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🌐 Brad-seq mRNA (for Shotgun or DGE) V.2

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protocol .



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Brad-seq mRNA

Yin-Tse Huang 2022. Brad-seq mRNA (for Shotgun or DGE). **protocols.io**
<https://protocols.io/view/brad-seq-mrna-for-shotgun-or-dge-bqznm5e>
Yin-Tse Huang



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- Lysis/binding buffer (LBB)

| A | B | C |
|------------------|-------------|-------------------|
| Component | Stock | Volumes for 50 ml |
| 100 mM Tris-HCl | 1 M pH 8 | 5 ml |
| 1M LiCl | 8 M | 6.25 ml |
| 10 mM EDTA | 500 mM pH 8 | 1 ml |
| 1% SDS (or LiDS) | 5% w/v | 10 ml |
| 5 mM DTT | 0.5 M | 500 µl |
| Antifoam A | | 750 ul |
| RNAse-free H2O | | Make up to 50 ml |

1. Add 5 µl/ml 2-Mercaptoethanol per ml before use.
2. Store at 4 C; warm up to RT by placing at 37 C before use (make sure salt crystals are all dissolved)
3. Shelf life: 1-2 months

■ Washing Buffer A (WBA)

| A | B | C |
|----------------|-------------|-------------------|
| Component | Stock | Volumes for 50 ml |
| 10 mM Tris-HCl | 1 M pH 8 | 500 µl |
| 150 mM LiCl | 8 M | 940 µl |
| 1 mM EDTA | 500 mM pH 8 | 100 µl |
| 0.1% SDS | 5% w/v | 500 µl |
| RNAse-free H2O | | Make up to 50 ml |

Store at 4 C and keep on ice prior to use

■ Washing Buffer B (WBB) (Store at 4 C and keep on ice prior to use)

| A | B | C |
|----------------|-------------|------------------|
| Component | Stock | Volume for 50 ml |
| 10 mM Tris-HCl | 1 M pH 8 | 500 µl |
| 150 mM LiCl | 8 M | 940 µl |
| 1 mM EDTA | 500 mM pH 8 | 100 µl |
| RNAse-free H2O | | Make up to 50 ml |

Store at 4 C and keep on ice prior to use

■ Low-salt Buffer (LSB) (Store at 4 C and keep on ice prior to use)

| A | B | C |
|----------------|-------------|------------------|
| Component | Stock | Volume for 50 ml |
| 10 mM Tris-HCl | 1 M pH 8 | 500 µl |
| 150 mM NaCl | 5 M | 1.5 ml |
| 1 mM EDTA | 500 mM pH 8 | 100 µl |
| RNAse-free H2O | | Make up to 50 ml |

Store at 4 C and keep on ice prior to use

- 10 mM Tris-HCl pH 8 (Store at room temperature)

| A | B | C |
|-----------------------------|----------|------------------|
| Component | Stock | Volume for 50 ml |
| 10 mM Tris-HCl | 1 M pH 8 | 500 µl |
| RNAse-free H ₂ O | | Make up to 50 ml |

Store at room temperature

- 1 M (1000 mM) 2-Mercaptoethanol

| A | B | C |
|-----------------------------|--------|------------------|
| Component | Stock | Volume for 50 ml |
| 1 M 2-Mercaptoethanol | 14.3 M | 7 µl |
| RNAse-free H ₂ O | | 93 µl |

Store at -20 °C immediately after use

- Ampure XP Bead Resuspension Buffer (ABR)

| A | B | C |
|-----------------------------|-------|---------------------------------------|
| Component | Stock | Volume for 5 ml (aimed concentration) |
| PEG 8000 | | 1.5 ml (15%) |
| NaCl | | 2.5 ml (2.5 M) |
| RNAse-free H ₂ O | | 1 ml |

Store at room temperature

Remake after a few months; Deteriorate over time, will not DNA well



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Tissue Lysis



5m

- 1 Add  5 µL of 2-Mercaptoethanol (2-ME) to  1 mL of LBB (for ratio, adjust it for the amount LBB used)

- 2 Wipe off RNA later from tissue; Place  **20 mg** of tissue in crushing tube with metal cone
- 3 Add  **200 µL** (100/10: LBB/Tissue ratio) in tube
- 4 Crush sample with multi-beads shocker at  **2000 rpm , 2-4 times**
- 5 Sit in  **Room temperature** for  **00:05:00** and remove metal cone 5m
- 6 Centrifuge at  **14000 rpm, 00:01:00**
- 7 Transfer all the lysate to a new 1.5 mL tube
- 8 Centrifuge at  **14000 rpm, 00:10:00**
- 9 Carefully transfer the supernatant to a new tube. Be careful not to carry over cell debris

Stop here and store samples in  **-80 °C** if needed

1st mRNA extraction 10m

- 10 Put  **100 µL** lysed sample in each well of 8-strip (Keep another half in  **-80 °C** just in case)

11 Add **2 μ L (6.25 μ m)** of biotin-20nt-20T oligo; Mix well by pipetting

12 Sit in **Room temperature** for **00:10:00** for incubation

10m

12.1 While waiting, prepare NEB magnetic Streptavidin beads
Resuspend beads well before use

12.2 Dispense **20 μ L** of Streptavidin beads into each well of a 8-strip (**1 mg : 1 μ L** beads/tissue ratio); Put 8-strip on magnetic rack and remove supernatant

12.3 Resuspend beads with **100 μ L** LBB to wash the beads;
Place 8-strip on magnet rack and remove supernatant
Beads are ready for use

13 Add biotin-incubated samples to washed beads

14 Slowly stir at **Room temperature** at **500 rpm** for **00:10:00**

10m





15 Place 8-strip on magnet rack and remove supernatant


If DNA is needed, keep the supernatant



2-ME in the solution

16 Wash with  150 µL of cold WBA (keep it as cool as possible) and Place 8-strip on magnet rack and remove supernatant

17 Wash with  150 µL of cold WBB (keep it as cool as possible) and Place 8-strip on magnet rack and remove supernatant




18 Wash with  150 µL of cold LSB (keep it as cool as possible) and Place 8-strip on magnet rack and remove supernatant

19 RNA elution buffer prep.

| A | B | C |
|-------------------|-------------|-----------------|
| Component | Stock conc. | Volume for 1 mL |
| Tris-HCl | 10 mM | 999 µl |
| 2-Mercaptoethanol | 1 M | 1 µl |

Freshly make every time before use

20 Resuspend beads in  17 µL RNA elution buffer

21 Warm at  80 °C for  00:02:00 in a thermal cycler. After that, cool it quickly on ice for  00:05:00^{7m}

Keep TTT away from AAA

22 Place 8-strip on magnet rack and **transfer supernatant** ( **17 µL**) to a new 8-strip

Supernatant here is GOOD for **DGE protocol**

22.1 DNAase treatment for Secondary mRNA Recovery (for SHO protocol)




22.2 RNase-Free DNase

[Set Qiagen Catalog #79254](#)

DANase prep.

| A | B |
|------------|---------|
| Component | Volume |
| RDD buffer | 1.85 uL |
| DNAase I | 0.46 uL |
| total | 2.31 uL |


DNAase (1500 K units): add 550 uL DEPC water, divided in small amount in tubes for use; shelf life 9 months















22.3 Add  **2.31 µL** DNAase to 1st RNA supernatant ( **17 µL**) =  **19.31 µL** in total

22.4 Sit at  **Room temperature** for  **00:15:00** 15m

22.5 Kill the activity of DNAase at  **70 °C** for  **00:10:00** 10m

2nd mRNA recovery 10m

23 Add  **150 µL** DEPC water to re-suspend the used beads;
Place 8-strip on magnet rack and remove supernatant

- 24 Add  **5 μ L** of 2-Mercaptoethanol (2-ME) to  **1 mL** of SBB (for ratio, adjust it for the amount SBB used)
- 25 Add  **150 μ L** SBB and Place 8-strip on magnet rack and remove supernatant (Wash beads)
- 26 Add in DNAase treated RNA ( **19.31 μ L**) to the washed beads +  **130 μ L** SBB
- 27 Sit at  **Room temperature** for  **00:10:00** 10m
- 28 Place 8-strip on magnet rack and remove supernatant
- 29 Wash with  **150 μ L** of cold WBA (keep it as cool as possible) and Place 8-strip on magnet rack and remove supernatant
- 30 Wash with  **150 μ L** of cold WBB (keep it as cool as possible) and Place 8-strip on magnet rack and remove supernatant
- 31 Wash with  **150 μ L** of cold LSB (keep it as cool as possible) and Place 8-strip on magnet rack and remove supernatant
- 32 Resuspend beads in  **17 μ L** RNA elution buffer
- 33 Warm at  **80 $^{\circ}$ C** for  **00:02:00** in a thermal cycler. After that, cool it quickly on ice for  **00:05:00**

Keep TTT away from AAA

- 34 Place 8-strip on magnet rack and **transfer supernatant (17 μ L) to a new 8-strip**

Supernatant here is GOOD for **SHO protocol**

Can be stored at **-20 °C** if needed

RNA fragmentation & 3-prime adaptor cDNA priming

5m

- 35 Make 3 strand priming **2.5 μ L**

| A | B |
|--------------------------------|-------------|
| Component | Volume |
| 5X Thermo Scientific RT buffer | 1.5 μ l |
| 3-prime priming adaptor | 1 μ l |

DGE 3' priming adaptor L-3ILL-20TV.2

GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTTTTTTTTTTTTTTTTTTTT

Shotgun 3' priming adaptor L-3ILL-N8.2 GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNNNN

- 36 Mix **2.5 μ L** 3 strand priming with **7.5 μ L** extracted RNA = **10 μ L** in total
Keep the remaining at **-20 °C**)

- 37 **10 μ L** of mixed in thermo cycler for RNA fragmentation

Fragmentation/Priming program: for DGE

(**25 °C 1s** , **94 °C 1.5 min** , **30 °C 1 min** , **20 °C 4 min** , **20 °C hold**)

Fragmentation/Priming program: for SHO

(⌚ 25 °C 1s , ⌚ 94 °C 1.5 min , ⌚ 4 °C 5 min , ⌚ 4 °C hold)

38  5 µL master mix +  10 µL fragmented RNA =  15 µL mixed solution

1st strand master mix

| A | B |
|--------------------------------|---------|
| Component | vol/rxn |
| 5X Thermo Scientific RT buffer | 1.5 µl |
| 0.5M DTT | 0.3 µl |
| H2O | 2.2 µl |
| 25mM dNTPs | 0.5 µl |
| RevertAid RT enzyme | 0.5 µl |

DTT: RNAase inhibitor



RevertAid RT enzyme add right before use

39 Mixed solution in thermo cycler for reverse transcription

Condition:

⌚ 25 °C 10min , ⌚ 42 °C 50min , ⌚ 50 °C 10min , ⌚ 70 °C 10min , ⌚ 4 °C hold

40 Use Ampure beads solution for cDNA purification;

 35 µL Ampure bead solution +  20 µL cDNA

Ampure bead solution


| A | B |
|-------------------|--------|
| Component | Volume |
| 50 mM EDTA pH 8.0 | 5 µl |
| Ampure beads | 30 µl |

Ampure bead = 1.5X sample (3:2)


41 Vortex for ⌚ 00:05:00 at ⌚ Room temperature

5m

42 Place 8-strip on magnet rack and remove supernatant



- 43 Wash with  **200 µL** ethanol without suspending; Dry the pellet
Repeat this process twice

Don't dry the pellet too much at 2nd time, otherwise it's hard to elute

Not recommend to store at  **-20 °C** at this stage

5-prime adapter sequence addition

15m

- 44 Add  **4 µL** of 10 µM 5-prime adapter directly to the pellet at  **Room temperature** to resuspend the bead pellet

- 45 Prepare the master mix in advance during cDNA synthesis, and add the enzyme just before using

 **6 µL** mater mix/rxn +  **4 µL** suspended cDNA pellet



mater mix recipe

| A | B |
|--------------------------|---------|
| Component | Volume |
| H2O | 3.5 µl |
| 10X Poll buffer | 1 µl |
| 250 mM MgCl ₂ | 1 µl |
| 25 mM dNTPs | 0.25 µl |
| DNA Pol I | 0.25 µl |

DNA Pol I add right before use

- 46 Sit at  **Room temperature** for  **00:15:00**

15m

- 47  **40 µL** Ampure bead solution +  **10 µL** cDNA


Ampure bead solution recipe

| A | B |
|---------------------------------------|--------|
| Component | Volume |
| 50 mM EDTA pH 8.0 | 10 µl |
| Ampure Bead Resuspension Buffer (ABR) | 30 µl |



Ampure bead = 1.5X sample (3:2)

48 Vortex for  **00:05:00** at  **Room temperature**


49 Place 8-strip on magnet rack and remove supernatant

50 Wash with  **200 µL** ethanol without suspending; Dry the pellet
Repeat this process twice

Don't dry the pellet too much at 2nd time, otherwise it's hard to elute

51 Elute the pellet in  **20 µL** of 10 mM Tris pH 8.0;
Let it sit for  **00:01:00**

1m

 **12 µL** when sample is little

52 Transfer the supernatant to new tubes

Enrichment and adapter extension

53  **12.2 µL** Enrichment master mix +  **2 µL** 1 µM ILL-INDEX primer +  **5.8 µL** cDNA

Enrichment master mix recipe

| A | B |
|---------------------------------|--------|
| Component | Volume |
| 2 X KAPA | 10 µl |
| 2 µM PE1 primer | 1 µl |
| 8 µM each EnrichS1 + S2 primers | 1 µl |
| 25mM dNTPs | 0.2 µl |

54 Mixed solution in thermo cycler




Condition for SHO

⬆ 98 °C 5 min , (⬆ 98 °C 20 s , ⬆ 65 °C 15 s , ⬆ 72 °C 15 s) X 18 cycles,
⬆ 72 °C 3 min , ⬆ 10 °C hold

Condition for DGE

⬆ 98 °C 5 min , (⬆ 98 °C 20 s , ⬆ 65 °C 15 s , ⬆ 72 °C 15 s) X 14 cycles,
⬆ 72 °C 3 min , ⬆ 10 °C hold

Final Cleanup

- 55  24 µL Ampure beads +  20 µL of enrichment product;
Mix well
- 56 Vortex for ⌚ 00:05:00 at ⬆ Room temperature
- 57 Place 8-strip on magnet rack and remove supernatant
- 58 Wash with  200 µL ethanol without suspending; Dry the pellet
Repeat this process twice

Don't dry the pellet too much at 2nd time, otherwise it's hard to elute



59 Elute the pellet in  12 μL of H₂O



60 Use Bioanalyzer for quantification for measuring the concentration of each sample

Library preparation 10m

61 Mix equimolar of samples in a 1.5 mL tube

62 Add same amount of AMPure XP as the DNA, mix well, and let it stand for  00:05:00 5m

63 Place on magnet rack for  00:05:00 , remove supernatant and wash twice with  70 μL 80% EtOH. Dry. 5m

64 Resuspend the beads by adding  22 μL of water, leave them in the magnet rack for  00:05:00 5m
Transfer the supernatant to a new 0.5 mL low bind tube.

65 Quantification with Qbit or Bioanalyzer