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**Protocol status:** Working We use this protocol and it's working.

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## Electrophoretic mobility shift assay (EMSA)

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#### **ABSTRACT**

This protocol details an experimental procedure that can be followed to assess the DNA binding activity of a purified protein sample (recombinant yeast Cbf1 used here as example) to different sequences. DNA probes containing the desired sequences are formed by annealing complementary ssDNA oligos. These oligos are ordered from IDT and with 3' 6-FAM tags for imaging.

- 1 Create storage stocks by resuspending ssDNA oligos to 100uM (100pmol/ul) in 1x IDTE buffer (10 mM Tris, 0.1 mM EDTA, pH 8.0).
- Create 10uM working stocks from the 100uM storage stocks. For 100uL of working stock:  $10uL \, ssDNA(100uM) + 90uL \, IDT \, duplex \, buffer = 100uL \, ssDNA(10uM)$
- Anneal labeled 1 units ssDNA probes to 1/1.2 units unlabeled probes in 100uL reactions. Using Thermocycler, heat reaction mixes to 95°C and incubate for ≥2min. Cool to 25°C/RT over 45min. Keep duplexed probes in the dark until they are needed.

Reagent	1:1 (uL)	1:1.2 (uL)
Labeled	10	10
Unlabeled	10	12
Duplex buffer	80	78

## **Protein binding reaction**

- 4 Place running tank on ice and pre-run TBE native gel (8-20%; stored in backup -20°C) at 200V for 60min or until current stabilizes.
- 5 Prepare 100uL of 5x protein (Cbf1) binding buffer:

Reagent	Volume (uL)	
10x PBS	10	
0.5M EDTA	2	
0.5M DTT	2	
20mg/mL BSA	2.5	
nfH20	83.5	

**6** Prepare individual binding reactions (20uL/reaction):

Reagent	Volume (uL)

Reagent	Volume (uL)
5x Cbf1 binding buffer	4
1ug/uL poly dl:dC	1
Purified Cbf1	1
Duplexed probe*	2
nfH2O	12

#### Note

\*IMPORTANT: Incubate the binding reactions for 10min at RT <u>prior</u> to adding the duplexed DNA probe. This sequesters general DNA-binding proteins onto competitor DNA (poly dl:dC), minimizing non-specific binding to duplexed probe.

7 Incubate at RT in the dark for 50min following the addition of duplexed probe.

### **Electrophoresis**

- 8 Add 4uL of orange G loading dye to each binding reaction (final volume = 24uL).
- 9 Load 10uL of each binding reaction into well on pre-run TBE gel. Run at 200V for ~30min. **Keep on ice** and in the dark.
- 10 Image gel using BioRad Chemidoc system in CIEMAS (same machine used for SDS-PAGE).

### Note

For publication quality images, image with FAM/Cy2 filter using the Typhoon imager in SANDS 250. Contact Dr. Seok-Yong Lee's lab for use.