TruSeq DNA PCR-Free Checklist

Fragment DNA

- ☐ 1 Normalize gDNA with RSB to 55 µl in the DNA plate.
 - ▶ 1 ug for a 350 bp insert size
 - ▶ 2 ug for a 550 bp insert size
- \square 2 [HS] Mix and centrifuge as follows.
 - a Shake at 1800 rpm for 2 minutes.
 - \square b Centrifuge at 280 × g for 1 minute.
- □3 [LS] Pipette to mix, and then centrifuge briefly.
- ☐ 4 Transfer 52.5 µl DNA to Covaris tubes.
- \square 5 Centrifuge at 280 × g for 5 seconds.
- ☐6 Fragment using the appropriate settings:

Table 1 350 bp Insert

Setting	M220	S220	S2	E210
Duty Cycle (%)	20	5	10	
Intensity	_	_	5	5.0
Power (W)	50	175	23	14
Cycles/Burst		200		
Duration (s)	65	50	4	45
Mode	_	Freque	ency sweeping	
Temperature (°C)	20		5.5–6	

Table 2 550 bp Insert

Setting	M220	S220	S2	E210
Duty Cycle (%)	20	5		10
Intensity	_	_		2.0
Power (W)	50	175	9	7
Cycles/Burst		200	0	
Duration (s)	45	25		45
Mode	_	Freque	ency sweeping	
Temperature (°C)	20		5.5–6	

Repair Ends and Select Library Size

]1]2]3	Centrifuge CTE at 600 × g for 5 seconds. Add 10 µl CTE or RSB. Add 40 µl ERP 2 or ERP 3 to each well.
4	 [HS] Mix, centrifuge, and incubate as follows. □ a Shake at 1800 rpm for 2 minutes. □ b Centrifuge at 280 x g for 1 minute. □ c Place on the 30°C microheating system
	for 30 minutes.
]5	[LS] Pipette to mix, centrifuge, and then place on the thermal cycler and run the ERP
	program.
]6	Vortex SPB.
	Using your calculations from the previous step, dilute SPB with PCR-grade water.
8	Vortex diluted SPB.
9	Add 160 µl diluted SPB to each well.
10	Mix as follows.
	► [HS] Shake at 1800 rpm for 2 minutes.
	[LS] Pipette up and down.
	Incubate at room temperature for 5 minutes.
	Centrifuge at 280 × g for 1 minute.
⊿ 13	Place on a magnetic stand until the liquid is
٠.,	clear.
_	Transfer 250 µl supernatant to the CEP plate.
	Vortex undiluted SPB.
	Add 30 µl undiluted SPB. Mix as follows.
	► [HS] Shake at 1800 rpm for 2 minutes.
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▶ [LS] Pipette up and down.

 \square 19 Centrifuge at 280 × g for 1 minute.

☐ 21 Remove and discard all supernatant.

clear.

☐ 18 Incubate at room temperature for 5 minutes.

☐ 20 Place on a magnetic stand until the liquid is

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\square 22 Wash two times with 200 μ l 80% EtOH.
□ 23 Use a 20 µl pipette to remove residual EtOH.
24 Air dry for 5 minutes.
\square 25 Add 17.5 μ I RSB, and then remove from the
magnetic stand.
☐ 26 Mix as follows.
▶ [HS] Shake at 1800 rpm for 2 minutes.
[LS] Pipette up and down.
\square 27 Incubate at room temperature for 2 minutes.
\square 28 Centrifuge at 280 × g for 1 minute.
\square 29 Place on a magnetic stand until the liquid is
clear.
\square 30 Transfer 15 μ l supernatant to the ALP plate.

SAFE STOPPING POINT

If you are stopping, seal the plate and store at -25°C to -15°C for up to 7 days.

Adenylate 3' Ends

\square	Centrifuge CTA at 600 × g for 5 seconds.
\square 2	Add 2.5 µl CTA.
\square 3	Centrifuge ATL or ATL 2 at 600 × g for
	5 seconds.
$\Box 4$	Add 12.5 µl ATL or ATL 2.
\Box 5	[HS] Mix and incubate as follows.
	☐a Shake at 1800 rpm for 2 minutes.
	☐ b Place on the 37°C microheating system
	for 30 minutes.
	C Move to the 70°C microheating system for
	5 minutes.
	d Place on ice for 5 minutes.
□6	[LS] Pipette to mix, and then place on the
	thermal cycler and run the ATAIL70 program.

Ligate Adapters

[HS] Centrifuge the DAP at 280 \times g for 1 minute.
[LS] Centrifuge the adapter tubes at $600 \times g$ for
5 seconds.
Remove LIG 2 from -25°C to -15°C storage.
In the order listed, add the following reagents:
CTL (2.5 μl)
▶ LIG 2 (2.5 μl)
DNA adapters (2.5 μl)
Mix as follows.
▶ [HS] Shake at 1800 rpm for 2 minutes.
[LS] Pipette up and down.
Centrifuge at 280 × g for 1 minute.
Incubate as follows.
► [HS] Place on the 30°C microheating
system for 10 minutes. Set aside on ice.
▶ [LS] Place on the thermal cycler and run
the LIG program.
Centrifuge the STL at $600 \times g$ for 5 seconds.
Add 5 µl STL to each well.
Mix as follows.
► [HS] Shake at 1800 rpm for 2 minutes.
► [LS] Pipette up and down.
Centrifuge at 280 × g for 1 minute.
Add SPB.
► Round 1 — 42.5 µl
▶ Round 2 —50 µl
Mix as follows.
► [HS] Shake at 1800 rpm for 2 minutes.
► [LS] Pipette up and down.
Incubate at room temperature for 5 minutes.

 \square 15 Centrifuge at 280 × g for 1 minute.

☐ 17 Remove and discard supernatant.☐ 18 Wash two times with 200 µl 80% EtOH.

clear.

☐ 16 Place on a magnetic stand until the liquid is



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\square 19 Use a 20 μ l pipette to remove residual EtOH.		
20 Air dry for 5 minutes.		
□21 Add RSB.		
▶ Round 1 − 52.5 μl		
▶ Round 2 — 22.5 µl		
\square 22 Remove from the magnetic stand, and then		
mix as follows.		
► [HS] Shake at 1800 rpm for 2 minutes.		
[LS] Pipette up and down.		
☐ 23 Incubate at room temperature for 2 minutes.		
\square 24 Centrifuge at 280 × g for 1 minute.		
☐ 25 Place on a magnetic stand until the liquid is clear.		
☐ 26 Transfer 50 µl supernatant to the CAP plate.		
☐ 27 Repeat steps 12 through 25 using the new plate and the Round 2 volumes.		
\square 28 Transfer 20 μ I supernatant to the TSP1 plate.		

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Check Libraries

- 1 Quantify libraries using qPCR, with the following modifications:
 - ▶ Use at least 2 µl of the original library stock.
 - Perform two additional dilutions.
 - Determine the concentration of the diluted library.
 - ▶ Make a size adjustment calculation.
 - Calculate the concentration of the undiluted library.
- 2 Verify fragment size by checking the library size distribution.
 - ☐ a Dilute the DNA library 1:5 with water.
 - □ b Run 1 µl diluted DNA library on a High Sensitivity DNA chip or NGS kit.

Normalize and Pool Libraries

	ransfer 5 µi library to the DCT plate.
2	Normalize to 4 nM using Tris-HCl 10 mM, pH 8.5
	with 0.1% Tween 20.

- \square 3 Mix as follows.
 - ▶ [HS] Shake at 1800 rpm for 2 minutes.
 - ▶ [LS] Pipette up and down.
- \square 4 Centrifuge at 280 × g for 1 minute.
- ☐ 5 Proceed to pooling or clustering:
 - lacksquare To pool libraries, proceed to .
 - To leave libraries unpooled, skip the remaining library prep steps and proceed to cluster generation.
- \square 6 Transfer 5 μ l to one well of the PDP plate.
- \square 7 Mix as follows.
 - ▶ [HS] Shake at 1800 rpm for 2 minutes.
 - ▶ [LS] Pipette up and down.
- \square 8 Centrifuge at 280 × g for 1 minute.
- ☐9 Proceed to cluster generation.
- □ 10 Transfer 5 µl to column 1 of the PDP plate.
- ☐ 11 Mix as follows.
 - ► [HS] Shake at 1800 rpm for 2 minutes.
 - ▶ [LS] Pipette up and down.
- \square 12 Centrifuge at 280 × g for 1 minute.
- ☐ 13 Transfer column 1 contents to well A2.
- ☐ 14 Mix as follows.
 - ► [HS] Shake at 1800 rpm for 2 minutes.
 - ▶ [LS] Pipette up and down.
- \square 15 Centrifuge at 280 × g for 1 minute.
- ☐ 16 Proceed to cluster generation.

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Acronyms

Definition
Adapter Ligation Plate
A-Tailing Mix
Clean Up ALP Plate
Clean Up End Repair Plate
Covaris Fragmentation Plate
Clean Up Sheared DNA Plate
A-Tailing Control
End Repair Control
Ligation Control
Diluted Cluster Template Plate
Customer Sample DNA Plate
End Repair Mix
High Sample
Illumina Experiment Manager
Insert Modification Plate
Ligation Mix
Local Run Manager
Low Sample
Pooled Dilution Plate
Resuspension Buffer
Sample Purification Beads
Stop Ligation Buffer
Target Sample Plate 1