

Probe Preparation

Cut 20-40 µg of plasmid DNA

I. Digest Rxns:

DNA	10-16µl
RE (20U/µl)	1µl
10X Buffer	2µl
20X BSA	1µl
dH2O	To volume
Total volume	20µl

-Incubate O/N at 37°C

Phenol/Chloroform Extraction

1. Extract DNA from samples by adding an equal volume of phenol/chloroform at 25:24 (bring samples to 450µl first to minimize DNA loss)
2. Vortex for 10 sec and spin for 5' at 12400xg (4C)
3. Transfer aqueous phase to a clean tube, add 450µl of chloroform, vortex for 10 seconds, and spin for 5' at 12400xg (4C).
4. Transfer the aqueous phase to a clean tube, add 1/10 volume NaOAC pH6.0 and 2.5x volume 100% EtOH
5. Invert several times and put in -80°C for 30'
6. Spin at 4°C at 12,400xg for 15'
7. Remove 100% EtOH and add 500µl of 70% EtOH
8. Spin for 10' at 15,000xg at 4°C
9. Remove EtOH, dry the pellet, and resuspend in 12µl DEPC dH₂O

OR

I. PCR:

Sigma H ₂ O	38.25µl
10X Titanium buffer	5µl
dNTP(10mM)	1µl
Plasmid DNA (1-15ng)	0.5µl
T7(10µM)	2.5µl
T3 or Sp6(10µM)	2.5µl
50X Titanium Taq	0.25µl

cycling parameters: 95°C for 1min, 35 cycles: 95°C for 30sec, 55°C for 30sec , 68°C for

1min/Kb

II. Transcription:

10X buffer	5 μ l
T7, T3, or Sp6(20U/ μ l)	4.5 μ l
Rnasin (40U/ μ l)	1 μ l
Dig mix	5 μ l
1-2.5 μ g linearized DNA	2-10 μ l
DEPC dH ₂ O	volume
Total volume	50μl

OR

10X buffer	5 μ l
T7, T3, or Sp6(20U/ μ l)	4.5 μ l
Rnasin (40U/ μ l)	1 μ l
Dig mix	5 μ l
PCR DNA	10 μ l
DEPC dH ₂ O	24.5 μ l
Total volume	50μl

1. Incubate samples at 37°C for 2 or more hours
2. Add 2 μ l (20U) DNase1 to each reaction and incubate at 37°C for an additional 10min
3. Add 52 μ l of stop buffer to each reaction
4. To remove excess Dig-labeled nucleotides transfer each sample to a Sephadex G50 column:
 - a. Add 800 μ l of DEPC dH₂O to the column after tapping it to ensure that the gel powder is on the bottom (also make sure to leave the end stopper on)
 - b. Replace the cap, vortex briefly, and incubate at RT for 30min
 - c. After incubating invert the column, tap sharply, and return to an upright position. This is to remove air bubbles
 - d. Remove the stopper and allow column fluid to flow into the catch container (~200-250 μ l). It may be necessary to spin at 750xg for a few seconds
 - e. Discard the flow-through and spin the columns at 750xg for 2min at 4°C.
 - f. Remove the catch tube and put the column into a clean tube, add the entire transcription reaction directly onto the top of the gel in a drop-wise fashion. Do this quickly so that the gels do not dry out
 - g. Spin the tubes at 750xg for 2min at 4°C being sure that the tubes are oriented in the same way as the first spin

5. After spinning EtOH precipitate the probes:
 - a. add 1/10 volume NaOAc pH6.0 and 2.5x volume 100% EtOH
 - b. Invert several times and put in -80°C for 15min
 - c. Spin at 4°C at 12,400xg for 15min
 - d. Remove 100% EtOH and add 500µl of 70% EtOH
 - e. Spin for 5min at 20,000xg at 4°C
 - f. Remove EtOH, dry the pellet, re-suspend in 50µl DEPC dH₂O
6. OD the probes and then dilute them to 1µg/ml in hybridization buffer
7. Remove 1µl of each sample to run on a 1% gel to check for correct probe MW.