

Trypsin Sensitivity Assay to Study the Folding Status of Proteins

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[Abstract] This protocol aims to evaluate folding status of proteins, utilizing trypsin sensitivity. Unfolded/misfolded proteins are more susceptible to trypsin than folded proteins, because trypsin easily accesses and cleaves loosely folded parts of proteins. This method is especially useful to compare tightness of the folding among wild-type and mutant proteins. As trypsin generally cleaves a peptide bond at the carboxyl-terminal side of the amino acids lysine or arginine, this method can be used to analyze the folding status of different types of proteins such as integral membrane or soluble proteins (Ninagawa *et al.*, 2015) and is applicable to cell lysates of any species and tissues as well as to recombinant proteins. You can use this technique with regular molecular and cell biology equipment.

Materials and Reagents

1. 6 well dish (Corning, Falcon®, catalog number: 353046)
2. PVDF membrane (GE Healthcare, catalog number: 10600023)
3. DT40 cell line (DT40 is a B cell line derived from an avian leukosis virus induced bursal lymphoma in a white leghorn chicken) (ATCC, catalog number: CRL-2111)
4. Homo sapiens colon colorectal carcinoma cell line (HCT116) (ATCC, catalog number: CCL-247)
5. Dulbecco's modified Eagle's medium (DMEM) (NACALAI TESQUE, catalog number: 08458-45)
6. Fetal bovine serum (FBS) (Thermo Fisher Scientific, Gibco™, catalog number: 10270-106)
7. 100 U/ml penicillin and 100 µg/ml streptomycin (NACALAI TESQUE)
8. RPMI (NACALAI TESQUE, catalog number: 30263-95)
9. Chicken serum (Thermo Fisher Scientific, Gibco™, catalog number: 16110-082)
10. Opti-mem (Thermo Fisher Scientific, Gibco™, catalog number: 31985-070)
11. Lipofectamine 2000 (Thermo Fisher Scientific, Invitrogen™, catalog number: 11668019)
12. 2.5 g/L trypsin (NACALAI TESQUE, catalog number: 32777-44)
13. Protease inhibitor cocktail (100x) (NACALAI TESQUE, catalog number: 25955-11) (for inhibition of various proteases' activity such as trypsin, in ddH₂O; stored at -20 °C)
14. NaCl (Wako Pure Chemical Industries, catalog number: 191-01665) (for DT40)
15. Na₂HPO₄
16. KCl
17. KH₂PO₄
18. Tris/HCl (pH 8.0) (Sigma-Aldrich, catalog number: T6791) (Stored at room temperature)
19. Glycerol

20. Bromophenol blue (BPB)
21. Sodium dodecyl sulfate
22. Nonidet P-40 (NACALAI TESQUE, catalog number: 23640-94) (for HCT116)
23. 20 mM carbobenzoxy-valyl-alanyl-aspartyl-[O-methyl]-fluoromethylketone (Z-VAD-fmk) (Promega, catalog number: G7231) (for inhibition of PNGase activity)
24. 10 mM Z-Leu-Leu-Leu-CHO (MG132) (PEPTIDE INSTITUTE, catalog number: 3175-v) (in DMSO; inhibition of proteasomal activity, stored at -20 °C)
25. 1 M dithiothreitol (DTT) (Wako Pure Chemical Industries, catalog number: 041-08976) (in water; for reduction of proteins)
26. Anti- β -actin antibody (Wako Pure Chemical Industries, catalog number: 017-24573)
27. Anti-myc antibody (MEDICAL & BIOLOGICAL LABORATORIES, catalog number: 562)
28. Phosphate buffered saline (PBS) (see Recipes)
29. 2x sodium dodecyl sulfate (SDS) sample buffer (pH 6.8) (see Recipes)
30. Buffer A (see Recipes)
31. Buffer B (see Recipes)

Equipment

1. High speed refrigerated micro centrifuge (Tomy, model: MX-301)
2. mPAGE (ATTO, model: AE-6530) (Using hand-made 10% gel)
3. Transfer equipment (ATTO, model: WSE-4020)
4. Heat block (TAITEC, model: DTU-1BN)
5. Micro porator (Digital Bio, model: MP-100)

Software

1. ImageJ

Procedure

1. Culture adherent HCT116 cells in Dulbecco's modified Eagle's medium (glucose 4.5 g/L) supplemented with 10% fetal bovine serum and antibiotics (100 U/ml penicillin and 100 μ g/ml streptomycin) at 37 °C in a humidified 5% CO₂/95% air atmosphere. Culture suspended DT40 cells at a density of 1×10^5 - 1×10^6 cells per ml in RPMI1640 medium supplemented with 10% fetal bovine serum, 1% chicken serum and antibiotics (100 U/ml penicillin and 100 μ g/ml streptomycin) at 39.5 °C in a humidified 5% CO₂/95% air atmosphere.
2. Transfect 8.0×10^5 HCT116 cells (40-60% confluent in 6-well plate) using 1 μ g of plasmid DNA, 300 μ l of Opti-mem and 10 μ l of Lipofectamine 2000 (Invitrogen) for one well according to the manufacturers' instructions (After the transfection medium is not replaced). To obtain 1.05×10^7

cells transfected cells, electroporate three times 3.5×10^6 DT40 cells with 8 μ g plasmid DNA with two pulses at 1,500 V for 15 msec according to the manufacturer's instructions.

3. Collect approximately 2.0×10^6 HCT116 cells or 4.0×10^6 DT40 cells for further analysis 24 h (HCT116 cell) or 16 h (DT40 cells) after transfection. For this the cells are washed with 1 ml PBS three times, scraped off and centrifuged at $3,000 \times g$ for 2 min at 4 °C. Discard the supernatants of samples and suspend cell pellets in 400 μ l (for HCT116 cells) or 250 μ l (for DT40 cells) buffer A and incubate for 20 min on ice to lyse the cells. We used smaller amount of buffer A for DT40 cells, because DT40 cells do not contain as much proteins as HCT116 cells. After this step, samples from HCT116 cells and DT40 cells are treated in the same manner.

Note: Important, the buffer A must NOT contain protease inhibitor cocktail as it would inhibit trypsin activity.

4. Clarify cell lysates by centrifugation at $17,800 \times g$ for 10 min at 4 °C and transfer the supernatants to new tubes. Pellets are discarded.
5. To allow trypsin processing of misfolded proteins, incubate 50 μ l aliquots (in case of protein concentrations of 1 mg/ml) of the cleared lysates with 1 μ l trypsin solution (Test different concentrations from 1 μ g/ml to 100 μ g/ml) for 15 min at 4 °C. To stop the trypsin reaction, mix samples with 40-50 μ l of buffer B supplemented with 10x protease inhibitor cocktail (We usually use high inhibitor concentrations: 5x final) and incubate at 100 °C for 5 min. Important, every 3 min four samples could be handled for equal incubation time with trypsin.
6. Separate 10 μ l of the samples (approximately 5 μ g) by SDS-PAGE followed by immunoblotting using PVDF membrane and probing with an antibody against the protein of interest. Percentage of SDS-PAGE depends on the molecular weight of the protein of interest. Band intensities can be analyzed and compared to each other by ImageJ or similar programs (Figure 1).

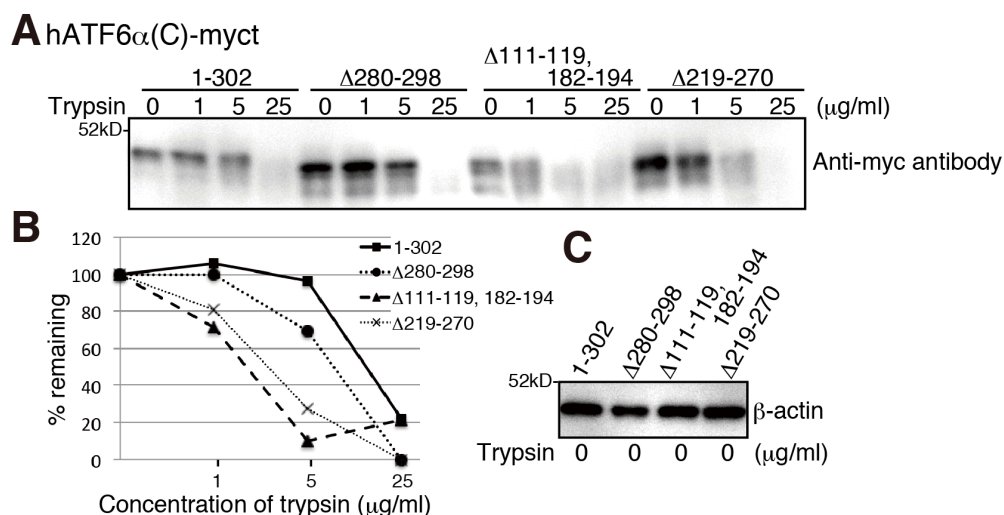


Figure 1. Severely misfolded mutants were more sensitive to trypsin. A. Chicken EDEM1/2/3 triple KO cells (gEDEM TKO) were described previously (Ninagawa *et al.*, 2015). The indicated myc-tagged hATF6 α (C) variants were transiently express in gEDEM-TKO cells

and subjected to trypsin sensitivity assay. After the indicated time points the trypsin reaction was stopped and samples analyzed by a 12% SDS-PAGE followed by immunoblotting with antibodies against the myc-tag. The data shown represents a single representative experiment out of three repeats. Less folded proteins are more sensitive to trypsin, so the hATF6 α (C)-myc mutants, Δ 111-119, 182-194 mutant and Δ 219-270 mutant were cleaved by lower trypsin concentrations. B. Quantification of band intensities shown in (A) by ImageJ and calculation of the percentage of remaining protein. C. Equal loading of cell lysates shown by immunoblotting of the lysates and probing with an antibody against β -actin.

Notes

1. Incubation time with trypsin and concentration of trypsin should be optimized for proteins of interests.
2. The lot and freshness of trypsin can affect the result and can change the outcome between independent experiments (We use trypsin stored at 4 °C). Anyway, you can compare the tightness of the folding among proteins of interests.
3. Dilute trypsin in PBS to obtain the expected concentration.
4. We used gEDEM TKO cells. This method is applicable for a wide variety of cells lines and recombinant proteins.
5. This method can be applied to recombinant proteins. In this case, you might reduce concentration of trypsin.
6. CPY and CPY* (Izawa *et al.*, 2012), and GFP and GFP variants (Xu *et al.*, 2013) were other examples used in this assay. We can provide plasmids to express hATF6 α (C)-myc 1-302 and Δ 280-298 for positive controls (Folded), and Δ 111-119, 182-194 and Δ 219-270 for negative controls (Less folded).

Recipes

1. Phosphate buffered saline (PBS)
137 mM NaCl
8.1 mM Na₂HPO₄
2.68 mM KCl
1.47 mM KH₂PO₄
2. 2x sodium dodecyl sulfate (SDS) sample buffer (pH 6.8)
100 mM Tris/HCl (pH 6.8)
20% glycerol
0.2% bromophenol blue (BPB)
4% sodium dodecyl sulfate
3. Buffer A
50 mM Tris/HCl (pH 8.0)

1% NP-40
150 mM NaCl
2 μ M Z-VAD-fmk (Add before use)
20 μ M MG132 (Add before use)
Stored at 4 °C.

4. Buffer B

1x SDS sample buffer
100 mM dithiothreitol (Add before use)
10x protease inhibitor cocktail (Add before use, and 10 times diluted from original protease inhibitor cocktail)
Stored at room temperature.

Acknowledgments

This protocol was adapted from and used in Ninagawa *et al.* (2015), Izawa *et al.* (2012) and Xu *et al.* (2013).

References

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3. Xu, C., Wang, S., Thibault, G. and Ng, D. T. (2013). [Futile protein folding cycles in the ER are terminated by the unfolded protein O-mannosylation pathway](#). *Science* 340(6135): 978-981.