

Materials List for:

Visualizing RNA Localization in *Xenopus* Oocytes

James A. Gagnon¹, Kimberly L. Mowry¹

¹Department of Molecular Biology, Cell Biology, and Biochemistry, Brown University

Correspondence to: Kimberly L. Mowry at Kimberly_Mowry@brown.edu

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Materials

10X Tx buffer
<ul style="list-style-type: none"> • 60 mM MgCl₂ • 400 mM Tris-HCl (pH 7.5) • 20 mM spermidine-HCl
20x cap/NTP mix
<ul style="list-style-type: none"> • 10 mM CTP • 10 mM ATP • 9 mM UTP • 2 mM GTP • 20 mM G(ppp)G Cap Analog (New England Biolabs)
G-50 column
<ul style="list-style-type: none"> • Hydrate 5 g Sephadex G-50 beads (Sigma Aldrich) in 100 ml deionized H₂O. DEPC-treat for 30 min. and autoclave. Store incomplete stock at room temperature. Before use, add the following RNase-free solutions: • 0.5 ml 0.2 M EDTA • 1 ml 1 M Tris pH 8.0 • 0.5 ml 20% SDS • Store complete G-50 solution at 4 ° C. • Remove and discard the plunger from a 3 ml syringe (BD Biosciences) and place the barrel of the syringe into a 15 ml conical tube (Corning). Plug the syringe with a small amount of glass wool (a plug about half the size of a penny). • Swirl complete G-50 solution to resuspend beads. • Add 2 ml G-50 solution to the empty column. • Spin for 1 minute at 1,000 x g in benchtop centrifuge. • Add 200 µl DEPC-treated deionized H₂O to each column. Spin. • Repeat wash twice more for a total of three washes. • Remove syringe barrel to a fresh 15 ml conical tube.
Collagenase solution
<ul style="list-style-type: none"> • 75 mg collagenase from <i>Clostridium histolyticum</i> (Sigma Aldrich) • 25 ml 0.1 M KPO₃⁺ (pH 7.4)
MBSH buffer
<ul style="list-style-type: none"> • 88 mM NaCl • 1 mM KCl • 2.4 mM NaHCO₃ • 0.82 mM MgSO₄ X 7H₂O • 0.33 mM Ca(NO₃)₂ X 4H₂O • 0.41 mM CaCl₂ X 6H₂O • 10 mM HEPES (pH 7.6)
Oocyte Culture Medium
<ul style="list-style-type: none"> • 50% L15 medium • 15 mM HEPES (pH 7.6) • 1 mg/ml insulin

- 100 mg/ml gentamicin
- 50 U/ml nystatin
- 50 U/ml penicillin
- 50 mg/ml streptomycin

MEMFA solution

- 0.1 M MOPS (pH 7.4)
- 2 mM EGTA
- 1 mM MgSO₄
- 3.7% formaldehyde

Computing RNA yield

- Determine CPM in "input" and "incorporated" samples using a standard scintillation counter.
- $\text{incorporation} = (\text{"incorporated"}) / (10 \times \text{"input"})$
- Typical incorporation values range between ~0.03 and 0.10.
- Maximum theoretical yields for different polymerases: T7, T3, SP6 - 2.64 μg
- $\text{Reaction yield in } \mu\text{g} = (\text{maximum yield of polymerase used}) \times (\text{incorporation})$
- $\text{Dilute RNA to } 50 \text{ nM} = (\mu\text{g RNA}) / 320 / (\text{length of RNA in bases}) / (5 \times 10^{-8})$
- The reaction usually yields ~50-100 μl of RNA at 50 nM.