues. When we expressed Green Falcans in HeLa cells, these indicators were applicable for visualization of cAMP dynamics in the low concentration range compared to the previously developed cAMP indicators, and visualized distinct kinetics of cAMP in various pathways with high spatiotemporal resolution in living cells. Furthermore, we demonstrated that Green Falcans are applicable to dual-color imaging with R-GECO, a red fluorescent Ca²⁺ indicator, in the cytoplasm and the nucleus. This study shows that Green Falcans open up a new avenue for understanding hierarchal and cooperative interactions with other molecules in various cAMP signaling pathways by multi-color imaging.

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1. Introduction

Cyclic adenosine 3',5'-monophosphate (cAMP) plays an important role as a second messenger with a wide range of concentrations from sub μM to μM in living cells.^{1–3} Many hormones or neurotransmitters stimulate G protein-coupled receptors in the cell membrane, leading to the activation of adenylyl cyclase which catalyzes the conversion of ATP to cAMP.⁴ In living cells, cAMP binds to protein kinase A (PKA),⁵ exchange proteins directly activated by cAMP (Epac),⁶ or cyclic nucleotide-gated (CNG) ion channels,⁷ and causes many cellular functions such as hormone secretion,⁸ cell motility,⁹ and memory formation,¹⁰ etc.

To date, many single-fluorescent protein (FP) based indicators have been developed as tools for detecting various

intracellular signaling molecules (Ca²⁺, cAMP, cGMP, etc.).^{11–17} Since single-FP based indicators require single wavelength emission for detection, it is suitable for multicolor imaging to simultaneously visualize the hierarchal and cooperative relationship with other molecules in the same or distinct intracellular compartments, with high spatiotemporal resolution. They show increased and decreased FI in the presence of analytes, called turn-on type and turn-off types, respectively. A few turn-off type green fluorescent cAMP indicators have been produced,^{14,15,18} but distinguishing photobleach or the degradation of the indicator from analyte decrease is problematic. Several turn-on green fluorescent cAMP indicators had been developed, but their utility was limited by low dynamic range.^{18–20} Very recently, new cAMP indicators with higher dynamic range were developed and this limitation is being resolved,²¹ but it is not sufficient to monitor a wide range of cAMP concentrations in living cells (Table S1†).

Here, we utilized our previously described method of semi-rational molecular design and molecular evolution²² in developing indicators such as Green cGull,¹⁷ MaLionG,²³ and Green Glifon,²⁴ to generate powerful turn-on type green fluorescent cAMP indicators with expanded dynamic range and various half maximal effective concentration (EC₅₀) employing PKA as a binding domain, named them Green Falcan 0.3, 1, 3, 10. We demonstrated that this useful method is not limited to cAMP

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indicators, but hopefully leads to the development of new indicators for a myriad of molecules.

2. Materials and methods

2.1. Chemicals

cAMP and cGMP were purchased from Sigma-Aldrich (St. Louis, MO, USA), forskolin from Tokyo Chemical Industry (Tokyo, Japan), 3-isobutyl-1-methylxanthine, 8-Br-cAMP and adenosine from Merck Millipore (Darmstadt, Germany), ATP, ADP and AMP from Oriental Yeast Co., Ltd. (Tokyo, Japan).

2.2. Plasmid construction

The DNA fragment of GFP, restructured from circularly permuted (cp) GFP in G-GECO,¹³ a green fluorescent genetically-encoded Ca²⁺ indicator, was modified by PCR to contain KpnI and BglII restriction enzyme sites between N144 and S146 and cloned into the pRSET-A vector (Thermo Fisher Scientific, Massachusetts, USA). The cDNA for the mouse cAMP-dependent protein kinase (PKA) type I-beta regulatory subunit²⁵ (NP_001240819.1) was obtained from adult mouse brain total RNA by RT-PCR. The region of the cDNA encoding the cAMP binding domain (amino acids 92–248) derived from site A, was ligated at the KpnI/BglII site of the GFP DNA. To improve the fluorescence intensity (FI), the N- and C-terminus linker lengths were optimized by inserting various lengths (N-terminus 0 to 9 amino acids; C-terminus 0 to 10 amino acids) of leucine zipper sequences²⁶ between the GFP and the cAMP binding domain sequences. The plasmids encoding mutant indicators with different linker lengths at the N- and C-terminus of the cAMP binding domain were transformed into *Escherichia coli* JM109 (DE3) (Promega, Madison, USA), and these lysates were subjected to examine the FI change. The plasmid DNA of the mutant showing the maximum response was selected as a template for subsequent random mutagenesis. To further improve the FI change by introducing random mutations into the amino acid sequences around the linkers, PCR was performed using two degenerate primers containing NNK and MNN mixtures of bases. Plasmids carrying site-directed random mutations around linkers were transformed into JM109 (DE3), and around 50 colonies (empirically ~15 kinds of amino acid coverage) were picked. The plasmid of the mutant with the maximum response was selected as the next template for the molecular evolution cycle of random mutagenesis, and this cycle continued until the response was more than 3-fold. Then, to adjust the half-maximal effective concentration (EC₅₀) of Green Falcans by changing the affinity to cAMP, PCR was performed using two primers containing NNK and MNN designed to the region around R211 of PKA, which is important for cAMP binding.²⁷ The mutants with various EC₅₀ (1, 3, 10 μM) were selected by the dose-response curve using a four-parameter logistic curve fitting. For live-cell imaging in mammalian cells, Green Falcans DNA sequences were cloned into the pcDNA3.1 (–) vector (Thermo Fisher Scientific, Massachusetts, USA) (Fig. S2A–D†). To improve the solubility of the expressed proteins, the super acidic region of mouse amyloid β precursor

protein (amino acids 190–286) was fused to the N-terminus of the indicators. For subcellular localization, the nuclear localization signal (nls; PKKKRKV) fused to the N-terminus of the indicators was derived from SV40. The cAMP indicator, cAMP_r plasmid was obtained from Addgene (#99143, Massachusetts, USA), and was cloned into the pcDNA3.1 (+) vector for better expression.

2.3. Protein production and purification

Green Falcans DNA in the pRSET-A vector were transformed into *Escherichia coli* JM109 (DE3) cells, cultured in 400 mL LB medium with 50 μg mL^{–1} ampicillin at 20 °C for 4 days, and harvested by centrifugation. The harvested cells were suspended in phosphate-buffered saline (PBS; pH 7.4) with 0.5% (v/v) Triton-X and 40 μg mL^{–1} lysozyme, and lysed by French press (Constant Systems Ltd, Daventry, UK). After centrifugation of the lysate, the supernatant containing the Green Falcans protein was purified by TALON® Metal Affinity Resin (TaKaRa-Bio, Shiga, Japan). After washing three times with 10 mM imidazole, the proteins were eluted from the resin with 300 mM imidazole. To remove the imidazole, the eluate was applied to PD-Miditrap G-25 (GE Healthcare, Chicago, USA) in 150 mM KCl, 50 mM HEPES-KOH (pH 7.4), 0.5% (v/v) Triton-X. The purified protein was stored at –80 °C.

2.4. SDS-PAGE

The purified Green Falcans proteins (1 μg) were mixed with an equal volume of 2 × SDS loading buffer, and heated at 95 °C for 5 min. These were loaded in 10% acrylamide gel. Precision Plus Protein Unstained Standards (Bio-Rad, California, USA) was also loaded as a protein marker. After electrophoresis, the gel was dyed by CBB staining. For taking a picture, luminographi (ATTO, Tokyo, Japan) was used.

2.5. In vitro spectroscopy

The excitation and emission spectra of Green Falcans were measured using a spectrophotometer (F2700, Hitachi, Tokyo, Japan). To examine the dose-response relationship, various concentrations of cAMP and cGMP (from 0.001 to 300 μM) were applied to 0.3 μM purified Green Falcans protein. Relative FI was calculated by dividing by the FI of the peak in the absence of cAMP. To obtain the EC₅₀ value and dynamic range of the Green Falcans for cAMP and cGMP, a four-parameter logistic curve fitting (eqn (1)) was performed using QtiPlot (IONDEV SRL, Bucharest, Romania).

$$y = d + \frac{a - d}{1 + \left(\frac{x}{c}\right)^b} \quad (1)$$

Here, a is the minimum value, b is the Hill coefficient, c is EC₅₀ and d is the maximum value. For cAMP, the minimum FI in fitting was normalized to 1, and then the maximum FI was used as the dynamic range. For cGMP, the minimum FI in the fitting was normalized to 1, and the maximum FI was normalized to the maximum FI for cAMP in the fitting. The working range for cAMP was calculated as a linear range using the second



derivative.²⁸ To examine the dose-response relationship for cAMP in the presence of cGMP, cGMP was applied at 10 times the concentration of each EC₅₀ value. To examine specificity, cAMP analogues were applied at 10 times the concentration of each EC₅₀ value to 0.3 μM purified Green Falcan proteins. The absorption spectra of 50 μM Green Falcan proteins were measured using a spectrophotometer (V-730BIO, JASCO Corporation, Tokyo, Japan) with or without 100 μM cAMP. To examine pH sensitivity, 0.3 μM purified Green Falcan proteins were applied in 150 mM KCl, 100 mM HEPES-KOH (pH 6–8.5).

2.6. Cell culture and transfection

HeLa cells were cultured in Dulbecco's modified eagle medium (high glucose) with L-glutamine, sodium pyruvate, penicillin and streptomycin (Wako, Osaka, Japan), 10% (v/v) heat-inactivated fetal bovine serum (GE Healthcare, Chicago, USA). These cells were subcultured in 35 mm glass-bottomed dishes at 37 °C with 5% CO₂ for live-cell imaging. Plasmids of each Green Falcan were transfected to these cells with Lipofectamine 3000 Transfection Reagent (Thermo Fisher Scientific, Massachusetts, USA) according to the manufacturer's instruction. Then, HeLa cells were cultured at 37 °C overnight. After changing the medium, HeLa cells were cultured at 30 °C to 32 °C for at least 30 h to allow maturation of indicator chromophores.

2.7. Live-cell imaging

Live-cell imaging was performed using a fluorescence inverted microscope (IX70, Olympus, Tokyo, Japan) equipped with an oil immersion ×40 objective lens (UApO/N 340, ×40, NA = 1.35, Olympus), a cooled CCD camera (Cool SNAP HQ2, Photometrics), a mercury lamp, an excitation filter of 460–480 nm, an emission filter of 495–540 nm and a dichroic mirror of 485 nm (U-MGFPHQ, Olympus). For dual-color imaging, the DA/FI/TR-3X3M-C Sedat filter set (OPTO-LINE, Inc., Saitama, Japan) is set by an excitation filter of 546–566 nm, an emission filter of 580.5–653.5 nm and triple band dichroic mirror of 403, 497 and 574 nm, was changed by using an HF110 high-speed filter wheel (Prior Scientific, Cambridge, UK). Fluorescence images were acquired every 5 s using Metafluor software (Molecular Devices, Sanjose, USA). The FI was measured by manually surrounding each cell with a region of interest (ROI) using ImageJ software (<https://imagej.net/software/fiji/downloads>).²⁹ After subtracting the FI of a cell-free area, relative FI was calculated by dividing by the mean of the FI captured over 1 or 2 min before administering reagents. For statistical analysis, a Student's *t*-test was performed using R (<https://www.r-project.org/>)³⁰ for Fig. 3F, S6B and S7C, F, † and one-way ANOVA was performed using EZR (<https://www.jichi.ac.jp/saitama-sct/SaitamaHP.files/statmed.html>)³¹ for Fig. 3C.

3. Results and discussion

3.1. Generation of Green Falcans

To generate turn-on type green fluorescent cAMP indicators with expanded dynamic range, we first constructed a prototype

indicator by inserting the cAMP binding domain (amino acids 92–248) derived from site A of mouse cAMP-dependent protein kinase (PKA) type I-beta regulatory subunit²⁷ (NP_001240819.1) into the vicinity of the chromophore of green fluorescent protein (GFP) (Fig. S1A†). When this prototype indicator was expressed in bacteria, the FI measured in the lysates increased slightly in the presence of cAMP. Since this response was insufficient for easy detection of minor changes in cAMP levels, we planned to improve the FI of prototype indicator by protein engineering. To begin, we selected the linker length between GFP and PKA (N-terminus: 4 amino acids, C-terminus: 3 amino acids) that displayed the largest FI change (1.3-fold) (Fig. S1B and C†). For further improvement of the dynamic range, we optimized the amino acid sequences around the N- and C-terminus linkers by site-directed random mutagenesis using PCR. After 40 cycles of random mutation, we obtained a green fluorescent cAMP indicator with more than 3-fold FI change in the presence of cAMP (Fig. S1D†). When the dose-response relationship to cAMP was examined, we found that the EC₅₀, apparent *K*_d, of this indicator was 0.3 μM (Fig. S4E†). Generally, cAMP is known to function in a wide range of concentrations from sub μM to μM range in living cells.¹ Therefore, to cover this range, we introduced random mutations around R211 of PKA, which is important for cAMP binding.²⁷ Consequently, by the selection based on the response to several concentrations of cAMP, we succeeded in obtaining additional three green fluorescent cAMP indicators with various EC₅₀ (1, 3, and 10 μM) (Fig. 1A, S4F–H†). The indicators, Green Falcan 0.3, 1, 3, and 10 were named after their EC₅₀ value and abbreviation of green fluorescent protein-based indicator visualizing cAMP dynamics (Fig. 1B). Additionally, utilizing the same molecular design such as insertion of a ligand-binding domain into the vicinity of an FP chromophore and optimization of linker region to expand FI change, we used to develop our previous indicators for cAMP,^{14–16} cGMP,¹⁷ ATP,²³ glucose,²⁴ pyruvate³³ and lactate³³

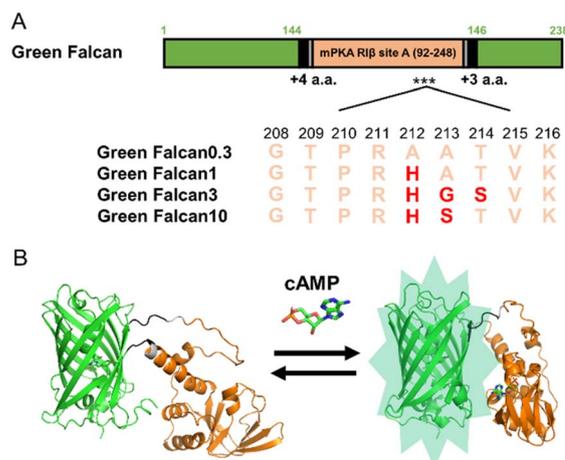


Fig. 1 Schematic design of Green Falcan proteins. (A) Diagram for Green Falcan proteins. Red letters indicate mutated amino acids. (B) Schematic 3D structures of Green Falcan unbound (left) and bound (right) to cAMP. The schematic 3D structures were created with AlphaFold2.³²



demonstrates its versatility for generating genetically-encoded indicators based on fluorescent protein.

3.2. Property of Green Falcons

To investigate the properties of the Green Falcons, we purified (Fig. S3†) and examined excitation and emission spectra for these indicators. All Green Falcons had two excitation peaks at around 400 nm and 498 nm with an emission peak at around 511 nm (Fig. 2A–D). The peaks in excitation at around 498 nm, and emission at around 511 nm, which are similar to the spectral properties of GFP,³⁴ were increased more than 3-fold in the presence of cAMP. In the presence of cAMP, the absorption spectra of all Green Falcons showed a little decreased and increased peaks at around 400 nm and 500 nm, respectively (Fig. S4A–D†). It has been demonstrated that the GFP absorption peak observed at around 400 nm corresponds to the protonated state of the chromophore and that observed at 500 nm, to the deprotonated state.³⁴ Thus, we think that a change in the population of the protonated/deprotonated state of chromophore contributes to the fluorescence intensity (FI) change. Interestingly, unlike other GFP-based indicators, Green Falcons have an excitation peak at around 400 nm, which increased in the presence of cAMP. Since the excitation peak derived from the protonated form of chromophore creates the fluorescence at around 511 nm by excited-state proton transfer,^{35,36} it seems that the proton transfer was also accelerated by the application of cAMP to induce FI change. However, the

measurement with 400 nm excitation is not practical because the dynamic range with 400 nm excitation is lower than that with 480 nm, and the excitation at short wavelength causes cell damage.

When the dose-response relationship of all Green Falcons was examined, we found that the EC₅₀ value of Green Falcon 0.3, 1, 3, 10 for cAMP was 0.30, 1.4, 2.6 and 11 μM, and the dynamic range in the presence of cAMP was 3.9, 5.6, 4.7, 6.5-fold, respectively. While we found that the EC₅₀ values of all Green Falcons for cGMP were higher than that for cAMP (Fig. 2E, S4E–H,† Table 1). However, since the EC₅₀ values of Green Falcons for cAMP are about 10–100 times lower than that for cGMP, the detection of cAMP is still affected in the presence of cGMP. Therefore, the dose-response relationship for cAMP in the presence of cGMP (10 times EC₅₀ for cAMP) was also examined. The responses of all Green Falcons for cAMP were smaller, while the EC₅₀ values of all Green Falcons were similar to those without cGMP, suggesting that Green Falcons function efficiently in the presence of cGMP (Fig. S4I–L†). To examine the specificity of all Green Falcons for cAMP, we applied cAMP structural analogues (ATP, ADP, AMP, adenosine, cGMP) at 10 times the concentrations of the EC₅₀ for each indicator. This demonstrated that all Green Falcons display high specificity for cAMP (Fig. 2F). The previously reported single-FP based indicators have shown pH sensitivity.^{14–17,23,24,33} Similarly, the FI of Green Falcons was increased depending on pH elevation (Fig. S4M–P†). These results suggest that by choosing the appropriate indicator, according to its EC₅₀, for the expected intracellular cAMP range, these indicators can be used to specifically visualize a wide range of cAMP levels in living cells during a variety of physiological events. Although we have employed a cAMP binding domain from mouse Epac1 for developing cAMP indicators such as Flamindo,¹⁴ Flamindo2¹⁵ and Pink Flamindo,¹⁶ that from PKA was done in this study. Since introducing a mutation in the binding domain usually decreases the affinity for ligands, we recommend exploiting a binding domain with a higher affinity to develop indicators covering a wide range of ligand concentrations.

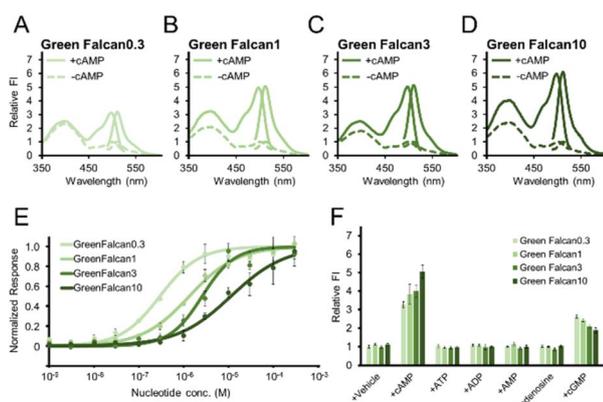


Fig. 2 Spectral and functional properties of Green Falcons (A–D) excitation and emission spectra of purified Green Falcon 0.3, 1, 3, and 10 in the presence (solid line) and absence (dashed line) of 100 μM cAMP. The relative fluorescence intensity (FI) was calculated by dividing by the FI of the peak in the absence of cAMP. (E) Dose-response curves of all Green Falcons. The FI was measured with excitation at 480 nm. The normalized response was calculated by dividing by the FI of the peak in the absence of cAMP, and the minimum and maximum FI in fitting was normalized to 0 and 1, respectively. The data represent the means ± standard deviation ($n = 3$). (F) Specificity of purified Green Falcons for cAMP structural analogues. The concentration of these analogues were 3, 10, 30 and 100 μM for Green Falcon 0.3, 1, 3 and 10, respectively. The FI was measured with excitation at 480 nm. Each relative FI was calculated by dividing by the FI of the peak in the absence of cAMP analogues. The data represent the means ± standard deviation ($n = 3$).

3.3. Applicability of Green Falcons to live-cell imaging

To investigate the applicability of Green Falcons to live-cell imaging, we expressed these indicators in HeLa cells, and monitored with several reagents increasing intracellular cAMP levels under a fluorescence microscope. Green Falcons were well expressed in HeLa cells, but had lower basal fluorescence intensity than HeLa cells expressing cAMP_r, a previously developed cAMP indicator (Fig. S5A and B†). After the administration of 100 μM forskolin, an adenylate cyclase activator, the fluorescence intensity (FI) of these cells expressing each Green Falcon was significantly increased (Fig. 3A–C, S6A–C†), suggesting that all Green Falcons are able to be used in visualizing cAMP dynamics in living cells. Moreover, when the administration of 5 μM forskolin to HeLa cells expressing Green Falcon 0.3, 10, cAMP_r or Pink Flamindo, previously developed cAMP indicators, the FI of Green Falcon 0.3 was significantly increased, but not of Green Falcon 10 and cAMP_r (Fig. S7A–C†).



Table 1 EC₅₀ of each Green Falcan

Indicator name	EC ₅₀ (cAMP) (μM)	EC ₅₀ (cGMP) (μM)	Working range (cAMP) (μM)
Green Falcan 0.3	0.30	2.8	8.1×10^{-2} –1.4
Green Falcan 1	1.4	1.0×10^2	0.30–9.3
Green Falcan 3	2.6	3.1×10^2	0.75–7.9
Green Falcan 10	11	8.7×10^2	4.9 – 2.6×10^{-2}

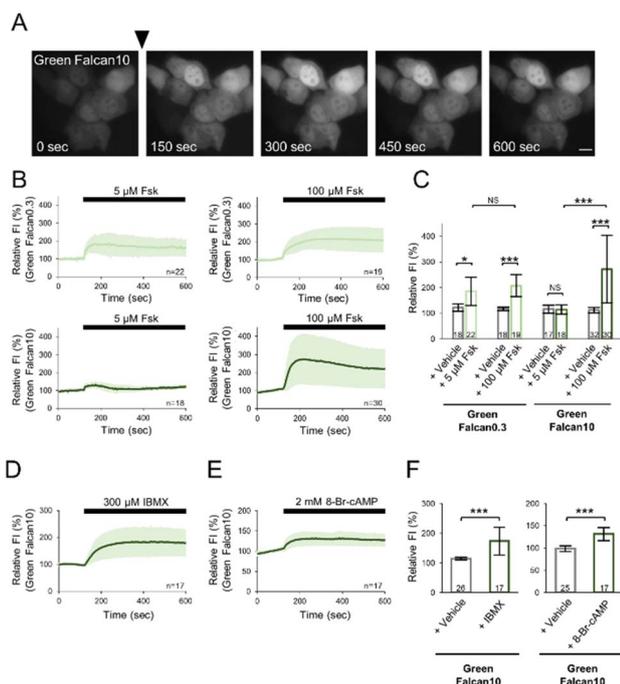


Fig. 3 Live-cell imaging using Green Falcans (A) sequential images of Green Falcan 10 expressing HeLa cells. Scale bar represents 10 μm. (B) Time courses of fluorescence intensity (FI) in HeLa cells expressing Green Falcan 0.3 and 10 in response to 5 or 100 μM forskolin. Relative FI was calculated by dividing by the mean of the FI during the 2 min prior to administration. The data represent the means ± standard deviation. The numbers in the lower right corner of the graphs represent the number of cells analyzed from three independent experiments. (C) Comparison of relative FI at 2 min after the administration of DMSO (vehicle) or forskolin to HeLa cells expressing Green Falcan 0.3 and 10. The data represent the means ± standard deviation. The numbers in bar graphs represent the number of cells analyzed from three independent experiments. **p* < 0.05, ****p* < 0.001. (D and E) Time course of FI in HeLa cells expressing Green Falcan 10 in response to 300 μM IBMX (D) or 2 mM 8-Br-cAMP (E). The data represent the means ± standard deviation. The numbers in the lower right corner of the graphs represent the number of cells analyzed from three independent experiments. (F) Comparison of relative FI at 2 min after the administration of DMSO (vehicle) or 300 μM IBMX, and after that of dH₂O (vehicle) or 2 mM 8-Br-cAMP to HeLa cells expressing Green Falcan 10. The data represent the means ± standard deviation. The numbers in bar graphs represent the number of cells analyzed from three independent experiments. ****p* < 0.001.

The FI of Pink Flamindo was also increased, but the change was smaller than that of Green Falcan 0.3 (Fig. S7D–F†). In addition, when the dose-response relationship for forskolin in HeLa cells

expressing Green Falcans, the FI of all Green Falcans by the administration of forskolin was increased in a dose dependent manner and Green Falcan 0.3 is most sensitive to forskolin (Fig. S6E–H†). These results suggest that a wide range of cAMP concentrations in a living cell is able to be monitored by choosing Green Falcan with a suitable working range. We further examined Green Falcan 10 carrying a similar EC₅₀ with previous cAMP indicators in live-cell imaging. After the administration of 300 μM 3-isobutyl 1-methylxanthine (IBMX, phosphodiesterase inhibitor) or 2 mM 8-Br-cAMP (membrane-permeable cAMP analogue) to HeLa cells expressing Green Falcan 10, the FI of these cells was significantly increased (Fig. 3D–F, S6D†). Interestingly, the kinetics and amplitude of FI change after the administration of forskolin, IBMX and 8-Br-cAMP differed from each other, reflecting the distinct kinetics of various mechanisms for achieving cAMP increase such as production, inhibition of degradation, and membrane permeation. Therefore, these results suggest that the developed indicators are adaptable for detecting a spatial and temporal change of cAMP dynamics in various physiological phenomena.

3.4. Dual-color imaging with Green Falcan and R-GECO

Many single-FP based indicators have been developed for visualizing with other signaling molecules. Since single-FP based indicators require single wavelength emission for detection, they are suitable for multicolor imaging. To demonstrate the advantage of single-FP indicators, we expressed both Green Falcan 10 and R-GECO,¹³ a red fluorescent Ca²⁺ indicator, in HeLa cells. After the administration of 100 μM forskolin and 10 μM histamine which causes Ca²⁺ elevation by stimulation of the histamine H1 receptor, the fluorescence intensity (FI) of both Green Falcan 10 and R-GECO was increased (Fig. 4A and C). We next expressed nuclear localization signal (nls)-Green Falcan 10 and R-GECO in HeLa cells to examine the applicability for visualization of dynamics of different molecules in a distinct intracellular compartment. After the administration of 100 μM forskolin and 10 μM histamine, the FI of both nls-Green Falcan 10 and R-GECO was increased (Fig. 4B and D). These results suggest that Green Falcans are applicable to dual-color imaging with other indicators, and to further hierarchical and cooperative analysis by multi-color imaging. Interestingly, the FI increase in the cytoplasm by the administration of forskolin and histamine was a little smaller than in the nucleus and was also smaller than by the administration of forskolin alone. These results could be ascribed to the accelerated degradation of cAMP by phosphodiesterase³⁷ in the cytoplasm due to calcium elevation.



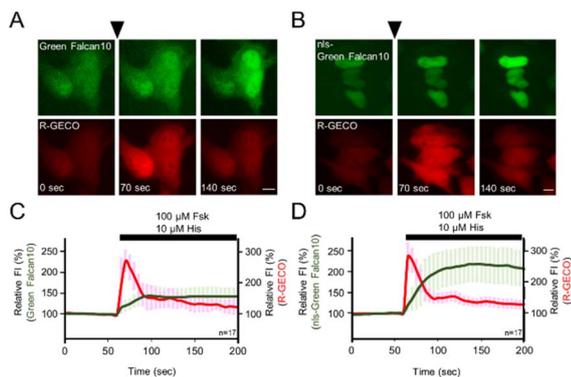


Fig. 4 Dual-color imaging of Green Falcan and R-GECO (A and B) sequential pseudo-color images of Green Falcan 10 and R-GECO (A), nls-Green Falcan 10 and R-GECO (B) expressing HeLa cells, scale bar represents 10 μm (C and D) time courses of fluorescence intensity (FI) in Green Falcan 10 (C)/nls-Green Falcan 10 (D) and R-GECO expressing HeLa cells in response to 100 μM forskolin and 10 μM histamine. Relative FI was calculated by dividing by the mean of the FI during 1 min prior to administration. The data represent the means ± standard deviation. The numbers in the lower right corner of the graphs represent the number of cells analyzed from three independent experiments.

4. Conclusion

In this study, we developed green fluorescent cAMP indicators with various EC_{50} by protein engineering, named Green Falcan 0.3, 1, 3 and 10. These indicators were able to detect cAMP dynamics in the cytoplasm and nucleus of living cells with high spatiotemporal resolution after the administration of several reagents and allowed dual-color imaging with R-GECO. Moreover, Green Falcan 0.3 was applicable to visualize a low concentration range of cAMP dynamics compared to previously developed cAMP indicators. Therefore, we expect that the Green Falcans will visualize unknown phenomena to unravel new physiological mechanism related to cAMP signaling in many organisms.^{38,39}

Conflicts of interest

There are no conflicts to declare.

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Notes and references

- 1 S. R. Adams, A. T. Harootunian, Y. J. Buechler, S. S. Taylor and R. Y. Tsien, *Nature*, 1991, **349**, 694–697.
- 2 A. Koschinski and M. Zaccolo, *Sci. Rep.*, 2017, **7**, 14090.
- 3 S. Börner, F. Schwede, A. Schlipp, F. Berisha, D. Calebiro, M. J. Lohse and V. O. Nikolaev, *Nat. Protoc.*, 2011, **6**, 427–438.
- 4 D. Willoughby and D. M. F. Cooper, *Physiol. Rev.*, 2007, **87**, 965–1010.
- 5 J. B. Shabb, *Chem. Rev.*, 2001, **101**, 2381–2411.
- 6 M. Grandoch, S. S. Roscioni and M. Schmidt, *Br. J. Pharmacol.*, 2010, **159**, 265–284.
- 7 U. B. Kaupp and R. Seifert, *Physiol. Rev.*, 2002, **82**, 769–824.
- 8 G. Tolhurst, Y. Zheng, H. E. Parker, A. M. Habib, F. Reimann and F. M. Gribble, *Endocrinol.*, 2011, **152**, 405–413.
- 9 A. Nakajima, S. Ishihara, D. Imoto and S. Sawai, *Nat. Commun.*, 2014, **5**, 5367.
- 10 E. R. Kandel, *Mol. Brain*, 2012, **5**, 14.
- 11 T. Nagai, A. Sawano, P. Eun Sun and A. Miyawaki, *Proc. Natl. Acad. Sci. U. S. A.*, 2001, **98**, 3197–3202.
- 12 J. Nakai, M. Ohkura and K. Imoto, *Nat. Biotechnol.*, 2001, **19**, 137–141.
- 13 Y. Zhao, S. Araki, J. Wu, T. Teramoto, Y. F. Chang, M. Nakano, A. S. Abdelfattah, M. Fujiwara, T. Ishihara, T. Nagai and R. E. Campbell, *Science*, 2011, **333**, 1888–1891.
- 14 T. Kitaguchi, M. Oya, Y. Wada, T. Tsuboi and A. Miyawaki, *Biochem. J.*, 2013, **450**, 363–373.
- 15 H. Odaka, S. Arai, T. Inoue and T. Kitaguchi, *PLoS One*, 2014, **9**, e100252.
- 16 K. Harada, M. Ito, X. Wang, M. Tanaka, D. Wongso, A. Konno, H. Hirai, H. Hirase, T. Tsuboi and T. Kitaguchi, *Sci. Rep.*, 2017, **7**, 7351.
- 17 S. Matsuda, K. Harada, M. Ito, M. Takizawa, D. Wongso, T. Tsuboi and T. Kitaguchi, *ACS Sens.*, 2017, **2**, 46–51.
- 18 P. H. Tewson, S. Martinka, N. C. Shaner, T. E. Hughes and A. M. Quinn, *J. Biomol. Screening*, 2016, **21**, 298–305.
- 19 C. R. Hackley, E. O. Mazzoni and J. Blau, *Sci. Signaling*, 2018, **11**, eaah3738.
- 20 S. Kawata, Y. Mukai, Y. Nishimura, T. Takahashi and N. Saitoh, *Proc. Natl. Acad. Sci. U. S. A.*, 2022, **119**, e2122618119.
- 21 L. Wang, C. Wu, W. Peng, Z. Zhou, J. Zeng, X. Li, Y. Yang, S. Yu, Y. Zou, M. Huang, C. Liu, Y. Chen, Y. Li, P. Ti, W. Liu, Y. Gao, W. Zheng, H. Zhong, S. Gao, Z. Lu, P. G. Ren, H. L. Ng, J. He, S. Chen, M. Xu, Y. Li and J. Chu, *Nat. Commun.*, 2022, **13**, 5363.
- 22 M. Mita, D. Wongso, H. Ueda, T. Tsuboi and T. Kitaguchi, *Methods Mol. Biol.*, 2021, **2274**, 89–100.
- 23 S. Arai, R. Kriszt, K. Harada, L. S. Looi, S. Matsuda, D. Wongso, S. Suo, S. Ishiura, Y. H. Tseng, M. Raghunath, T. Ito, T. Tsuboi and T. Kitaguchi, *Angew. Chem., Int. Ed.*, 2018, **57**, 10873–10878.
- 24 M. Mita, M. Ito, K. Harada, I. Sugawara, H. Ueda, T. Tsuboi and T. Kitaguchi, *Anal. Chem.*, 2019, **91**, 4821–4830.
- 25 C. H. Clegg, G. G. Cadd and G. S. McKnight, *Proc. Natl. Acad. Sci. U. S. A.*, 1988, **85**, 3703–3707.



- 26 I. Ghosh, A. D. Hamilton and L. Regan, *J. Am. Chem. Soc.*, 2000, **122**, 5658–5659.
- 27 R. A. Steinberg, M. M. Symcox, S. Sollid and D. Øgreid, *J. Biol. Chem.*, 1996, **271**, 27630–27636.
- 28 J. L. Sebaugh and P. D. McCray, *Pharm. Stat.*, 2003, **2**, 167–174.
- 29 W. S. Rasband, *ImageJ*, U. S. National Institutes of Health, Bethesda, Maryland, USA, 2018, <https://imagej.nih.gov/ij/>.
- 30 R Core Team, *R: A Language and Environment for Statistical Computing*, R Foundation for Statistical Computing, Vienna, Austria, 2022, <https://www.R-project.org/>.
- 31 Y. Kanda, *Bone Marrow Transplant.*, 2013, **48**, 452–458. <https://www.jichi.ac.jp/saitama-sct/SaitamaHP.files/statmed.html>.
- 32 J. Jumper, R. Evans, A. Pritzel, T. Green, M. Figurnov, O. Ronneberger, K. Tunyasuvunakool, R. Bates, A. Židek, A. Potapenko, A. Bridgland, C. Meyer, S. A. A. Kohl, A. J. Ballard, A. Cowie, B. Romera-Paredes, S. Nikolov, R. Jain, J. Adler, T. Back, S. Petersen, D. Reiman, E. Clancy, M. Zielinski, M. Steinegger, M. Pacholska, T. Berghammer, S. Bodenstein, D. Silver, O. Vinyals, A. W. Senior, K. Kavukcuoglu, P. Kohli and D. Hassabis, *Nature*, 2021, **596**, 583–589.
- 33 K. Harada, T. Chihara, Y. Hayasaka, M. Mita, M. Takizawa, K. Ishida, M. Arai, S. Tsuno, M. Matsumoto, T. Ishihara, H. Ueda, T. Kitaguchi and T. Tsuboi, *Sci. Rep.*, 2020, **10**, 19562.
- 34 R. Heim, D. C. Prasher and R. Y. Tsien, *Proc. Natl. Acad. Sci. U. S. A.*, 1994, **91**, 12501–12504.
- 35 D. Stoner-Ma, A. A. Jaye, K. L. Ronayne, J. Nappa, S. R. Meech and P. J. Tonge, *J. Am. Chem. Soc.*, 2008, **130**, 1227–1235.
- 36 K. Brejc, T. K. Sixma, P. A. Kitts, S. R. Kain, R. Y. Tsien, M. Ormö and S. J. Remington, *Proc. Natl. Acad. Sci. U. S. A.*, 1997, **94**, 2306–2311.
- 37 T. A. Goraya and D. M. F. Cooper, *Cell. Signal.*, 2005, **17**, 789–797.
- 38 I. Uno, K. Matsumoto and T. Ishikawa, *J. Biol. Chem.*, 1983, **258**, 3539–3542.
- 39 P. Milanese, A. Arce-Rodríguez, A. Muñoz, B. Calles and V. de Lorenzo, *Environ. Microbiol.*, 2011, **13**, 324–339.

