

Feb 28, 2023

# Whole-cell radioligand saturation binding

PLOS One

DOI

dx.doi.org/10.17504/protocols.io.ewov1oxe2lr2/v1

Angus Li<sup>1</sup>, Samuel Liu<sup>1</sup>, Rennica Huang<sup>2</sup>, Seungkirl Ahn<sup>1</sup>, Robert J Lefkowitz<sup>1,2,3</sup>

<sup>1</sup>Department of Medicine, Duke University Medical Center, Durham, North Carolina, United States of America;

<sup>2</sup>Department of Biochemistry, Duke University Medical Center, Durham, North Carolina, United States of America:

<sup>3</sup>Howard Hughes Medical Institute, Duke University Medical Center, Durham, North Carolina, United States of America



Angus Li

### Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN ACCESS



DOI: https://dx.doi.org/10.17504/protocols.io.ewov1oxe2lr2/v1

External link: <a href="https://doi.org/10.1371/journal.pone.0283477">https://doi.org/10.1371/journal.pone.0283477</a>



**Protocol Citation:** Angus Li, Samuel Liu, Rennica Huang, Seungkirl Ahn, Robert J Lefkowitz 2023. Whole-cell radioligand saturation binding. **protocols.io** <a href="https://dx.doi.org/10.17504/protocols.io.ewov10xe2lr2/v1">https://dx.doi.org/10.17504/protocols.io.ewov10xe2lr2/v1</a>

#### Manuscript citation:

Li A, Liu S, Huang R, Ahn S, Lefkowitz RJ (2023) Loss of biased signaling at a G protein-coupled receptor in overexpressed systems. PLoS ONE 18(3): e0283477. doi: 10.1371/journal.pone.0283477

**License:** This is an open access protocol distributed under the terms of the **Creative Commons Attribution License**, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

Protocol status: Working

We use this protocol and it's working

Created: February 28, 2023

Last Modified: February 28, 2023

**Protocol Integer ID: 77745** 

Keywords: cell radioligand saturation, cell radioligand, receptor, coupled receptor, overexpressed system

Funders Acknowledgements:
Howard Hughes Medical Institute

#### **Abstract**

This protocol details an experimental procedure used to generate results described in the manuscript Li, A., Liu, S., Huang, R., Ahn, S., & Lefkowitz, R. J. (2023). Loss of biased signaling at a G protein-coupled receptor in overexpressed systems.

### **Troubleshooting**



### Day 1

- Grow cells on 150 mm dish to ~70% confluency
- 2 Wash twice with 10 mL PBS
- 3 Detach cells with 1 mL trypsin-EDTA + 11 mL media. Trypsinize for 5 minutes and check under microscope for complete detachment; tap dish if necessary
- 4 Count cells with hematocytometer and dilute collected cells to 164,000 cells/mL
- 5 Pipet 1 mL into each well of poly-D-lysine coated 12-well plate. Make duplicate plates for all conditions: one plate for binding and one for BCA assay

# Day 2

6 Add 250 mL of 5x doxycycline for all desired concentrations to cells 14 hours before assay

# Day 3 (BCA)

- 7 Aspirate media from BCA assay plates, wash each well with 1 mL cold PBS with MqCl2 and CaCl2, add 200 uL glycerol lysis buffer using repeater dispensing pipette. Shake in cold room
- 8 Fast cool centrifuge to 4 °C
- 9 Scrape each well with cell scraper, wiping with KimWipe and washing in cold PBS between wells. Pipet 150 uL lysate up and down 5x to wash wells and collect in Eppendorf tubes on ice
- 10 Centrifuge lysate for 10 minutes at 4 °C and 13,000 rpm
- 11 Collect 100 uL supernatant into new Eppendorf tubes on ice



12 Measure protein concentration using standard BCA assay protocol

## Day 3 (Binding)

- 13 Prepare saturating concentration of radioligand in media on ice. For example, 12.5 uL 3H-Angll in 5 mL media, or 4 uL 3H-olmesartan in 5.5 mL media. Prepare at least 5 mL per plate. Vortex briefly to mix
- 14 Take 1.5 mL media with radioligand per plate and add high concentration (orders of magnitude above radioligand concentration) of nonlabelled antagonist (e.g. 3 uL of 100 mM candesartan) for nonspecific binding measurement. Vortex briefly to mix
- 15 Dump media from plates into sink and invert plate onto paper towel to remove remaining liquid. Add 300 uL media with radioligand per well on ice. Use top two rows of plate for total binding duplicates and bottom row for nonspecific binding. Wait ≥ 1.5 hours on ice
- 16 Pre-chill 50 mL per plate PBS with MgCl2 and CaCl2 on ice and pre-warm 0.1% SDS + 0.5 N NaOH in 37 °C water bath
- 17 Remove radioligand media into dedicated radioactive liquid waste vial using P1000 on ice. Remove nonspecific binding media last
- 18 Wash wells with 1 mL cold PBS with MgCl2 and CaCl2 on ice and collect using P1000 into radioactive liquid waste vial. Wash 4 times total
- 19 Remove plates from ice and add 500 uL warm 0.1% SDS + 0.5 N NaOH per well. If there is precipitate in the solution, allow it to warm for longer. After adding, wait for  $\geq 15$ minutes
- 20 Collect samples from wells using P1000 and add to labeled scintillation vials. Add 300 uL of each radioligand media mix to two scintillation vials each for total counts. Add 5 mL scintillation cocktail (Lefko-Fluor) to each vial. Cap vials
- 21 Wipe vials with moist KimWipe and place in racks for scintillation counter. Place racks in counter and count using 3h\_cpm\_3\_min protocol
- 22 Perform second count using different protocol flag on next day

# **Analysis**



- 23 Use GraphPad radioactivity calculator to calculate specific activity of radioligand. The counter efficiency of the beta counter is approximately 65%; check the latest CPA results for an exact value
- 24 Copy binding counts into Excel spreadsheet and label
- 25 Transfer protein concentrations from BCA assay into spreadsheet
- 26 Use GraphPad radioactivity calculator and data in spreadsheet to calculate receptor concentration in fmol/mg
- 27 Use GraphPad radioactivity calculator and total counts from radioligand media to calculate radioligand concentration in nM