





Sep 06, 2022

Detection of Tau ubiquitylation

ltika Saha¹, F. Ulrich Hartl^{2,3}, Mark S. Hipp^{2,3}

¹Department of Cellular Biochemistry, Max Planck Institute of Biochemistry, Am Klopferspitz 18, 821 52 Martinsried, Germany;

²Department of Biomedical Sciences of Cells and Systems, University Medical Center Groningen, University of Groningen, Antonius Deusinglaan, 1, 9713 AV Groningen, The Netherlands;

³School of Medicine and Health Sciences, Carl von Ossietzky University Oldenburg, 26129 Oldenburg, Germany

1 Works for me Share

dx.doi.org/10.17504/protocols.io.3byl4je4rlo5/v1

Felix Kraus

ABSTRACT

This protocol describes the detection of ubiquitylated Tau from HEK293 cells stably expressing and propagating aggregates of Tau repeat domain fused to YFP (Sanders et al. Neuron, 2014; Saha et al, BioRxiv, 2022).

DOI

dx.doi.org/10.17504/protocols.io.3byl4je4rlo5/v1

EXTERNAL LINK

https://www.biorxiv.org/content/10.1101/2022.02.18.481043v1.full

PROTOCOL CITATION

Itika Saha, F. Ulrich Hartl, Mark S. Hipp 2022. Detection of Tau ubiquitylation. **protocols.io**

https://protocols.io/view/detection-of-tau-ubiquitylation-cf7vtrn6

K

MANUSCRIPT CITATION please remember to cite the following publication along with this protocol

The AAA+ chaperone VCP disaggregates Tau fibrils and generates aggregate seeds Itika Saha, Patricia Yuste-Checa, Miguel Da Silva Padilha, Qiang Guo, Roman Körner, Hauke Holthusen, Victoria A. Trinkaus, Irina Dudanova, Rubén Fernández-Busnadiego, Wolfgang Baumeister, David W. Sanders, Saurabh Gautam, Marc I. Diamond, F. Ulrich Hartl, Mark S. Hipp bioRxiv 2022.02.18.481043; doi: https://doi.org/10.1101/2022.02.18.481043

		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 are credited	,
		CREATED Sep 05, 2022	
		LAST MODIFIED	
		Sep 06, 2022	
		PROTOCOL INTEGER ID 69589	
1	Harvest cell	ls from 2 confluent wells of a 6 well plate.	
2	Lyse aggregate-containing cells by vortexing in cold RIPA buffer (Thermo) supplemented with protease inhibitor cocktail (Roche) and DNase. 20 mM N-ethylmalemide should be included in the lysis buffer to inhibit the activity of deubiquitinating enzymes.		
3	Briefly sonic	cate lysate and centrifuge at 2,000 x g for 5 min to remove cell debris.	5m
4	Collect supe	ernatant, determine protein concentration and normalize across samples.	
•	1111111		
5	Dilute 1 mg	protein in a total volume of 600 μL RIPA buffer.	
6	Add 50 µL a	nti-GFP bead slurry (µMACS GFP Isolation kit, Miltenyi Biotec) to diluted lysat	e.
7	Incubate for	r 1 h at 4 °C in a rotating wheel at 10 rpm.	1h

LICENSE

8	Before the end of 1 h, place μ -columns (Miltenyi Biotec) in the magnetic field of a μ MACS Separator (Miltenyi Biotec) and equilibrate columns by applying 250 μ L RIPA buffer. Allow complete flow-through.
9	Apply cell lysates and beads to μ-columns. Allow complete flow-through.
10	Wash columns 4 times with 1 mL 0.1% SDS/PBS.
11	Elute by applying 50 µL pre-heated (95 °C) 1x SDS sample buffer.
12	Analyze eluates by immunoblotting with antibodies against GFP or ubiquitin.