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BACKGROUND

Ubiquitin and Ubiquitin-Like Proteins

In cells, proteins are tagged for degradation by ubiquitin and sent to the proteasome. In contrast, covalent modification of cellular proteins by the ubiquitin-like modifier SUMO (small ubiquitin-like modifier) regulates various cellular processes, such as nuclear transport and signal transduction. The ubiquitin family of proteins fall into two classes: the first class, ubiquitin-like proteins (UBLs) function as modifiers in a manner analogous to that of ubiquitin. Examples of UBLs are SUMO, Nedd8 (also called Rub1), ISG15, Apg8, Apg12, and Fat10. Proteins of the second class include parkin, RAD23 and DSK2, are designated ubiquitin-domain proteins (UDPs). These proteins contain domains that are related to ubiquitin but are otherwise unrelated.

Ubiquitin Conjugation Machinery

The conjugation of Ub/UBLs to target proteins requires an orchestrated addition of Ub/UBLs to lysine residues in the target protein by E1 (ubiquitin activating enzyme), E2 (ubiquitin conjugating enzyme), and E3 (ubiquitin ligase) in an ATP dependent manner. The enzymes form an isopeptide bond between the carboxy-terminus of the UBL and the ϵ -amino group of the lysine residue of target proteins.

Deubiquitinating Enzymes

Ubiquitination is a reversible process in which deconjugation is performed in cells by deubiquitinating enzymes, otherwise known as isopeptidases. Isopeptidases are cysteine proteases that can be divided into five families, depending on sequence homology. These five families are the ubiquitin-C terminal hydrolases (UCH), the ubiquitin specific processing proteases (UBPs), the Machado-Joseph Disease domain proteases, the Otubain proteases (Otu), and JAMM domain proteases. The roles of isopeptidases include recycling of fused ubiquitin/UBL and processing pro-ubiquitin/UBL by cleavage to the mature form. Removal of ubiquitin or UBL moieties can affect cellular physiology in a number of ways, and several isopeptidases have been linked to pathologies such as cancer and cardiovascular disease.

ABOUT THE ASSAY

The Ub-CHOP2-Reporter Deubiquitination Assay consists of ubiquitin fused to a reporter enzyme, as well as a separate reagent substrate for the reporter enzyme. When fused to ubiquitin, the reporter is rendered catalytically inactive. Following cleavage of the Ub-reporter system by the isopeptidase, the free (and now active) reporter subsequently acts upon its substrate. Thus, in this coupled assay, the signal generated by cleavage of the reporter enzyme's substrate is a quantitative measure of isopeptidase activity.

BENEFITS

- 1. Superior to ubiquitin-AMC and FRET-based assays.
- 2. Rapid and robust readout for deubiquitylation activity within 45 minutes; signal to background ratio >20.
- **3.** Reporter substrates are non-radioactive.
- 4. Amenable to high throughput screening (HTS) and miniaturization.
- 5. Assay tests deconjugation of ubiquitin/UBL from a more physiologically relevant protein.
- Unlike ubiquitin-AMC, the CHOP2-Reporter system does not require excitation in the UV range (consequently reducing the incidence of false positives).

SUGGESTED USES

- **1.** Demonstration of novel isopeptidase activity.
- 2. High throughput screening (HTS) for antagonists or agonists of isopeptidase activity.

COMPONENTS

1. Ub-CHOP2-Reporter (Reporter System)

Size: 1 x 375 µl (4µM)

Buffer: 20 mM Tris (pH 8.0), 150 mM NaCl, 10% glycerol Storage: -80° C, avoid cycles of freezing and thawing

2. USP2 (Control Isopeptidase)

Size: 2 x 10 µl (2µM)

Buffer: 20 mM Tris (pH 8.0), 150 mM NaCl, 10% glycerol, 200μg/ml BSA, 2 mM β-

mercaptoethanol

Storage: -80° C, avoid cycles of freezing and thawing

3. Reporter Substrate 2

Size: 1 x 30 µl (25 µM)

Buffer: DMSO Storage: -80° C

Misc.: light sensitive

excitation/emission wavelengths are 485/531 nm

4. Control Fluorophore 2 (Please refer to Fluorophore Control section, Pg 11 for information concerning the use of this reagent for plate reader validation)

Size: 1 x 20µl (1mM)

Buffer: DMSO Storage: -80° C

ADDITIONAL ITEMS REQUIRED

1. Assay Buffer (for use in Step 1 of Protocol)

20 mM Tris-HCl (pH 8.0), 2 mM CaCl $_2$, 2 mM $\beta\text{-mercaptoethanol},~0.05\%$ CHAPS

Prepare fresh assay buffer on a daily basis

2. Fluorescence plate reader

Appropriate filters required for excitation and emission wavelengths

Excitation: 485 nm Emission: 531 nm

- 3. Black 96 well plate (or desired template)
- 4. 15 ml tube
- 5. 1.5 ml snap cap tubes

SOLUTIONS

USP2 (Control Isopeptidase) (For Use In Step 2 of Protocol)

- 1. Add 990 µl of assay buffer to one of the tubes labeled USP2.
- 2. Mix thoroughly by vortexing..
- 3. You are now ready to add 50 µl aliquots (20nM USP2) into the wells of a black 96 well plate.

Reporter Substrate Solution (for use in Step 4 of Protocol)

- 1. Perform this step in a timely manner as to minimize exposure of the reporter substrate 2 to light.
- Dilute Reporter Substrate 2 in a 1:10 ratio with DMSO: Add 1ml reporter substrate 2 to 9ml of DMSO.
- 3. Combine 3.2 μl of diluted reporter substrate 2, and 196.8 μl of assay buffer in a 1.5 ml snap cap
- 4. Mix thoroughly by vortexing .
- You are now ready to add 50 μl aliquots into the black 96 well plate.
- 6. Each 50µl aliquot contains 40nM of reporter substrate 2.

Ub-CHOP2-Reporter (Reporter System) and Reporter Substrate Solution (For Use In Step 5 of Protocol)

- 1. Perform this step in a timely manner as to minimize exposure of the reporter substrate 2 to light.
- Combine 150 μl of Ub-CHOP2-Reporter, 9.6 μl reporter substrate 2, and 5.86 ml of assay buffer in a 15 ml Falcon Tube.
- **3.** Mix throughy by vortexing.
- **4.** You are now ready to add 50 μl aliquots into the black 96 well plate.
- 5. Each 50 µl aliquot contains 100nM of Ub-CHOP2-Reporter and 40 nM of reporter substrate 2.

Test Isopeptidase (For Use In Step 3 of Protocol)

We suggest an initial dilution series of your test isopeptidase to optimize its concentration/activity. As a positive control, the kit includes the catalytic core domain of the deubiquitinating enzyme USP2, to be used at a concentration of 20nM in the final reaction. Make the dilutions of your test isopeptidase in assay buffer such that a final volume of 50 μ l is added to each well. A suggested protocol is listed below.

1. Label seven 1.5 ml snap cap tubes T1 through T7.

- 2. Place 200 μl of assay buffer in tubes T2 through T7.
- 3. Dilute the test isopeptidase to a concentration of 800nM in 400µl Assay Buffer in Tube T1
- **4.** Vortex Tube T1 and perform a 2-fold dilution by transferring 200 μl of solution from Tube T1 into Tube T2.
- 5. Perform another 2 fold dilution by taking 200 μl from Tube T2 and placing it into Tube T3.
- **6.** Repeat for Tubes T4 through T7.

tube	Concentration of test isopeptidase in 50 µl (nM)	Concentration of test isopeptidase in 100 µl (nM)
T1	800	400
T2	400	200
Т3	200	100
T4	100	50
T5	50	25
T6	25	12.5
T7	12.5	6.25

PROTOCOL

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Optimization of Your Test Isopeptidase, Suggested Protocol

- 1. Add 50 µl of Assay Buffer in triplicate to columns 1 and 2.
- 2. Add 50 μ l of USP2 (control isopeptidase) in triplicate to column 3.
- 3. Add 50 μl of Tubes T1-T7 of your test isopeptidase in triplicate to columns 4 through 10.
- 4. Add 50 μl of reporter substrate solution in triplicate to column 1.
- 5. Add 50 µl of reporter and reporter substrate solution in triplicate to columns 2 through 10.
- **6.** Below gives a representation of the 96 well plate layout.

'	2	3	4	3	0	'	0	9	10	- 11	12
50 μl Assay I	Buffer 50 µl Assay Buffe	er 50 µl USP2	50 μl Tube T1	50 µl Tube T2	50 µl Tube T3	50 µl Tube T4	50 µl Tube T5	50 µl Tube T6	50 µl Tube T7		
50 µl Repo		2- 50 μl Ub-CHOP2- Reporter &	50 μl Ub-CHOP2- Reporter &	50 μl Ub-CHOP2- Reporter &	50 µl Ub-CHOP2- Reporter &						
50 µl Assay l + 50 µl Repo Substrat	rter 50 µl Ub-CHOP2	+	50 µl Tube T1 + 50 µl Ub-CHOP2- Reporter &	50 µl Tube T2 + 50 µl Ub-CHOP2- Reporter &	50 µl Tube T3 + 50 µl Ub-CHOP2- Reporter &	50 µl Tube T4 + 50 µl Ub-CHOP2- Reporter &	50 µl Tube T5 + 50 µl Ub-CHOP2- Reporter &	50 µl Tube T6 + 50 µl Ub-CHOP2- Reporter &	50 µl Tube T7 + 50 µl Ub-CHOP2- Reporter &		
50 μl Assay l + 50 μl Repo Substrat	Buffer 50 µl Assay Buffer trter 50 µl Ub-CHOP2	+	50 µl Tube T1 50 µl Ub-CHOP2- Reporter &	50 µl Tube T2 + 50 µl Ub-CHOP2- Reporter &	50 µl Tube T3 + 50 µl Ub-CHOP2- Reporter &	50 µl Tube T4 50 µl Ub-CHOP2- Reporter &	50 µl Tube T5 + 50 µl Ub-CHOP2- Reporter &	50 µl Tube T6 50 µl Ub-CHOP2- Reporter &	50 µl Tube T7 + 50 µl Ub-CHOP2- Reporter &		

- 7. There are two options for fluorescence detection. First, the plate can be incubated at room temperature for 30 minutes, protected from light, and a single time point reading can be measured using the appropriate excitation and emission filters, 485 and 531 nm, respectively. Alternatively, an in-plate kinetic reading can be performed by measuring the change in fluorescence over time.
- **8.** Determine the mean plate blank value (column 1 in the representative example) and subtract from each data point to give a true representation of the change in fluorescence intensity. It is recommended that each experimental condition is measured in triplicate.

FLUOROPHORE CONTROL

Validation of fluorometric plate reader settings with fluorophore control (if desired)

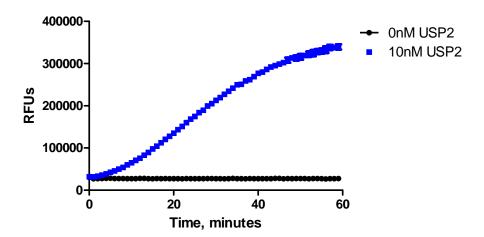
- Combine 2µI of Control Fluorophore 2 with 4mL of Assay Buffer. (Final concentration of 500nM for fluorophore solution)
- 2. Prepare a set of two-fold serial dilutions by combining 500µl of diluted fluorophore with 500µl of assay buffer. (This will result in concentrations of 250, 125, 62.5, 31.2, 15.6, 7.81, and 0nM).
- 3. Place 100µl of each dilution into the wells of 96-well plate in triplicate.
- Read plate on fluorescent plate reader with excitation and emission filters compatible with measurements of 485nm and 531nm respectively.

QUALITY CONTROL

Quality Control

In a 96 well plate, 50nM Ub-CHOP2-Reporter and 20nM reporter substrate were incubated in the presence and absence of 10nM USP2. Liberation of the fluorescent reporter substrate was monitored on a fluorescent plate reader using an excitation wavelength of 485 nm and an emission wavelength of 531 nm. Net RFU was determined by subtracting the blank (reporter substrate alone) from each data point. Results are shown in the figure below.

Representative experiment. Data = mean \pm Standard deviation



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