Montana Molecular

Green cADDis[™] cAMP Assays

Product Info & Protocol

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Green cADDis™ cAMP Assays

About these Assays

Cyclic AMP (cAMP) is an essential second messenger for many cellular processes. The messages carried by cAMP are tightly regulated within cells. The cADDis assays detect changes in cAMP in living cells. We also have cADDis assays that can be targeted to specific populations of cells in mixed cultures or to subcellular compartments; contact us to discuss these additional tools. The cADDis assays for cAMP can be combined with different colored sensors to measure multiple signals simultaneously.

Depending upon the sensor in your kit (Downward or Upward), fluorescence either decreases or increases. For example, #D0250G Green Down cADDis **decreases in fluorescence** when cAMP levels **increase** in the cell. Both cADDis sensor versions are bright, robust, and easy to detect on fluorescence plate readers (Fig. 6). For sensitive and/or difficult to transduce cell types, such as primary cultures, we offer purified, high-titer BacMam stock. Please contact us to discuss whether purified BacMam is the right product for you.

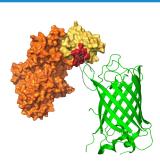


Figure 1. cADDis sensor is comprised of a circularly permuted mNeonGreen fluorescent protein fused to the hinge region of EPAC-2. EPAC-2 hinge domain shown in red, regulatory doman in yellow, catalytic domain in orange.

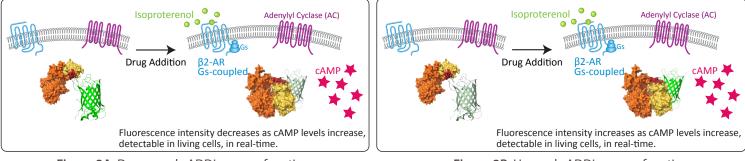
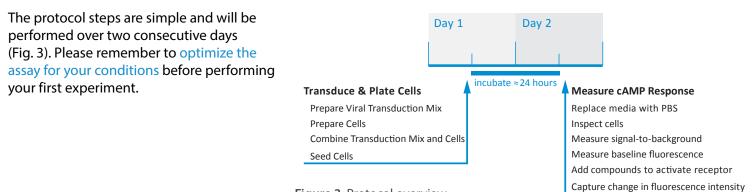


Figure 2A. Downward cADDis sensor function

Figure 2B. Upward cADDis sensor function

The cADDis Assay protocol is optimized for measuring cAMP responses in rapidly dividing, immortalized cell lines on a 96-well plate, and has been validated in live HEK 293T cells [1]. cADDis users have published results in a wide variety of cell types. This assay is very robust and can be used for live-cell imaging or for screening on automated fluorescence plate readers. For use in iPSC-derived or adherent cells, see Suggestions for Assays in Adherent Cells section. For use in difficult to transduce cell types, see Alternative Spinoculation Protocol section.





| Table 1. c | ADDis cAMP Assay Kits | | |
|------------|--------------------------|----------|---|
| Product | Description | Promoter | Recommended Use |
| #D0200G | Green Down cADDis cAMP | CMV | Fluorescence imaging and plate reader assay ($Z' > 0.9$) |
| #D0250G | Green Down cADDis cAMP * | CMV | Fluorescence imaging and plate reader assay ($Z' > 0.9$) |
| #D0205G | Green Down cADDis cAMP | CAG | Fluorescence imaging and plate reader assay |
| #U0200G | Green Up cADDis cAMP | CMV | Fluorescence imaging and plate reader assay ($Z' > 0.85$) |
| #U0250G | Green Up cADDis cAMP* | CMV | Fluorescence imaging and plate reader assay ($Z' > 0.85$) |
| #U0205G | Green Up cADDis cAMP | CAG | Fluorescence imaging and plate reader assay |

*These products have been packaged in Montana Molecular's improved Big Sky BacMam vector

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Assay Kit Materials and Storage

BacMam stocks should be stored at 4°C protected from light in the original package. Store control agonist at -20°C. Avoid repeated freeze/thaw cycles. We recommend re-testing BacMam stock after storing for more than 12 months. If your BacMam stock has been purified, use it within 30 days for best results. Store sodium butyrate at 4°C.

| Table 2. | Materials in Kit | Details | Storage |
|----------------------------|--|--|---------|
| $\approx 2 \times 10^{10}$ | AMP sensor BacMam /G/mL in ESF 921 Insect Culture Medium on Systems, product #96-001-01) | Green fluorescent sensor that changes in fluorescence intensity in response to increases in cAMP. VG/mL is the number of viral genes per milliliter (see Biosafety Considerations section). | 4°C |
| SB | sodium butyrate (Sigma Aldrich product #B5887) 500 mM in H ₂ O | Sodium butyrate is added to the culture to maintain BacMam expression. Other HDAC inhibitors may work as well. | 4°C |
| Rs | β2 adrenergic receptor BacMam in ESF 921 Insect Culture Medium (Expres- sion Systems, product #96-001-01) | A Gs-coupled receptor provided as a positive control for the purpose of assay optimization. Contains a separate red fluorescent protein that is tar- geted to the nucleus. (The control receptor included in CAG-promoter kits does not have a separate fluorescent protein.) | 4°C |
| As | isoproterenol 10 mM (Sigma Aldrich Product Number I2760) in 10 mM HCI | Isoproterenol can be used to stimulate Gs signaling through the positive control, the $\beta 2$ adrenergic receptor. | -20°C |

Additional Materials Required (not included in kit)

- 1. Black, clear bottom microplate coated with a cell attachment factor. We recommend the following plates; 96-well Greiner Bio-One (#655946), 96-well Thermo Fisher Scientific (#152037), 384-well Greiner Bio-One: (#781946), 384-well Corning: (#354663).
- 2. Dulbecco's Phosphate Buffered Saline 1X with Ca²⁺ and Mg²⁺ (10X solution from Gibco #14080055) [2].
- 3. Cells and cell media. We recommend media with low autofluorescence such as EMEM, McCoy's 5A, and F12K culture media.

Biosafety Considerations

The BacMam vector carrying the fluorescent biosensor in these assays is a modified baculovirus, used for delivery to, and expression in, a wide variety of mammalian cell types including primary cultures.

BacMam is a modified baculovirus, *Autographa californica*, AcMNPV. The natural host of baculovirus is larvae of the order *Lepidoptera*. The BacMam vector in the kit is produced in Sf9 insect cells and is pseudotyped to infect mammalian cells. In mammalian cells, the baculovirus genome is silent and it cannot replicate to produce new virus. While it should be handled carefully, in a sterile environment, it is classified as a Biosafety Level 1 (BSL-1) reagent [9].

Other types of viruses are quantified in terms of plaque forming units (PFU) in cells from the natural host. Since BacMam is modified to produce expression in mammalian cells, we quantify the virus by measuring viral genes (VG) per milliliter (mL). Viral samples are prepared to release viral genomic DNA, then multiple dilutions of the preparation are run in qPCR using primers that are specific to the VSVG gene in the BacMam genome. Results are compared against a standard curve to generate an average titer for each viral stock. Check the label on the tube to find VG/mL for your stock.

This product is for research use only and is not for use or sale in human or animal diagnostic or therapeutic products.

Terms of Sale

All materials in this kit are provided without warranty, express or implied. User is responsible for making sure product use complies with applicable regulations. No right to resell products or any components of these products is conveyed. Use of materials is restricted to the intended purpose described in this protocol. Reverse engineering or modification of materials is not permitted. User agrees to accept these Terms of Sale before using materials.

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Protocol for Use

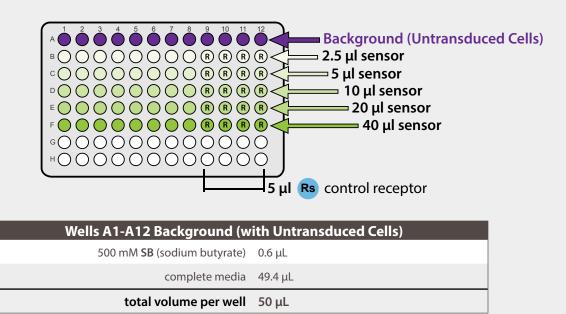
This protocol is optimized for use in HEK 293T cells, however, it can be adjusted for use with practically any cell type. **Take the time to optimize the assay for your cell type and your particular conditions.** See our suggestions for Assays in Adherent Cell Types, Scaling for 384-well Plates, and Alternative Spinoculation Protocol.

Before your first experiment: Optimize Your Assay by Titrating the Sensor

We recommend using a titration series to determine the best combination of signal above background, cell health, and sensor expression. Ideally, the signal in each well before drug addition should be at least 5 times above background.

Transduce and Plate Cells for Optimization

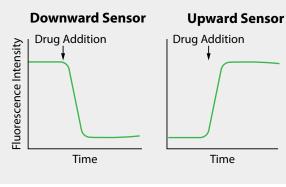
- i. Set up your plate. Be sure to include control wells (untransduced cells) in order to calculate signal-to-background. (For details on preparing cells and transduction mix, refer to Day 1 procedure on page 5).
- ii. Perform titration to determine optimal sensor volume for your cells, as detailed in plate diagram and tables below.
- iii. For each sensor volume, include a subset of wells that includes 5 µl of **Rs** control receptor included in your kit. These wells will be used for your positive control.



| | | total volume per well | 50 μL |
|------------------------------|---------|---|--------|
| total volume per well | 50 μL | L complete media 41.9 | |
| complete media | 46.9 μL | Rs (β2 adrenergic control receptor)) | 5 µL |
| 500 mM SB (sodium butyrate) | 0.6 μL | 500 mM SB (sodium butyrate) | 0.6 µL |
| cADDis cAMP Sensor | 2.5 μL | cADDis cAMP Sensor | 2.5 μL |
| Wells B1-B8 Transduction Mix | | Wells B9-B12 Transduction Mix | |

Measure Parameters to Determine Optimal Conditions

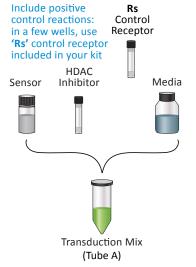
iv. Determine optimal sensor volume by analyzing fluorescence above background, cell health, and response to drug. (For details on measuring fluorescence, refer to Day 2 procedure.)





Day 1 – Transduce and Plate Cells for your Experiment

- A. Prepare Viral Transduction Mix (Tube A): For each transduction reaction (i.e. one well in a 96-well plate), prepare the transduction mix as detailed in table at right (using the optimal volume of sensor that was determined in your optimization experiment). Mix gently.
- B. **Prepare Cells (Tube B):** Detach cells using standard trypsinization protocol. Resuspend cells in complete culture media and determine cell count.
- C. Prepare a dilution of cells at your desired concentration. (100 μL of this cell resuspension will be required for a single well in a 96-well plate, so prepare enough of the dilution to seed the desired number of wells in the plate).
- D. Combine Transduction Mix and Cells: Combine Tube A and Tube B (50 μL Tube A + 100 μL Tube B). Mix by pipetting up and down gently, and seed 150 μL per well on the 96-well plate.
- E. Cover plate to protect from light and let rest at room temperature for 45-60 minutes.
- F. Incubate \approx 20-24 hours under normal cell growth conditions (5% CO₂ and 37°C), protected from light.



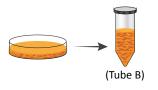
Tip: When preparing a master transduction mix, to avoid coming up short, scale up by 10-15% of the number of wells needed.

| Transduction Mix (T | ube A) |
|--------------------------------|------------|
| cADDis cAMP Sensor | Variable |
| SB 500 mM sodium butyrate | 0.6 µL |
| Rs control receptor (if using) | 5 μL |
| complete media | (To 50 μL) |
| | |

total volume per well 50 μL

| Cells (Tube B) | |
|----------------|--|
| | |

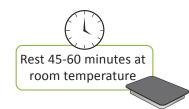
cells per well 100 µL



| Transduction Mix + (Tube A) + (Tube I | |
|--|--------|
| transduction mix | 50 µL |
| cell suspension | 100 μL |

total volume per well $150 \ \mu L$

Tip: 90% cell confluence after 24 hours is ideal. For HEK 293T cells 50,000 cells per well works well but plating density is cell type dependent

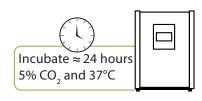


Tube

Α

Tube

R



Suggestions for Assays in Adherent Cells

This protocol is optimized for rapidly dividing immortalized cells.

However, cADDis biosensor assays can be used with primary cultures of cells as well as iPSC-derived cells. In the case of nondividing, or differentiated cells, the transduction should be done with adherent cells, and the media should be exchanged the following day. Specific details of the protocol will vary by cell type, so it is important to take the time to titrate BacMam for optimal results. For expression in rare cell types, or specific cells in mixed cultures, Cre-dependent and specific promoter systems are available for many of our sensors.

*When working with adherent cell cultures, add an extra day to seed the cells prior to transduction.

Adherent Cells Day 1:

• Seed the cells the day before you add transduction mix, incubate 24 hours

Adherent Cells Day 2:

- Perform transduction as directed on page 5, but add the transduction reaction directly to the plated cells (no aspiration of cell medium necessary). Gently rock the plate 4-5 times in each direction to mix throughout the well. Incubate the cells under normal growth conditions (5% CO₂ and 37°C), protected from light, for 20-24 hours.
- Optional step (cell type dependent): After 4-8 hour incubation with sensor BacMam (6 hours is optimal), very gently aspirate transduction solution (we recommend using a plate washer). Add 100 µL complete growth media with sodium butyrate at a concentration of 2 mM. If cells will not tolerate a full media exchange, partial media exchanges can be done.

Adherent Cells Day 3:

• Measure Fluorescence as as detailed in Day 2 procedure.

Scaling for 384 Wells (1 plate)

To set up the assay in a 384-well plate, follow all of the protocol steps, adjusting reagent volumes as follows:

| Transduction Mix (Tu (384-well plate) | |
|--|---------|
| cADDis cAMP Sensor | 2 µL |
| 500 mM SB sodium butyrate | 0.1 μL |
| Rs control receptor | 1 μL |
| complete media | 9.4 μL |
| total volume per well | 12.5 μL |

Cells (Tube B) (384-well plate)

cells per well 12.5 µL

12.5 μ L of the cell resuspension will be required for a single well in a 384-well plate. A plating density of 7,500 cells per well is a good starting point, so prepare the cell suspension at 600,000 cells/mL. Depending on the cell type and plate type, 5,000-15,000 cells per well may be optimal.

| Transduction Mix + Cells (Tube A) + (Tube B) (384-well plate) |
|---|
| transduction mix 12.5 µL |
| cell suspension 12.5 µL |

total volume per well 25 µL

Tip: When scaling for 384-well plates, the volume of the cell suspension per well can be increased to $50 \ \mu$ L to improve cell health. Make sure that the well is still receiving 7,500 cells and that you increase the volume of sodium butryate to 0.2 μ L per well. 7

Alternative Spinoculation Protocol

This alternative protocol may also be useful for particularly difficult-to-transduce or low expressing cell types:

- 1. Prepare transduction mix (detailed in table at right).
- 2. Detach cells using standard trypsinization protocol. Resuspend cells in complete culture media and perform cell count.
- 3. Prepare a dilution of cells at your desired concentration (22,500 cells/well in a 96-well plate is a good starting point, but will ultimately depend on the cell type being used). 50 μL of this cell resuspension will be required for a single well in a 96-well plate, so prepare enough of the dilution to seed the desired number of wells in the plate.
- Combine the transduction mix with the cell suspension
 (50 μL transduction mix + 50 μL cells). Mix gently, then seed
 100 μL of this mix per well on a 96-well plate.
- 5. Let cells sit at room temperature, protected from light, for 20 minutes.
- 6. Spin the plate at 1,500 x g for 1.5-2 hours at room temperature.
- * We recommend sealing the plate with Breathe-Easy[®] (Cat. No. 70536-10) during this step to avoid contamination.
- After spinning the plate, remove the transduction mix and replace with fresh media containing 0.6 μL sodium butyrate (2 mM per well).
- 9. Return plate to normal growth conditions and incubate for 48 hours.

| see optimization page | |
|-----------------------|--|
| | |
| | |
| | |

Tip: *Titrate the sensor to*

determine optimal volume;

If the above protocol does not result in acceptable expression levels, we recommend using System Biosciences' Spinoculation protocol, which adds SBI's TransDux Max and MAX enhancer reagents to the transduction reaction. Please make the following adjustments to your transduction reaction:

| Transduction Mix + Enhancer | Reagents |
|-----------------------------------|------------|
| cADDis cAMP Sensor | Variable |
| TransDux | 0.34 μL |
| Max Enhancer | 17 μL |
| 500 mM SB sodium butyrate) | 0.6 µL |
| 1M HEPES (pH 7.4) | 0.7 μL |
| cell culture media | (To 50 μL) |
| total volume per well | 50 μL |

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Alternative Transduction Mix

| total volume per well | 50 μL |
|---------------------------|------------|
| cell culture media | (To 50 μL) |
| 1M HEPES (pH 7.4) | 0.7 μL |
| 500 mM SB sodium butyrate | 0.6 µL |
| cADDis cAMP Sensor | Variable |
| | |

Day 2– Measure cAMP Response

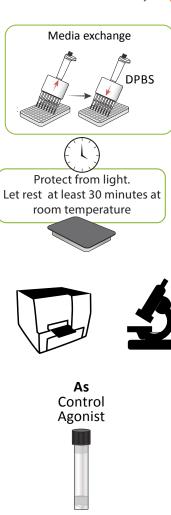
- G. Prior to measuring fluorescence, replace culture media with DPBS (1X, containing Ca²⁺ and Mg²⁺). Wash gently 4-5 times, careful not to dislodge cells. We recommend using a plate washer. (We recommend the BioTek 405 TS.)
- H. Cover the cells. Allow to equilibrate at room temperature at least 30 mintues before measuring fluorescence. Experiments can be performed at room temperature
- I. Visually inspect cells on microscope to confirm cell health. Transduced cells should be at least 5× brighter compared to untransduced cells. (If you have high back-ground fluorescence, gently wash cell culture media again.)

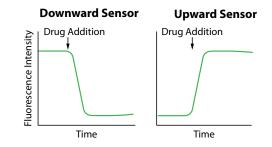
Measure Fluorescence on Plate Reader or Imaging System

- J. Measure fluorescence in transduced and untransduced cells. Transduced cells should be at least 5× brighter compared to untransduced cells.
- K. Acquire 10-20 baseline fluorescence reads before adding compounds.
- L. Activate the receptor with agonist to affect levels of cAMP. Resume measurement immediately after adding drug. Measure continuously for a minimum of 30 minutes, (and up to 5 hours at 15-60 second intervals on a plate reader). A change in fluorescence intensity under standard GFP excitation and emission will indicate a change in cAMP levels. For wells transduced with the β 2 adrenergic control receptor, add 10 μ M As (isoproterenol, final concentration in well) to activate the receptor. The optimal dose of agonist may need to be determined for a given cell line.
 - * **Important:** Make sure that the final concentration of any drug solvents in the well (e.g. DMSO) does not exceed 1%. The ideal concentration for DMSO is 0.1%.

Tip: Add agonists at a concentration of $3-4 \times$ (in a volume of 50 µL DPBS) to wells containing 100-150 µL DPBS. This will result in a 1:3 or 1:4 dilution, and allow for adequate diffusion.

Tip: Always test the addition of the vehicle alone (i.e. DPBS without drug).





Fluorescence Detection

Our assays are compatible with automated fluorescent plate readers and imaging systems.

We have validated on:

- Agilent (BioTek) Synergy
- Agilent (BioTek) Cytation[™]
- BMG CLARIOstar®
- Agilent (BioTek) Lionheart
- Agilent (BioTek) Neo
- Hamamatsu FDSS®
- Epifluorescence microscopes

| Our customers have reported good results on: |
|--|
|--|

- Molecular Devices FLIPR[®]
- Molecular Devices Flexstation®
- Perkin Elmer Enspire®
- Perkin Elmer Opera Phenix ®

Tip: Below are the specific settings recommended for use with our assays for instruments we use at Montana Molecular. To determine the best settings for fluorescence detection on your instrument, please consult the manufacturer.

| Table 3. CLARIOstar [®] Recommendations | | |
|--|------------------------|--|
| Instrument Settings | | |
| Detection Mode: FI (Bottom) | | |
| Detection Method: Plate Mode, Kinetic | | |
| Scan Mode: Orbital Averaging | | |
| Scan Diameter (mm): 4 | | |
| Gain/Focal Height: | Adjusted prior to test | |
| Optical Settings | | |
| Excitation: | F 482-16 | |
| Dichroic: | LP 504 | |
| Emission: | F 530-40 | |

| Table 4. Neo & MX™ Recommendations | | | |
|------------------------------------|----------------------------|--|--|
| Instrument Settings | | | |
| Detection Method: | Fluorescence Intensity | | |
| Read Type: | Endpoint/Kinetic | | |
| Optics: | Monochromators | | |
| Excitation: | 485 nm | | |
| Emission: | 528 nm | | |
| Bandwidth: | 20 nm (for both ex and em) | | |
| Optics Position: | Bottom | | |
| Gain: | 100 | | |

| Table 5. Cytation™ & Lionheart Recommendations | | |
|--|-------------------------------|--|
| Image Preprocessing | | |
| Image Set: | GFP | |
| Background: | Dark | |
| Rolling Ball Diameter: Automatic | | |
| Image Smoothing: | 0 cycles | |
| Cellular Analysis Parameters | | |
| Channel: | Tsf [GFP] | |
| Threshold: | 7,000 | |
| Background: | Dark | |
| Split Touching Objects: | Checked | |
| Fill Holes in Mask: | Checked | |
| Minimum Object Size: | 5 μm | |
| Maximum Object Size: | 1,000 μm | |
| Include Primary Edge Objects: | Checked | |
| Analyze Entire Image: | Checked | |
| Advanced Detection Options | | |
| Rolling Ball Diameter: | Automatic | |
| Image Smoothing Strength: | 1 cycle of 3×3 average filter | |
| Evaluate Background On: | 5% | |
| Primary Mask: | Use threshold mask | |



Fluorescence Properties

Green cADDis is constructed with the very bright, mNeonGreen fluorescent protein [6]. While the peak excitation and emission wavelengths are 506 nm and 517 nm, respectively, a range of 485-505 nm (excitation) and 515-535 nm (emission) may be used if your instrument does not allow measurement at the peak ex/ em. For example, on the BioTek Synergy MX[™], the preferred ex/em is 488/525. If using filters, we recommend Chroma's Catalog set #49003 for optimal results.

Tip: We offer mNeon BacMam Kits

[products **#P1000G (CMV)**, **#P1005G (CAG)** and **#P1001G** (Synapsin)]. These test kits are a good way to determine BacMam transduction efficiency, evaluate promoter systems, and optimize expression in your cells of choice.

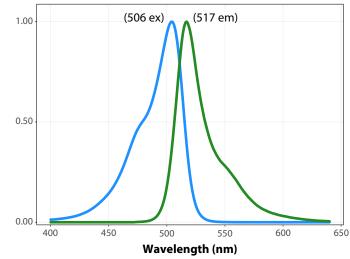


Figure 4. Absorption and emission properties of the mNeon-Green fluorescent protein plotted as a function of wavelength.

Timing

Unlike many assays that measure accumulation of cAMP in cell lysates, the cADDis assay measures cAMP in living cells, in real time. For best results, be sure to capture changes in cAMP during the peak response. In Figure 5, fluorescence was captured from cells before the addition of the drug and then sampled at regular intervals. The maximal response is reached at 5-10 minutes after the addition of the drug.

Data Analysis

Check out our helpful reference document "Biosensor Analysis Overview"

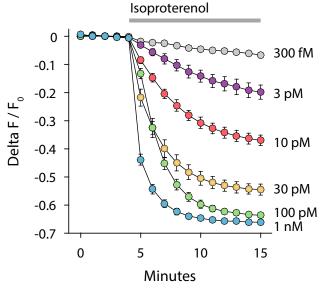


Figure 5. HEK 293T cells transduced with 20 μ L of BacMam carrying Green Down cADDis cAMP biosensor(#D0200G), activated with isoproterenol. The graph shows the expected timing and duration of cAMP when using the positive controls provided in your kit.

Assay Optimization

Optimizing Fluorescence

Twenty-four hours after transduction, check your cells for fluorescence. Wells that were transduced with the sensor should be 5-10 times brighter than control wells that were not treated with the sensor.

HDAC inhibitors may be used to maintain expression of the sensors. While BacMam transduction alone will result in sensor expression, sodium butyrate or another HDAC inhibitor, such as valproic acid (VPA) or trichostatin A (TSA), will generate higher levels of expression and will maintain this level of expression [4]. If cells look unhealthy, use lower concentrations or no HDAC inhibitor.

The type of cell culture media used in your experiment can affect the transduction efficiency of BacMam. Our assays have been validated in EMEM, McCoy's 5A, and F12K culture media Cell culture media with high sodium bicarbonate content can interfere with transduction efficiency [8]. If your cell culture media of choice is affecting transduction efficiency, transduction can be conducted in DPBS for 6 hours and then the cells can be returned to complete cell culture media containing 2 mM sodium butyrate.

Optimizing Expression for Your Cell Type

To determine optimal conditions for your cell type, prepare a dilution series of transduction reactions by varying the amount of sensor. For example, a range of 2.5 μ L to 40 μ L is a good starting range in a 96-well plate (Fig. 6). Choose the range that gives you at least 5-10 times above background and gives you the highest amount of fluorescence change with your Rs control receptor.

Varying the cell density, concentration of sodium butyrate, or trying a new HDAC inhibitor (VPA or TSA) may boost expression as well. optimizing to avoid sensor saturation

Please contact us if you would like to use the sensor under the control of a specific promoter system. Sensors under weak promoters may be limited to detection on imaging systems. To maintain strong expression in specific cell types, we recommend ordering a Cre-inducible, floxed sensor.

Purified viral preparations, which can increase expression in particularly sensitive or difficult to transduce cell types, are available upon request.

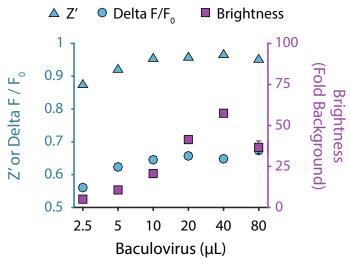


Figure 6. As the amount of cADDis added to the wells increases, so does the baseline fluorescence, plotted in purple. The change in fluorescence when cAMP changes also increases with more virus, but reaches the maximum possible change and remains constant over a broad range of virus concentration (circles). Z' is high over a broad range of virus concentration (triangles).

Use the Positive Control

If the cells are expressing the sensor, and fluorescence is detectable on your instrument, then evaluate sensor performance using the positive control receptor included in your kit. Add 5 μ L of the **Rs** (β 2 adrenergic control receptor) to a set of control wells and activate with isoproterenol included in the kit (Fig. 5).

Optimizing Receptor Expression

If you have titrated the cADDis sensor and determined the optimal volume, but fail to see a receptor mediated signal, the receptor expression level may be the issue. Try titrating the receptor with a fixed amount of the cADDis sensor.

If you need further help, see the Troubleshooting Guide or let us know, we're happy to help!

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Troubleshooting Guide

| Problem | Possible Cause | Solution |
|---|---|--|
| Low sensor expression and/or poor transduction efficiency | Suboptimal sensor BacMam volume is being used. | Perform titration of the sensor BacMam stock, testing a large range (i.e. 2.5-40 μL in 96-well plate format) to identify optimal volume. Too little can result in low expression, too much can cause cells to become sick. |
| | Transducing adherent cells. | Transduce cells while in suspension. If this isn't possible, try doing a media exchange on adherent cells after 4-6 hours, in addition to leaving the virus on overnight. See Suggestions for Assays in Adherent Cells. |
| | Suboptimal cell density; too few or too many cells added. | Transduce cells so that they are between 80-90% confluent at the time ofthe experi- ment. Also, when transducing cells in suspension, make sure that cells in the source flask are < 100% confluent (approximately 80% confluent is ideal). |
| | | Add HDAC inhibitor at the proper concentration: |
| | HDAC inhibitor was not | sodium butyrate - 2mM final concentration |
| | added to the transduction mix, or the concentration | valproic acid - 5mM final concentration |
| | was wrong. | trichostatin A - 0.25µM final concentration |
| | | * Perform a titration to determine optimal concentration for the cell type being used. |
| | HDAC inhibitor being used is not optimal for cell type. | Test other HDAC inhibitors (e.g. sodium butyrate, valproic acid, trichostatin A.) |
| | | After adding transduction mix to cells, let cells sit at room temperature for 45-60 min. before placing back in incubator (longer incubation times at room temperature may further improve transduction). |
| | | Perform media exchange after various incubation times with the transduction mix, in addition to leaving the virus on overnight. |
| | Cell type being used trans- duces poorly. | Try high-titer, purified BacMam stock. |
| | | Validate assay in a different cell type (e.g. HEK 293T cells) |
| | | • Transduce cells multiple times (e.g. on Day 1, and again on Day 2). |
| | | Incubate cells for 48 hours post transduction, before performing assay. |
| | | Consider using a different viral vector, such as lentivirus or AAV. |
| | Cell culture media is inhib- iting transduction. | Remove media during transduction, preparing the transduction mix in DPBS and adding to cells. Replace transduction mix with media after 2-6 hours. |
| | BacMam stock was not stored properly (i.e. not stored at 4°C, exposed to light for long periods, subjected to multiple freeze-thaw cycles), or the shelf life has been exceeded. | Follow guidelines for product storage. BacMam stocks are stable for at least 12 months when stored properly. After this time period, the stock should be re-evaluated and compared to previous experiments. Purified BacMam stocks should be used within 30 days for best results. |
| | BacMam stock was not mixed adequately before transducing cells | Mix BacMam stock thoroughly before transduction, especially after being stored for long periods |
| | Promoter is not optimal for cell type being used. | Identify promoters that work best in the cell type being used. If promoter is not on product list, consult Montana Molecular for custom production services |
| | Cells are contaminated | Monitor cells for bacteria, fungi, etc. |

Green cADDis™ cAMP Assays

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| Problem | Possible Cause | Solution |
|---|---|--|
| Low fluorescence signal on microscope/plate reader. | Low sensor expression, low transduction efficiency. | See solutions for Problem 1. |
| | Excitation/emission settings are not optimal for the sensor being used. | Refer to protocol for the fluorescence spectra of the sensor. Make sure that filter sets or monochromators are aligned with the peak excitation and emission wavelengths of the sensor. |
| | Cells are in cell culture media, and the media is producing a large fluorescent signal (autofluorescence). | Exchange media so that cells are in DPBS at the time of experi- ment. If cell culture media must be used, try using FluoroBrite media. |
| | Wrong microplate type is being used. | Use black, clear-bottom microplates with low autofluorescence. See our recommended plates. |
| | Exposure time or gain setting on instrument is suboptimal. | Test different exposure and gain settings, monitoring how the signal-to-background and noise in the measurement changes. Too high can result in saturation and/or photobleaching; too low can result in noisy data. |
| | Cells were dislodged during media exchange/plate washing. | Make sure that media exchange or plate washing is done gently and does not dislodge cells. Confirm with visual inspection on a microscope . |
| Signal-to-background is low (i.e. cells/wells with sensor are not much brighter than control cells/ wells without sensor). | Low sensor expression, low transduction efficiency. | See solutions for Problem 1. |
| | Excitation/emission settings are not optimal for the sensor being used. | Refer to protocol for the fluorescence spectra of the sensor. Make sure that filter sets or monochromators are aligned with the peak excitation and emission wavelengths of the sensor. |
| | Exposure time or gain setting on instrument is suboptimal. | Test different exposure and gain settings, monitoring how the signal-to-background and noise in the measurement changes. Too high can result in saturation and/or photobleach- ing; too low can result in noisy data. |
| | Media exchange was not performed before running the assay; cells are in media rather than DPBS. Cell culture media being used has high autofluorescence. | Perform media exchange so that cells are in DPBS at the time of experiment. If cell culture media must be used, try using Fluoro- Brite media. |
| | Cells were dislodged during media exchange/plate washing. | Make sure that media exchange or plate washing is done gently and does not dislodge cells. Confirm with visual inspection on a microscope. |
| Signal is noisy. | Low sensor expression, low transduction efficiency. | See solutions for Problem 1. |
| • | Gain setting or exposure time on instru- ment is too low. | Increase gain setting or exposure time. |
| | Media exchange was not performed, or plate washing was inadequate causing high well-to-well variability. Cells are not in DDPBS at the time of experiment. | Exchange media so that cells are in DDPBS at the time of experi- ment. If cell culture media must be used, try using FluoroBrite media. Make sure that plate washing is highly consistent from well to well. |
| | Cells were dislodged during media exchange/plate washing. | Make sure that media exchange or plate washing is done gently and does not dislodge cells. Confirm with visual inspection on a microscope . |
| | Cells are detaching from the plate. | Coat the plate with poly-D lysine or other appropriate cell attach- ment factor. See our recommended plates |

14 Troubleshooting

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| Problem | Possible Cause | Solution |
|--|--|---|
| Good fluorescence signal, but sensor is not responding to drug as expected. No change in fluo- rescence observed, or signal is in the wrong direction. | Photobleaching | Reduce exposure time, sampling rate, and/or light intensity. |
| | Drug is at the wrong concentration | Confirm drug concentration and solubility. |
| | Drug was not stored properly. | Confirm drug storage conditions. |
| | Drug was added to the cells in a volume that was too low relative to the volume of DDPBS/media in the well, resulting in improper mixing. | Add drug in a volume that will allow for sufficient diffusion (i.e. 1:3 or 1:4 drug to total volume) |
| | Drug was not added in the same solution as the solution in the well/culture dish. | Make sure that the drug preparation and cells are in the same solution. |
| | Drug addition is producing an artifact. | Make sure to add a vehicle-only control. Make sure drug is added in a solution that is the same as the solution in the well. Do not exceed 1% DMSO final in the well (0.1% or less is ideal). |
| | Compounds being tested are fluorescent. | Scan compounds for fluorescence to confirm. If possible, dilute compounds in order to reduce the fluorescence artifact of the compound. |
| | Drug addition was too forceful and dislodged cells. | Add drugs manually or with an on-board dispense function, but do so gently, so as not to dislodge cells. |
| | Baseline reads were not acquired before adding drug. | Acquire 10-20 baseline fluorescence reads before adding drug. Monitor for a change in fluorescence intensity upon addi- tion of drug. |
| | Gain setting on instrument is too high, and signal is saturating. Gain setting is too low, and signal cannot be detected. | Adjust gain setting. |
| | Too much sensor has been added to cells and the signal is saturated (i.e. not enough analyte for the amount of sensor in the cell). | Titrate the amount of sensor to determine maximum signal for your cell type. See protocol recommendations for HEK293T cells |
| | Target receptor was not added, or expres- sion levels are suboptimal (too little or too much, or receptor has high level of constitu- tive activity). | Titrate the amount of receptor to optimize the signal for your cell type and receptor. |
| | Sampling rate is not consistent with sensor kinetics. | Acquire 10-20 baseline reads before adding drug. Resume mea- surement quickly after adding drug (within 5-10 seconds for DAG/PIP ₂ , 60 seconds for cADDis and cGMP, and 1-2 seconds for GECO Ca ²⁺). Measure long enough to capture max response of sensor. |
| Poor cell health, cells detaching from plate. | Too much BacMam stock was added to cells (e.g. sensor, receptor, Gs mutant). | Titrate lower amounts of BacMam stock to identify the optimal volume for your cells. |
| | | Confirm concentration of HDAC inhibitor being used. Make new stock solution. Try a different HDAC inhibitor. Confirm that they are being used at the proper concentration: |
| | Concentration of HDAC inhibitor is too high, | sodium butyrate - 2mM |
| | or cells are sensitive to the HDAC inhibitor being used. | valproic acid - 5mM |
| | being used. | trichostatin A - 0.25µM |
| | | * Perform a titration to determine optimal concentration for the cell type being used. |
| | Plate surface is not coated with a cell attachment factor. | Coat plates with a cell attachment factor (e.g. PDL, laminin, col- lagen, fibronectin etc.) to enhance attachment. |
| | Edge wells are being used, and cells in the edge wells may be subject to conditions that are not conducive to growth. | Do not use edge wells. |

15 Troubleshooting / References

Green cADDis™ cAMP Assays

| Problem | Possible Cause | Solution |
|---|---|--|
| Poor cell health, cells detaching from plate. | Too much BacMam stock was added to cells (e.g. sensor, receptor, Gs mutant). | Titrate lower amounts of BacMam stock to identify the optimal volume for your cells. |
| | | Confirm concentration of HDAC inhibitor being used. Make new stock solution. Try a different HDAC inhibitor. Confirm that they are being used at the proper concentration: |
| | Concentration of HDAC inhibitor is too high, | sodium butyrate - 2mM |
| | or cells are sensitive to the HDAC inhibitor being used. | valproic acid - 5mM |
| | | trichostatin A - 0.25μM |
| | | * Perform a titration to determine optimal concentration for the cell type being used. |
| | Plate surface is not coated with a cell attachment factor. | Coat plates with a cell attachment factor (e.g. PDL, laminin, col- lagen, fibronectin etc.) to enhance attachment. |
| | Edge wells are being used, and cells in the edge wells may be subject to conditions that are not conducive to growth. | Do not use edge wells. |
| | Cells were dislodged during media exchange/plate washing. | Make sure that media exchange or plate washing is done gently and does not dislodge cells. Confirm with visual inspection on a microscope.z |
| | DDPBS being used does not contain Ca ²⁺ and Mg ²⁺ . | Use DDPBS containing Ca ²⁺ and Mg ²⁺ . |
| | Cells are contaminated. | Monitor cells for bacteria, fungi, mycoplasma. |
| | Cells were not grown under proper growth conditions (i.e. 5% CO ₂ , 37°C). | Incubate transduced cells at 37°C, in 5% CO ₂ . |
| | Cells are sensitive to acidity from the insect cell culture media present in BacMam virus. The insect cell culture media is more acidic than typical mammalian cell culture media. | Wash transduction mix off of cells after 2-6 hours. Replace with fresh cell culture media, maintaining concentration of sodium butyrate or other HDAC inhibitor. |

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| Related Products | | | |
|------------------|---------------------------|----------|--|
| Product | Description | Promoter | Recommended Use |
| #U0200R | Red Up cADDis cAMP | CMV | Fluorescence imaging and plate reader assay ($Z' < 0.5$) |
| #X0200G | Gi cADDis cAMP | CMV | Fluorescence imaging and plate reader assay ($Z' > 0.7$) |
| #D0300R | Red Down DAG | CMV | Fluorescence imaging and plate reader assay ($Z' > 0.7$) |
| #U0300R | Red Up DAG | CMV | Fluorescence imaging and plate reader assay ($Z' > 0.5$) |
| #U0600R | Red GECO Ca ²⁺ | CMV | Fluorescence imaging and plate reader assay ($Z' > 0.5$) |

Contact Us

If you have any questions about the protocols described here, or if you have ideas about how we can improve these tools, then we want to hear from you. Your feedback is extremely valuable. Please send an email to: info@montanamolecular.com or call us at +1 406-200-8321 and we'll respond as quickly as we can.



US Patent 11,366,114 B2 European Patent Number: EP3065754B1

Our goal is to make your



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workflow easy and reproducible.



We also have a Troubleshooting Guide at the end of this document

