

May 10, 2024

Alternative method to visualize receptor dynamics in cell membranes

DOI

dx.doi.org/10.17504/protocols.io.rm7vzj8z8lx1/v1

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DOI: dx.doi.org/10.17504/protocols.io.rm7vzj8z8lx1/v1

Protocol Citation: Ravelli Cosetta, Corsini Michela, Ventura Anna, Domenichini Mattia, Grillo Elisabetta, Mitola Stefania 2024.

Alternative method to visualize receptor dynamics in cell membranes. **protocols.io**

<https://dx.doi.org/10.17504/protocols.io.rm7vzj8z8lx1/v1>

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Protocol status: Working

We use this protocol and it's working

Created: April 05, 2024

Last Modified: May 10, 2024

Protocol Integer ID: 98426

Keywords: receptor dynamics, VEGFR2, microscopy



Funders Acknowledgement:

Stefania Mitola

Grant ID: AIRC grant IG17276

Stefania Mitola

Grant ID: Consorzio

Interuniversitario per le

Bioteχνologie (CIB)

Abstract

This protocol details alternative method to visualize receptor dynamics in cell membranes.



Cell transfection (day 1 and 2)

1d 4h

- 1 The day before transfection, seed $2 \times 10^4/\text{cm}^2$ of CHO cells in 6 well plate in F12 medium supplemented with 100 IU/mL penicillin and 100 $\mu\text{g}/\text{mL}$ streptomycin and 10% FCS (complete medium) and culture under normal conditions at 37 °C in 5% CO₂. When using different cell lines, ensure a cell density to allow 60-70% of confluence the day of the transfection.
- 2 After 24:00:00 , transfect each well of CHO cells with 4 μg of pBE-hVEGFR2-eYFP and 8 ng of PEI (1 $\mu\text{g}/\mu\text{L}$) in serum and antibiotics free F12 medium.
 - After 04:00:00 , replace medium with complete medium.

1d 4h

Protein immobilization to glass coverslip (day 3)

1d 9h

- 3 Incubate the 2 well chambered glass coverslips with 100 μL of sterile PBS containing 2 $\mu\text{g}/\text{mL}$ of human VEGF-A for 16:00:00 at 4 °C . The coating is carried out by placing a drop containing the recombinant protein in the center of the well in order to obtain a 10 mm diameter coating spot. VEGF-A can be replaced by other ligands able to recruit other specific receptors.
- 4 After 16:00:00 , remove unbound ligand and wash the coverslips 3 times with cold and sterile PBS.
 - 4.1 Wash the coverslips with cold and sterile PBS. (1/3)
 - 4.2 Wash the coverslips with cold and sterile PBS. (2/3)
 - 4.3 Wash the coverslips with cold and sterile PBS. (3/3)
 - Under these conditions, ligand binds to the coverslip in a dose-dependent manner, with maximal binding at coating concentrations $\geq 2 \mu\text{g}/\text{mL}$. Using this concentration, it is possible to have a spot with a high concentration of ligands.
 - Substratum-immobilized ligand is resistant to high molar salt (2 mol/L NaCl) and detergent (0.2% Triton X-100) washes [1,2].

16h



16h





- 5 Block nonspecific binding sites with 1 mg/mL of BSA for 01:00:00 Room temperature .

1h

Cell preparation (day 3)

1d 16h

- 6 Put glass coverslips on the bottom of a 24 well plate and ensure it remains to the bottom of the well while seeding the cells.
- 7 24 hours after cell transfection, plate CHO at the density of 75.000/cm² in complete medium on the coverslips and culture under normal conditions for 16:00:00 . When using different cell lines, ensure a cell density to allow 50-80% of confluence the day of image acquisition.

16h

Image acquisition (day 4)

6h

- 8 Replace the complete medium of transfected cells with F12 (without phenol red) 1% FCS and culture under normal conditions for 02:00:00 .
- 9 After 02:00:00 of starvation, flip upside-down the cell-plated coverslips on immobilized-VEGF chambered in F12 1% FCS.
- 10 Put the sample in the microscope incubator at 37 °C and 5% CO₂.
- To analyze VEGFR2 recruitment we acquired Z-stack images for 02:00:00 .
- 11 Acquire images using YFP fluorescence filter set (excitation: 500/20; dichroic: long pass 512; emission: 535/30).

2h

2h

2h



Note

CRITICAL STEP: It is very important to acquire all the images with the same camera exposure. This allows you to compare different images or different experimental conditions, when necessary.

- 12 Acquire imaging with a PlanApochromat 63X/1.4NA Oil objective and Apotome structured illumination that allow a sectioning of 1.3 µm. Set an overlap of 50% between two consecutive stacks. On average, a whole CHO cell is acquired in 10-12 slices, with a total thickness of acquisition of 13-15.6 µm.





- 13 Process images without deconvolution.

Image quantification and data analysis (day 5)

- 14 Open image series in Fiji as hyperstacks. A sequence of images open, each representing a stack.

- 15 Convert image stack in 8 bits.

- 16 Adjust brightness and contrast in order to clearly see cells in each stack.

Note

That any adjustment in brightness and contrast for the visualization won't modify fluorescence quantification.

- 17 In Analyze > Set Measurements select Area and Area percentage options.

- 18 Open the threshold menu and set threshold manually in order to clearly see the specific fluorescence standing out from the background.



Note

CRITICAL STEP: It is very important to set the same threshold for all the images. This allows you to compare different images or different experimental conditions, when necessary.

- 19 Draw, using freehand selection, the projection of the cell. Analyze one cell at a time. Scroll through the image sequence measuring Area and Area percentage in every stack.



- 20 Save data for the analysis.

21 Calculate the number of pixels positive for VEGFR2 associated fluorescence using the formula:

$$\text{N}^{\circ} \text{ of VEGFR2-positive pixels} = \text{Area percentage} * (\text{Area}/100)$$

22 Sum all the pixel from each Z-stack to obtain the total amount of VEGFR2-positive pixels for cell

23 Calculate the distribution of VEGFR2 in each stack using the formula:

$$\% \text{ of VEGFR2 area} = (\text{N}^{\circ} \text{ of VEGFR2-positive pixels} / \text{total amount of VEGFR2-positive pixels}) * 100$$

Note

That the sum of % of VEGFR2 area from all the stacks should be 100.

3D reconstruction

24 Select a region of interest (ROI) that includes one cell or more. Save the image.

25 Create orthogonal projection by choosing, from Image > Stacks the Orthogonal Views command.

26 Create a 3D image using “3D viewer” plugin.



Protocol references

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