

Polysome Preparation, RNA Isolation and Analysis

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[Abstract] During mRNA translation, 40S and 60S ribosomal subunits bind to target mRNA forming into an 80S complex (monosome). This ribosome moves along the mRNA during translational elongation to facilitate tRNA reading codon, where translation is activated and many monosome can bind the same mRNA simultaneously, which forms polysomes. Polysomes can be size-fractionated by sucrose density gradient centrifugation. The more specific mRNA in polysomes implies more active translational status of the mRNA.

Materials and Reagents

1. Cells (Neuroblastoma cell line SKN-SH)
2. Protease inhibitors (Sigma-Aldrich, catalog number: P8340-5ML)
3. DTT (Sigma-Aldrich, catalog number: 43815)
4. Cycloheximide (CHX) (Sigma-Aldrich, catalog number: C7698)
5. Heparin (Sigma-Aldrich, catalog number: H3149)
6. Sucrose (Sigma-Aldrich, catalog number: S1888)
7. Phenol/chloroform/isoamyl alcohol (Life Technologies, Invitrogen™, catalog number: 15593-031)
8. Sodium acetate (Thermo Fisher Scientific, catalog number: S209-500)
9. 1x PBS
10. 1x Trypsin-EDTA, 0.05% Trypsin/0.53 mM EDTA (Cellgro, catalog number: 25-052-CV)
11. RPMI-1640 (Hyclone, catalog number: SH30096.01)
12. Tris-Base (Thermo Fisher Scientific, catalog number: BP-152-1)
13. KCl (Thermo Fisher Scientific, catalog number: BP-366-500)
14. MgCl₂ (Sigma-Aldrich, catalog number: M-2393)
15. Triton X-100 (Sigma-Aldrich, catalog number: T8787-250ML)
16. DNAase/RNAase free Ethanol (Sigma-Aldrich, catalog number: E7023-500ML)
17. DNAase/RNAase free water (BioExpress, catalog number: UPW-1000)
18. Ultracentrifuge tubes (14 x 89 mm) (Beckman Coulter, catalog number: 344059)
19. Polysome extraction buffer (PEB) (see Recipes)
20. Sucrose solutions (see Recipes)

Equipment

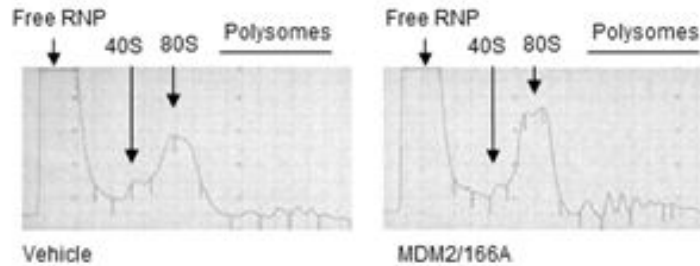
1. BR-188 Density gradient fractionation system (Brandel)
2. Beckman optima L-70 ultracentrifuge (Beckman)
3. 7500 Real-time PCR system (Applied Biosystems)
4. Boekel scientific orbitron rotator I, 115 V (Boekel Scientific)

Procedure

1. Prepare neuroblastoma cell line SKN-SH cells $2.5-5 \times 10^7$ /group, normally 2 to 3 150 cm² flasks with 50 ml of RPMI-1640 medium.
2. Before harvest, cells are incubated in complete RPMI-1640 medium (RPMI-1640 medium with FBS, Pen/Strep and Glutamine) containing a final concentration of 100 µg/ml CHX for 10 min at 37 °C.
3. Discard medium andrinse cells twice with cold 10 ml 1x PBS containing 100 µg/ml CHX.
4. Harvest cells, briefly add 7 ml of trypsin into each flask, incubate at 37 °C for 3 min, then collect cells with 10 ml PBS containing 100 µg/ml CHX (this step is not needed for suspension cells).
5. After centrifugation, resuspend cells in 1 ml of 1x PBS, then transfer them into 1.7 ml micro centrifuge tube.
6. Lyse cells with 1 ml PEB Buffer containing 1% Triton-X100, vortex 15 sec and keep on ice for 30 min.
7. Centrifuge at 14,000 rpm for 30 min at 4 °C and collect supernatants that are ready to be loaded on sucrose gradients of 10%-50% consistency.
8. Make sucrose solutions according to the recipe step.
9. Make 10 ml gradient by adding 2.0 ml of each solution (50% on bottom, 10% on top) and load into ultracentrifuge tube.
10. Add 1 ml of cell extract to the top of each gradient. Cover each tube with parafilm, put it in a SW41 rotor andspin samples at 38,000 rpm for 120 min at 4 °C.
11. Collect 13 fractions from the top to the bottom of the tube using BR-188 density gradient fractionation system.
12. Machine setting options: fraction collection time: 1 min and 15 sec/fraction; pump speed, 0.75 ml/min; chart speed, 60 cm/h).
13. Keep all fractions at -80 °C, it's good for extraction of RNA in 1 month.
14. For RNA extraction, thaw all samples on ice.
15. Pick 0.5 ml of each sample into a new 2 ml-microcentrifuge tube, keep on ice.
16. Mix the water saturation Phenol/chloroform/isoamyl alcohol (25:24:1) completely.

17. Add 500 μ l of the mixed Phenol/chloroform/isoamyl alcohol into each tube, when pipetting keep shaking with hand to avoid water separating out.
18. Vortex for 30 sec and then keep rotating at 4 $^{\circ}$ C for 5 min.
19. Centrifuge at 14,000 rpm for 10 min at 4 $^{\circ}$ C, at the same time prepare new 2 ml-microcentrifuge tube for each sample, add 3 M Sodium Acetate (pH 5.2) 50 μ l/tube.

A, The 254 nm traces obtained during collection of fractions.



B, Relative distributions of mRNA.

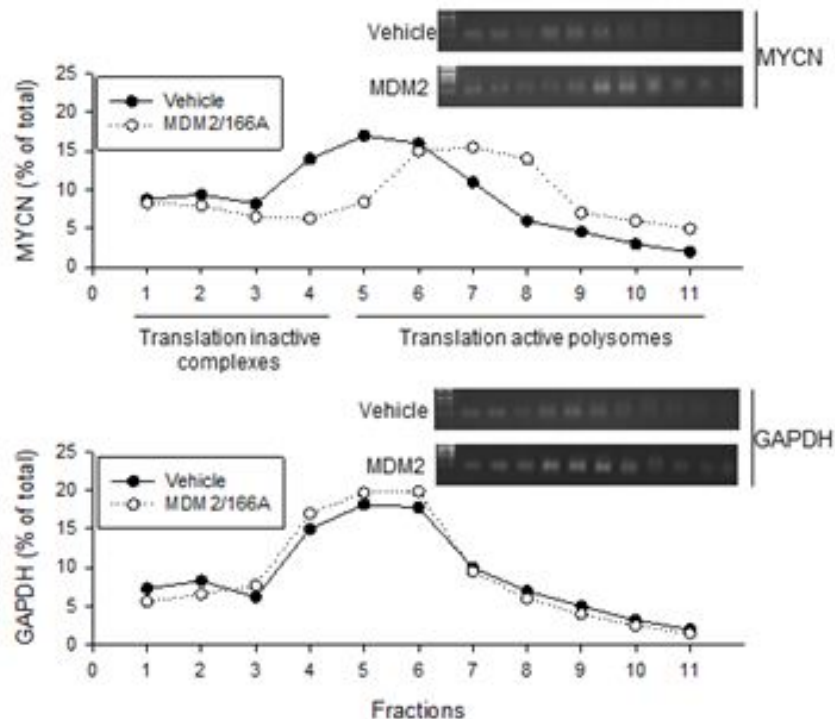


Figure 1. MDM2 regulates MYCN translation. A. Representative polysomal profiles from control vector- and MDM2/166A-transfected SK-N-SH cells. The 254-nm traces obtained during collection of fractions are shown. B. Relative distributions of MYCN and GAPDH mRNA in SK-N-SH cells transfected either with vehicle or MDM2/166A.

20. Very carefully pipet out all the aqueous phase and put it into the prepared fresh 2 ml-microcentrifuge tube.
21. Then add 1.5 ml (3 volumes) of 100% ethanol, vortex 30 sec.
22. Keep the samples at -80 °C for 3 h.
23. Centrifuge at 14,000 rpm for 30 min at 4 °C.
24. Wash pellet with cold 70% RNase free ethanol by rotating for 5 min.
25. Centrifuge pellet at 14,000 rpm for 5 min at 4 °C.
26. Discard supernatants and remain the pellet in the tubes.
27. Put tubes on ice to let all the liquid evaporated.
28. Add 30-50 µl of DNase/RNase free H₂O and subjected to quantitated RT-PCR.

Recipes

1. Polysome extraction buffer (PEB)
(Triton-X100 Free)
20 mM Tris-HCl (pH 7.5)
50 mM KCl
10 mM MgCl₂
1 mM DTT
100 µg/ml CHX
200µg/ml Heparin
2. Preparation of 50%, 40%, 30%, 20%, 10% sucrose solutions as following
Make 50% Sucrose solution (25 g sucrose + PEB without Triton-X100 to 50 ml)
Dilute 50% sucrose solution into the solutions of 40%, 30%, 20% and 10% in PEB buffer

Acknowledgements

This work was supported by the National Institutes of Health (R01 CA123490 and R01CA143107 to MZ) and CURE (MZ and LG).

References

1. Gu, L., Zhang, H., He, J., Li, J., Huang, M. and Zhou, M. (2012). [MDM2 regulates MYCN mRNA stabilization and translation in human neuroblastoma cells.](#) *Oncogene* 31(11): 1342-1353.