

mRNA Display Protocol

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1. Obtain DNA library
2. Using NEB T7 Quick High Yield RNA Synthesis Kit
 - a. For transcripts > 0.3 kb in a 20 uL reaction volume:
 - i. Nuclease-free water – X uL
 - ii. NTP Buffer Mix – 10 uL (10 mM each NTP final)
 - iii. Template DNA – X uL (1 ug)
 - iv. T7 RNA Polymerase Mix – 2 uL
 - b. Mix and incubate at 37C for 2 hrs. Recommended to use a dry air incubator or thermocycler to prevent evaporation
 - c. Optional: DNase treatment to remove DNA template
 - i. Add 30 uL nuclease-free water to 20 uL RNA synthesis reaction.
 - ii. Add 2 uL of DNase I (RNase-free)
 - iii. Mix and incubate at 37C for 15 minutes
3. Purification of synthesized RNA
 - a. Using Zymoclean RNA Clean and Concentrator -5 Kit, follow provided instructions. Ensure that sample has at least 50 uL volume. Resuspend sample in a low, but manageable volume in order to maximize the concentration of RNA.
 - b. Take an OD measurement of the RNA. The RNA concentration in uM or pmoles/uL is approximately the absorbance at 260 nm (A-260) divided by 3.2
4. Ligation of splint/linker to RNA using NEB T4 DNA Ligase Kit
 - a. Will need roughly 100-200 uL RXN to have sufficient amounts by the end of mRNA display. This can be broken into smaller volume RXNs.
 - b. Prepare preliminary mix
 - i. 8-10 uM RNA; 11 uM splint; 12 uM puromycin-linker; ddH2O as necessary
 1. Lower amounts (~8 uM) of RNA seemed to provide the greatest efficiency
 - c. Heat at 65C for 1-3 minutes to remove RNA tertiary structures. DO NOT ADD T4 DNA LIGASE OR LIGASE BUFFER YET!
 - i. Allow to cool for 1-3 minutes at RT
 - d. Add T4 DNA Ligase Buffer and Ligase as appropriate
 - e. Incubate at room temperature for 30-60 minutes
5. Analysis of ligated transcripts by gel electrophoresis
 - a. Prepare 2X RNA Loading Buffer

- i. 95% Formamide; 0.025% SDS; 0.025% Bromophenol Blue; 0.025% Xylene Cyanol; 0.5 mM EDTA
 - ii. Mix well and dissolve all SDS
 - b. Run sample on agarose gel with ethidium bromide.
 - i. Run at a lower voltage than with DNA
6. Gel Purification
 - a. Extract appropriate band according to expected RNA size
 - b. Several methods of purification are available such as LiCl/EtOH precipitation, PCR-cleanup column, PH:CL extraction/EtOH precipitation, or gel purification
 - i. When high purity RNA transcript is desired, gel purification is recommended
7. Affinity purification using oligo dT cellulose to remove unligated transcripts
 - a. Prepare oligo dT Binding Buffer
 - i. 50 mM Tris, pH 8; 1 M NaCl; 10 mM EDTA; 0.1% Triton x-100
 - b. Use 2 mg oligo dT cellulose per 100 uL product (~0.4 uM RNA)
 - c. Dilute sample 10X in Binding Buffer
 - d. Transfer to spin-x filter.
 - e. Wash 3X with Binding Buffer at RT
 - f. Wash 1X with ice cold TBS with 0.05% Tween
 - g. Elute with RT or warm 5 mM Tris, 3 x 25 uL
8. Translation
 - a. 50-100 uL Translation volume, using Ambion Retic Lysate for 0.4 uM ligated RNA template. Follow kit instructions
 - b. Incubate for 1 hour at 30C
 - c. Add salt to facilitate fusion formation. Per 50 uL RXN:
 - i. 12 uL 3 M KCl; 4 uL 1 M MgCl₂
 - ii. Incubate for 30 minutes at RT
9. FLAG Pre-selection
 - a. Wash 10 uL M2 anti-FLAG agarose beads with PBS with 0.02% Tween-20
 - b. Add 10 uL 0.5 M EDTA to translation sample. Dilute to 500 uL with TBS, 0.02% Tween-20 and add to 10 uL M2 agarose beads
 - i. Can do binding in spin-x cup
 - c. Bind 1 hour at 4C, then wash 3X with cold TBS – tween
 - d. Wash twice with cold 1XRT buffer (SSII First Strand Buffer)
 - e. Elute off beads by incubating with 100 uL of 1XRT buffer with 0.2 mg/ml 3XFLAG peptide for 30 minutes at RT
10. Reverse Transcription
 - a. 100 uL Reaction volume

- i. 2 μ L 25 mM dNTP mix; 150 pmoles [1.5 μ L of 100 μ M] of reverse primer; 1ng-5 μ g RNA
 - ii. Heat sample for 2 minutes at 42C, then add 1 μ L of superscript II enzyme
 - iii. Incubate for one hour at 42C
- b. Test on a gel and gel purify