

Mayr lab

3'-seq protocol, October 2013

Time line:

Day -1: DNase treat your RNA samples (optional)

Day 1: Prepare beads, anneal oligo to beads, 1st, 2nd strand synthesis

Day 2: Introduce nick (Rnase HII), nick translation, T7 exo, mung bean nuclease, blunt-end, ligate adapter o/n (over night)

Day 3: Test-PCR, ethanol precipitate, run gel, cut out bands, elute o/n

Day 4: Process all libraries as above

Day 5: Resuspend library

Or:

Day1: Day1+Day2

Day2: Day3+Day4

Day3: Resuspend library

PROTOCOL

DNase treatment of RNA samples:

DNase treatment of 10 ug total RNA with Ambion (DNA-free kit) without using the Inactivation reagent (white solution), instead, after incubation with the DNase enzyme for 30 min at 37°C (volume 50 ul)

Next:

- add 50 ul H₂O, now total volume is 100 ul
- add 100 uL Phenol/Chloroform/Isoamylalcohol (25:24:1), pH=4.5 to each sample
- vortex 30s
- on ice, 10 min
- spin 15 min, 14K, 4°C
- transfer upper phase to new eppi
- add same volume of chloroform to it
- vortex 30s
- on ice, 10 min
- spin 14K, 15 min, 4°C
- take again the upper phase and put it into a new eppi
- add 1/10 of the volume 3M NaOAc and 2.5x volume of 100% EtOH
- put at -20°C, 2 hours
- spin, 20 min, 14K, 4°C
- wash pellet with 300ul 70% EtOH
- spin, 14K, 5 min
- resuspend in 10 ul water
- spec with Nanodrop

Attach oligo(dT) primer to beads:

Prepare beads:

Use a magnetic stand. After the RNA was incubated with the beads we **DO NOT** vortex samples during each wash just wash off side of the tube while pipetting in the wash solution or other solutions; use magnet for 1 min for each step, unless indicated otherwise.

Beads: Dynabeads M280 streptavidin (Invitrogen). 10 mg/ml

Use 50 ul beads/sample

1. resuspend beads by vortexing.
2. transfer 50 ul of beads to siliconized tube
3. magnet 1 min, wash with 500 ul 1x B&W buffer
4. repeat 2x
5. wash 2x in Solution A (200 ul), 2 min
6. wash 2x in solution B.
8. wash 2x in solution 1x BB
9. resuspend in 2x BB buffer (100 ul)

Anneal biotinylated oligo:

Amount of oligo:

Stock: 10 uM of PS193 (/5Biosg/CAGACGTGTGCTCTTCCGATCTTTTTTTTTTrUTTTTTTTTTVN)

Dilute 5 ul Oligo + 95 ul H₂O (per sample)

1. add 100 ul above to 100 ul beads in 2x BB (this dilutes BB to 1x)
2. rotate 15 minutes at room temp
3. magnet 2 min, remove SN
4. wash 8x with 1x B&W buffer (600 ul) (extensive washing to prevent oligo carryover)
5. wash 2x in 1x BB
6. resuspend in 75 ul 2xBB buffer

Bind RNA to the beads:

Set Eppendorf Thermomixer to 65°C

use 2 ug DNase treated total RNA/library

dilute RNA in 75 ul H₂O

1. heat RNA at 65°C, 5 min in Thermomixer
2. warm beads for 1-2 min in Thermomixer
3. mix RNA and beads
4. incubate at 45°C in water bath for 10 min
5. magnet 1 min, remove SN
6. wash 3x with 1x BB buffer (750 ul)
7. wash with 100 uL 1x First strand buffer
8. repeat, have beads in 100 uL 1x First strand buffer

9. change Thermomixer to 50°C

First strand synthesis:

1. prepare First strand mix:

	1x	4.5x	9x	18x	6.5x	5.5x
Water	69					
10X RT buffer	10					
DTT (0.1M)	10					
dNTP mix (10 mM)	5					
RNAse OUT	2					
Total	100					

2. magnet 1 min, discard SN

3. resuspend beads in 96 uL of mix

4. incubate beads with mix at 50°C, 5 min

5. add 4 ul Superscript III, pipette up and down

6. incubate in Thermomixer at 50°C, interval shaking, 1300 rpm x 15 sec, stand x 2 min, 1 hr

7. prepare 2nd Thermomixer at 70°C

8. 70°C, 1300 rpm x 15 s, stand x 2 min, 15 min

9. set first Thermomixer at 16°C

10. put samples on ice, 5 min

Second Strand Synthesis:

1. prepare Second strand mix:

	1x	4.5x	9x
5 X Second strand buffer	27.5		
dNTP mix (10 mM)	3	13.5	27
total	30.5		

2. add Sec. strand mix to RT on ice

3. mix, 5 min on ice

4. prepare enzyme mix

	1x	4.5x	9x
DNA pol I (NEB)	5	22.5	45
RNase H	1	4.5	9
E. coli ligase	1	4.5	9
total	7		
Total SS synthesis	137.5		

5. add enzyme mix to sample

6. spin, mix well

7. 16°C, 1400 rpm x 15 s, stand x 2 min, 2.5 hr

Prepare cleaning solution:

You will need it for 6 steps

	1x	9x	10x	5x
Cleaning sol.	98.6	887.4	986	493
Pronase	1.4	12.6	14	7

Buffer exchange:

1. magnet, discard SN
2. 750 ul buffer C
3. magnet, discard SN
4. 100 ul cleaning solution, spin
5. 37C, 1300 rpm x 15 s, stand x 2 min, 15 min
6. magnet, discard SN
7. 750 ul buffer D wash
8. repeat wash 3x with buffer D
9. have 750 ul buffer D + beads. can stop O/N

From this point on steps 1-8 will be summarized as:
buffer C, cleaning solution, buffer D

Introduce Nick:

1. magnet, discard SN
2. resuspend in 100 ul 1x ThermoPol buffer
3. magnet, discard SN
4. resuspend in 100 ul 1x ThermoPol buffer, spin
5. prepare mix:

	1x	4.5x	9x
10x ThermoPol buffer	10	45	90
RNase HII	(4)		
Water	86	387	774
total	96		

1. magnet, discard SN
2. add 96 ul mix to beads
3. add 4 ul RNaseHII
4. mix by pipetting up and down
5. 37C, 1200 rpm x 15 s, stand x 2 min, 1 hr
NO ICE here....

-buffer C, cleaning solution, buffer D

Nick Translation:

1. magnet, discard SN
2. wash 2x 100 ul 1x NEB buffer 2 (cold)
3. spin
4. make mix:

	1x	4.5x	9x
dNTP (10 mM)	5	22.5	45
NEB2 (10x)	10	45	90
Water	77	346.5	693
total	92		

5. resuspend beads in mix
6. transfer beads into PCR tubes at 8°C in the PCR machine
7. place new PCR tubes also at 8°C into the PCR machine
8. add 8 ul of E.coli DNA pol I (NEB) into empty PCR tubes and incubate for 2 min
9. add beads to enzyme, mix well
10. incubate exactly 8 min at 8°C
11. prepare EDTA stop solution (100 mM)
12. add 100 ul EDTA stop solution (final conc, 50 mM)
13. magnet, discard SN
14. transfer to new siliconized tube

-buffer C, cleaning solution, buffer D
-change Thermomixer to 25°C

-wash 2x 100 ul 1x NEB Buffer 4

T7 exonuclease digestion:

1. prepare mix:

	1x	4.5x	9x
10x NEB 4	10	45	90
T7 exonuclease	(5)		
Water	85	382.5	765
total	95		

2. magnet, discard SN
3. add 95 ul of mix
4. add 5 ul T7 exonuclease
5. mix well by pipetting up and down
6. 25°C, 1400 rpm x 15 s, stand x 2 min, 45 min
7. change Thermomixer to 37°C

-buffer C, cleaning solution, buffer D

-wash 2x 100 ul 1x mung bean nuclease buffer

Mung Bean Nuclease digestion:

1. prepare mix:

	1x	4.5x	9x
10x mung bean nucl buffer	10	45	90
Mung bean nuclease	0.1	0.45	0.9
Water	90	405	810
total	100		

2. magnet, discard SN

3. add 100 ul of the mix to the beads

4. 15°C, 1400 rpm x 15 s, stand x 2 min, 30 min

-buffer C, cleaning solution, set mixer to 25°C, buffer D

-wash 2x 100 ul 1x NEB buffer 2

Klenow Blunt End reaction:

1. Make 1x NEB buffer 2 with 33 uM of each dNTP, you will need 100 ul of this mix/sample:

	1x	10x
NEB buffer 2	10	100
dNTP (10mM)	0.33	3.3
Water	89.7	896.7
total	100	1000

2. magnet, discard SN

3. add 99 ul of the mix to the beads

4. add 1 ul Klenow enzyme

5. 25°C, 1400 rpm x 15 s, stand x 2 min, 30 min

-buffer C, cleaning solution, set mixer to 16°C, buffer D

-wash 2x 100 ul 1x T4 ligase buffer

-transfer to new siliconized tube

Ligate adapter:

1. prepare mix:

	1x	9x
10x T4 buffer	10	90
T4 ligase	(5)	
50 uM PS185+PS186*	4.15	37.35
Water	80.85	727.65
total	95	

(*) PS185: AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCT
 PS186: AGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGTAGATCTCGGTGGTCGCCGTAT
 PS185+PS186 are pre-annealed, gel-purified Illumina Truseq universal Ad1 and Ad2

2. magnet, discard SN
3. add 95 ul of the mix to the beads
4. add 5 ul T4 ligase
5. mix by pipetting up and down
5. 16°C, 1400 rpm x 15 s, stand x 2 min, o/n
6. 25°C, 1400 rpm x 15 s, stand x 2 min, 1 hr

-buffer C, cleaning solution, buffer D

-8x wash in buffer D (this is to ensure removal of free adapter)

-wash 2x 100 ul 1x Phusion buffer, spin

-transfer to new siliconized tube

-magnet, discard SN

-resuspend in 39 ul of 1x Phusion buffer

(if storing beads at 4°C, spin beads down and make sure cap is tight so beads don't dry out)

Phusion PCR:

1. 98C x 2 min
 2. 98C x 20 s
 3. 63C x 30 s
 4. 72C x 15 s
- cycle to step 2 - 14x cycles
5. 72C x 10 min
 6. hold at 4°C

To check the library: total volume: 100 ul, use 5 ul beads

	1x	4.5x	9x	12.5x
Beads	(5)			
Water	68	306	612	850
5x Phusion buffer	20	90	190	250
PS192*, 10 uM	2.5	11.25	22.5	31.25
Barcode prim., 10 uM	(2.5)			
25 mM dNTP	1	4.5	9	12.5
HotStar Phusion	1	4.5	9	12.5
total	92.5			

(*): PS192: AATGATACGGCGACCACCGAGATC
 PS192 is three nucleotides longer than Truseq universal PCR primer R1

EtOH precipitation:

1. Use 100 ul SN
2. add 250 ul 100% EtOH
3. vortex
4. -20°C, 2 h
5. spin 14K, 15 min, 4°C
6. air dry the pellet, 6-8 min
7. resuspend in 8 ul water
8. let dissolve for 10 min, then resuspend, keep on ice
9. add 3 ul of 5x Ficoll dye, mix

Run Gel:

Novex 8% precast TBE PAGE gels

Use 1x TBE running buffer

Ladder: Low range Fermentas ladder (1 ul ladder, 3 ul 5x Ficoll dye, 7 ul water)

Load 11 ul per lane

1. run gel 200V x 35 minutes
2. stain for 15 minutes in 50 ml TBE + 10 ul Ethidium Bromide (can use top of 1 mL pipette tip box- do not re-use). Light shaking
3. destain 2 x 10 minutes in H₂O. Light shaking
4. put gel on saran wrap.
5. take picture with UV imager

Phusion PCR – large scale:

1. As above, 5 ul of beads, 100 ul total volume, x6 for each library
2. Ethanol Precipitation, combine 3x 100 ul + 30 ul 3M NaOAc + 840 ul 100% EtOH (2 eppis/sample)
3. incubate at -20°C o/n
4. spin 14K, 20 min, 4°C
5. discard SN, wash 1x with 70% EtOH
6. spin, 14K, 10 min, 4°C
7. air dry pellet, 6-8 min
8. add 24 ul H₂O to pellet, let stand for 1 hour, add 9 ul Ficoll-dye
9. load 11 ul per lane (empty lane, Marker, 6x sample, marker, empty lane)
10. Run the gel, 200 V, 45-50 min
11. Stain, 15 min, destain 2x 10 min
12. Put the Gel in saran wrap
13. Take picture and save it
14. Mark the smear: between 150 – 210 bp
15. Cut out bands (cut 2 gel slabs containing 3 lanes each and place into 1.5 ml eppi) (Careful not to touch ladder lane with blade. Use new blade if it touches ladder.)
16. add 400 uL Lonza elution buffer

17. rotate o/n at 4°C
18. remove SN from gel, transfer to new eppi
19. add 1000 ul (2.5x volume) of 100% EtOH, add 40 ul 3M NaOAc, 1ul glycogen
20. -20°C, 2 hr
21. spin, 14K, 20 min, 4°C
22. discard SN, wash with 700 ul 70% EtOH
23. spin, 14K, 10 min, 4°C
24. air dry pellet 5-8 min
25. resuspend in 8 ul of water
26. let dissolve on ice for 15 min, then pipette up and down
27. spec 1.5 ul
28. bring to Genomics facility

Appendix

Oligos used:

PS193_Hiseq_17dT_HPLC
/5Biosg/CAGACGTGTGCTCTTCCGATCTTTTTTTTTrUTTTTTTTTTVN

For Adapter:
PS185 + PS186

For PCR:
Use PS192 as reverse primer and any of the barcode forward primers (PS187-PS190, PS197-198)
PS197: BC1; CAAGCAGAAGACGGCATAACGAGATCGTGATGTGACTGGAGTTCAGACGTGTGCTCTTCCGATC
PS187: BC2; CAAGCAGAAGACGGCATAACGAGATACATCGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATC
PS198: BC3; CAAGCAGAAGACGGCATAACGAGATGCCTAAGTGACTGGAGTTCAGACGTGTGCTCTTCCGATC
PS188: BC6; CAAGCAGAAGACGGCATAACGAGATATTGGCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATC
PS189: BC7; CAAGCAGAAGACGGCATAACGAGATGATCTGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATC
PS190: BC9; CAAGCAGAAGACGGCATAACGAGATCTGATCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATC

The adapter oligos and the forward PCR primers are PAGE purified.

Anneal adapters:

10X annealing buffer

Reagent	Final conc	Stock	10 mL
Tris-HCL pH 7.5	100 mM	1M Tris pH 7.5	1 mL
NaCl	500 mM	2M NaCl	2.5 mL
EDTA	10 mM	0.5 M EDTA pH 8	0.2 mL
Water			6.3 mL

Stock: 50 uM of each oligo in water

-Combine 45 ul Oligo 1 + 45 ul Oligo 2 + 10 ul 10x annealing buffer

-Heat in heat block 95C, 5 min

-remove block from heater, put another hot block on top, let warm to room temp

Bind oligo(dT) oligo to beads:

Binding capacity for ss oligos: 200 pmol (=0.2 nmol) for 1 mg of beads

We have 10 mg/ml beads

If we use 50 ul of beads = 0.5 mg beads/sample

Binding capacity/sample: 100 pmol

Oligo: have 10 uM solution = 10 pmol/ul, therefore use 10 ul oligo / sample

Solutions:

Use 0.025% Tween 20 in all wash solutions

0.1% Tw20: 500 ul in 500 ml

0.01% Tw20: 50 ul in 500 ml

0.025% Tw20: 125 ul in 500 ml

2x Binding & Washing (B&W) buffer

FINAL	STOCK	50 ml
10 mM Tris-HCl pH 7.5	1 M	0.5 ml
1 mM EDTA pH 8	0.5 M	100 ul
2M NaCl	2 M	49.6 ml
Tween 20 (0.05%)		25 ul

Solution A

Final	Stock	50 ml
0.1 M NaOH	3 M	1.67 ml
0.05 M NaCl	5 M	0.5 ml
Water		47.9 ml

Solution B

Final	Stock	50 ml
0.1 M NaCl	5 M	1 ml
Water		49 ml

2x Binding buffer (BB):

FINAL	STOCK	50 ml
20 mM Tris-HCl pH 7.5	1 M	1 ml
1 M LiCl	1 M	48.8 ml
2 mM EDTA pH 8.0	0.5 M	0.2 ml

Buffer C

Final	STOCK	50 ml		
1x PBS	1x PBS	50 ml		
0.025% Tween 20		12.5 ul		

1X TE

10 mM Tris-HCL pH 8.0
2 mM EDTA

Buffer D

FINAL	STOCK	500 ml
10 mM Tris-HCl pH 8	1 M Tris	5
2 mM EDTA	0.5 M EDTA	2
0.025% Tween		125 ul
water		493

Cleaning Solution (without pronase)

FINAL	STOCK	50 ml		
1x PBS		50 ml		
1 mM CaCl ₂	1 M	50 ul		

Pronase- stable at 4°C for 6 months when resuspended (resuspend in water to 10 mg/ml)

25 mM each dNTPs

Have 100 mM dATP, dGTP, TTP, dCTP
Mix 50 ul + 50 ul + 50 ul + 50 ul

10 mM each dNTPs

Have 100 mM dATP, dGTP, TTP, dCTP
Mix 50 ul + 50 ul + 50 ul + 50 ul + 300 ul H₂O

NEB buffer 2 plus 33 uM each dNTPs

Use 10 mM dNTP

Add 3.3 ul dNTPs + 996.7 ul H₂O

5X Ficoll loading dye (high density TBE sample buffer)

Final
18 mM Tris base
18 mM Boric Acid
0.4 mM EDTA
3% Ficoll type 400
0.02% Bromo Blue
0.02% Xylene Cyanol
water

Reagent	10 mL
10X TBE	1 mL
Ficoll Type 400	1.5 g
Bromo Blue- 0.25%	4 mL
Xylene Cyanol- 1%	1 mL
water	4 mL

-store at 4°C for up to 6 months

10X TBE

Reagent	500 mL
Tris Base	54 g
Boric Acid	27.5 g
EDTA 0.5 M pH 8	20 mL
water	

-stable at room temp up to 6 months.

Lonza Elution Buffer

Ammonium acetate 3.85 g

Magnesium acetate 0.215 g

0.5 M EDTA 0.2 mL

10% SDS 1mL

Up to 100 mL with H₂O

Reagents

Company

Invitrogen
Fermentas
Invitrogen
Invitrogen
Roche
NEB
Invitrogen
NEB
NEB
NEB
NEB
NEB
NEB
NEB
NEB
NEB
NEB
Eppendorf
Bioplas/Applied
biosystems
Millipore

Reagent

Dynabeads M-280 streptavidin
GeneRuler Low Range DNA Ladder, 25-700 bp
Superscript III RT
dNTPs, 10mM
Pronase 10 mg/ml
Phusion high-fidelity DNA Polymerase
8% TBE Gels 1.0 mm, 10 well
T4 DNA ligase
Rnase H
T7 exonuclease
DNA Pol I (E.coli)
Klenow, DNA Pol I, Lg (Klenow) fragment
Rnase HII
Mung bean nuclease
Buffer 2
Buffer 4
Thermomixer
1.5ml Flat Top MCT, Natural, 500 Tubes/Pack
(siliconized)
PureProteome Magnetic Stand

Cat

112-05D
SM1191
18080-044
AM8200
10165921001
F-540L
EC6215BOX
M0202M
M0297L
M0263L
M0209L
M0210L
M0288L
M0250S

4165SL or 4030