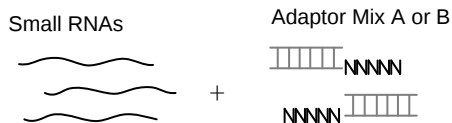


Workflow

Starting material

- Small RNA, purified using the flashPAGE™ Fractionator or PAGE, **or**
- Total RNA that contains the small RNA fraction **and**
- Adaptor Mix A: for sequencing the 5' ends of small RNAs, **or**
- Adaptor Mix B: for sequencing the reverse complement of the RNA



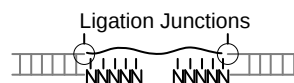
Hybridization and ligation to adaptor mix (8 hr)

On ice, mix RNA, Adaptor Mix, and Hybridization Solution.

Incubate the sample at 65°C for 10 min, then at 16°C for 5 min.

Add ligation reagents to each sample.

Incubate the sample at 16 °C for >8 hr in a thermal *cycler*

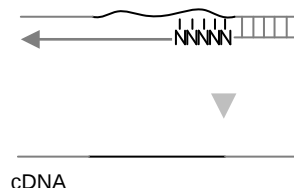


Reverse transcription and RNase H digestion (1 hr)

On ice, add 20 µ L RT Master Mix to each sample.

Incubate at 42°C for 30min

Add 1 µ L RNase H to 10 µ L cDNA and incubate at 37 °C for 30 min



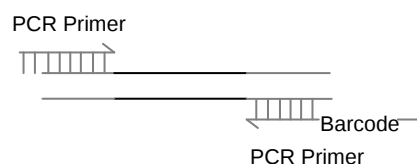
cDNA library amplification (1 to 1.5 hr)

Dispense PCR Master Mix into wells of a PCR plate or tubes.

Add RNase H-treated cDNA to each reaction mix, then run the PCR

Run 5-10 µ L PCR product on a native 6% polyacrylamide gel.

Evaluate the PCR products.



Amplified library cleanup and size selection by PAGE(3 to 4 hr)



SOLiD™ Sample Preparation and Sequencing

SOLiD™ System: start at the “Templated Bead Preparation” section of the SOLiD™ System instructions(emulsion PCR)