

Procedure & Checklist – cDNA Capture Using IDT xGen® Lockdown® Probes

Before You Begin

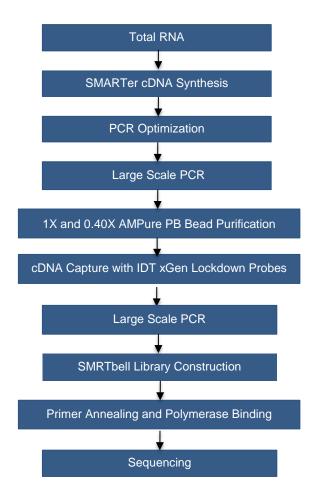
This document describes the process for capturing cDNA prepared with the SMARTer PCR cDNA Synthesis Kit (Clontech) and pulled-down using xGen Lockdown Probes/Panels from IDT.

To perform this procedure, you must have reviewed the <u>Procedure & Checklist – Iso-Seq Template Preparation for Sequel Systems</u>.

Workflow

The workflow includes the following:

- 1. Preparing the cDNA library using the SMARTer PCR cDNA Synthesis Kit.
- 2. Capturing cDNA with the IDT xGen Lockdown Probes (biotinylated probes).
- 3. Constructing SMRTbell® libraries
- 4. Sequencing using the PacBio System.



Materials Needed

Item	Vendor	Part Number
cDNA Library		
SMARTer PCR cDNA Synthesis Kit	KAPA Biosystems	KK8503
Lonza flash gel	Lonza	
PCR/Target Capture		
PolyT blocker Oligo (5' TTT TTT TTT TTT TTT TTT TTT TTT TTT / 3InvdT/3')	IDT	N/A
xGen Hybridization and Wash Kit (16 or 96 reaction)	IDT	
xGen Lockdown Panels/Probes (target probes)*	IDT	
Takara LA Taq DNA Polymerase Hot-Start version	Clontech	RR042A
SMARTer PCR Oligo (5' AAG CAG TGG TAT CAA CGC AGA GTA C 3')	IDT	N/A
Dynabeads® M-270 Streptavidin	Life Technologies	65305
AMPure® PB beads	РасВіо	
SMRTbell Library Construction and Sequencing		
Template Prep Kit	PacBio	
DNA/Polymerase Binding Kit	PacBio	
DNA Sequencing Kit	PacBio	
AMPure® PB beads		

^{*}If the xGen Lockdown Probes were received dry, resuspend them in IDTE pH 8.0 to a final concentration of 0.75pmol/µL. For additional support regarding resuspension of Lockdown Probes pools, visit

www.idtdna.com/xgen ► xGen Lockdown Probes ► Support tab ► expand Number of Reactions and Resuspension Volume.

Prepare cDNA Library

To prepare a cDNA library, refer to pages 1 – 11 (<u>Procedure & Checklist – Iso-Seq Template Preparation for Sequel Systems</u>).

- 1. Prepare 1st-strand synthesis using the SMARTer PCR cDNA Synthesis Kit.
- 2. Enrich by:
 - a. Optimizing PCR cycles.
 - b. Performing large-scale PCR.

STEP	/	Prepare the Hybridization	n Sample	Notes		
		 In this section, you will need the following: SMARTer PCR oligo (IDT) PolyT blocker (IDT) 2X Hybridization Buffer contained in xGen Lockdown Hybridization and Wash Kit Hybridization Buffer Enhancer in xGen Lockdown Hybridization and Wash Kit xGen Lockdown Panels/Probes (target probes) 				
1		Add 1 – 1.5 µg cDNA to a new 0.5 mL LoBind tub	De.			
2		Add 1 μL of SMARTer PCR oligo and 1 μL PolyT μM) to the tube containing the cDNA.	blocker (both at 1000			
3		Close the tube's lid and puncture a hole in the cap with an 18 – 20 gauge or smaller needle.				
4		Dry the cDNA Sample Library/SMARTer PCR oligo/PolyT blocker completely in a LoBind tube using a DNA vacuum concentrator (speed vac). Do not leave tubes in the speed vac once they have dried. This will result in over drying the tube contents.				
5		To the dried-down sample add:				
		Component	Volume			
		2X Hybridization Buffer	8.5 µL			
		Hybridization Buffer Enhancer	2.7 µL			
		Nuclease Free Water	1.8 µL			
6		Seal the hole in the tube's cap with a laboratory to	ape.			
7		Mix the reaction by tapping the tube, followed by	a quick spin.			
8		Place the tube in a +95℃ h eat block for 10 minut	es to denature the cDNA.			
9		Quick spin at maximum speed, allowing the mix to cool to room temperature before addition of probes. Probes should never be added while at 95°C.				
10		Leave the PCR tube on the bench for approximately 30 seconds, then add 4 μL of xGen Lockdown Panel/Probe for a total volume of 17 μL.				
11		Mix and quick spin.				
12		Incubate in a thermocycler at +65℃ for 4 hours. Ilid should be turned on and set to maintain +75℃ hybridization temperature).				

STEP	/	Preparinç	g Beads	for Ca	pture			Notes	
1		In this section, you will need the following: Wash buffers contained in the xGen Lockdown Hybridization and Wash Kit Dynabeads M-270 Streptavidin Prepare Wash Buffers: a. Prepare 1X working solutions of the buffers listed in the below table. The							
		total volume of 1X buffer in the table is for a single experiment. Scale up accordingly when multiple samples are required. Ensure that 10x Wash Buffer I is in solution before use. Any solids can be dissolved by warming with your hands or incubate at 65℃ heat block and vortexing.							
		Buffer Stock	Conc.	Vol. Buffer	Water	Volume*	Final Conc.		
		Wash Buffer I (tube 1)	10X	30 µL	270 µL	300 µL	1X		
		Wash Buffer II (tube 2)	10X	20 μL	180 µL	200 µL	1X		
		Wash Buffer III (tube 3)	10X	20 μL	180 µL	200 µL	1X		
		Stringent Wash Buffer (tube S)	10X	40 µL	360 µL	400 µL	1X		
		Bead Wash Buffer *Store working solutions at room temperature.	2X	250 µL	250 μL	500 µL	1X		
		are calculated for a single capture; scale up b. Preheat the following wash to the follo	o according ouffers to Buffer I (T	ly if multiple > +65℃ in ube1)	hybridizatior a heat b	n reactions wil	I be processed.		
2		Prepare the capture beads: a. Allow the Dynabeads M-2 temperature for 30 minute b. Mix the beads thoroughly c. For a single sample, aliqued Scale up volume for multiple d. Place the LoBind tube in a remove and discard the subeads. Any remaining trace wash steps. Note: Allow the before removing the supercollect to the side of the tube. While the LoBind tube is in Wash Buffer. For multiple f. Remove the tube from the in solution. g. Quickly spin and place the collect the beads to the side the liquid. h. Repeat steps e - g for a to i. Resuspend by vortexing the multiple samples, scale up j. Place the tube in the magniform multiple samples, tranclear, remove and discard k. The washed beads are no immediately to the next stem Small amounts of residual binding of DNA to the cap	s prior to byvortex of 100 µl ole sample imagnet in pernatant. The imagnet in the magnet in the magnet in the magnet in the magnet in the samples in according to the super weady the super in the super	ing for 15 beads in les. ic rack. What being could will be eads to so he Dyna gnetic rack, wash was rack and tube back tube. On washes, in 100 plangly, a to collect plangly aliquoto o bind the ot allow thash Buffer ash Buffer washes.	is seconds ato a 1.5 I hen the stareful no removed the for a beads are k, add 20 ith 200 µld vortex k in the mode clear, L of 1X B at beads to the into never experience capture and a 1.5	s. mL LoBind supernatar t to disturb d with subs at least 1-2 e "filmy" ar 0 µL of 1X L x X samp until the be agnetic ra remove ar ead Wash o the side v LoBind to d DNA. Pro e beads to	ont is clear, of the sequent s		

STEP	Binding cDNA to Beads and Wash	Notes
1	Bind cDNA to the capture beads:	
	a. To the washed capture beads, transfer the 17 µL hybridized probe/sample mixture prepared in the "Preparing hybridization section".	
	 b. Mix by tapping the tube until the sample is homogeneous. 	
	c. Incubate in a thermomixer set to +65℃ for 45 minutes or transfer the mix to a PCR tube and incubate in a thermocycler (heated lid set to +75℃). Hand mix periodically by gently tapping the tube to keep the beads in suspension.	
2	Wash the captured cDNA:	
	a. Pre-heat 1X Wash Buffer (tube 1) and 1X Stringent Wash Buffer (Tube S) to 65℃,	
	b. After 45 minutes of incubation, remove the tube from the 65°C thermomixer and add 100 μL pre-heated 1X Wash Buffer I (Tube 1).	
	c. Mix thoroughly by tapping the tube until the sample is homogeneous.	
	d. If using a PCR tube, transfer the sample to a 1.5 mL LoBind tube.	
	 e. Place the tube in the magnetic rack to collect the beads to the side of the tube. Remove and discard the liquid once clear. 	
	f. Remove the tube from the magnetic rack and add 200 μL of 1X Stringent Wash Buffer (TubeS) heated to +65°C. Mix by tapping the tube until the sample is homogeneous. Work quickly so that the temperature does not drop below +65°C.	
	g. Incubate at +65℃ for 5 minutes.	
	h. Repeat steps d - f for a total of two washes using 1X Stringent Wash Buffer (TubeS) heated to +65℃.	
	 Place the tubes in the magnetic rack to collect the beads to the side of the tube. Remove and discard the liquid once clear. 	
	 j. Add 200 μL of room temperature 1X Wash Buffer I (Tube1). Hand mix by gently tapping the tube. Quick spin. 	
	 k. Place the tube in the magnetic rack to collect the beads to the side of the tube. Remove and discard the liquid once clear. 	
	 I. Add 200 μL of room temperature 1X Wash Buffer II (Tube2) and mix thoroughly by tapping the tube until Sample is homogeneous. Quick Spin. 	
	 m. Place the tube in the magnetic rack to collect the beads to the side of the tube. Remove and discard the liquid once clear. 	
	 n. Add 200 μL of room temperature 1X Wash Buffer III (Tube 3) and mix by tapping the tube until sample is homogeneous. Quick Spin. 	
	 Place the tubes in the magnetic rack to collect the beads to the side of the tube. Remove and discard the liquid once clear. 	
	p. Remove the tubes from the magnetic rack and add 50 μL of EB to each tube of bead-bound captured sample. This is enough for two PCR reactions required in the next section.	
	q. Store the beads plus captured samples at -15 to -25℃ or proceed to the next step. It is not necessary to separate the beads from the eluted DNA. The bead/sample mix can be added to the PCR reaction directly.	

STEP	✓		Amplifica	tion of Capture	ed DNA Sample		Notes	
		• Taka Clon • SMA	In this section, you will need the following: Takara LA Taq DNA Polymerase Hot-Start Version from Clontech SMARTer PCR Oligos (from the Clontech Kit)					
1		version.		y Takara LA Taq D owing PCR reactio	NA Polymerase Hone	nt-Start		
			Compo	onent	Volume			
		Water			104.5 μL			
		10x LA I	PCR Buffer		20 μL			
		2.5 mM	each dNTPs		16 µL			
		SMART	er PCR Oligos	(12 µM each)	8.3 µL			
		Takara I	_A Taq DNA p	olymerase	1.2 µL			
		Capture	d Library		50 μL			
		Total			200 μL			
		PCR	reaction in 10	into two tubes, 10 00 μL volumes. ollowing PCR prot	0 μL each. It is bes	t to perform the		
		Step	Temp	Time				
		1	95℃	2 mi	inutes			
		2	95℃	20 se	econds			
		3	68℃	1 0 m	inutes			
		4	Repeat step	Repeat steps 2-3, 7 to10 times for a total of 8 to 11 cycles				
		5	72℃	1 0 m	inutes			
		6	4℃	Н	old			
2		·	•	e 100 µL reactions ure PB beads.	s and proceed to th	e next step to		

STEP	/	Post Amplification Clean Up	Notes
1		Add 1X AMPure PB beads to the pooled PCR product.	
2		Mix by tapping the LoBind tube until the sample is homogeneous.	
3		Incubate at room temperature for 10 minutes.	
4		Place on magnetic rack until solution clears. Remove and discard supernatant.	
6		With the tube still on magnet, add 200 µL freshly prepared 70% ethanol to the tube containing beads plus DNA.	
7		Remove and discard 70% ethanol.	
8		Repeat steps 5 to 6 for total of two washes with 70% ethanol.	
9		Let beads air dry for 1 minute. (Note - over drying the beads will result in reduced DNA yield.)	
10		Add 27 µL EB and remove the tube from the magnet. Mix by tapping the tube until the sample is homogeneous. Then incubate at room temperature for 2 minutes.	
11		Place back on magnet. When the solution clears, remove 25 µL supernatant into new 1.5 mL LoBind tube.	
12		Determine concentration using Qubit device or similar quantification assay.	
13		Run 1 µL of sample on Agilent DNA 12000 chip according to manufacturer's instructions.	
14		The captured cDNA is now ready for SMRTbell library construction.	

Repair DNA Damage

Use the following table to repair any DNA damage. If preparing larger amounts of DNA, scale the reaction volumes accordingly (i.e., for 10 μ g of DNA scale the total volume to 100 μ L). Do not exceed 100 μ L of DNA in the final reaction.

1. In a LoBind microcentrifuge tube, add the following reagents:

Reagent	Cap Color	Stock Conc.	Volume	Final Conc.	✓	Notes
cDNA			μL for 5.0 μg			
DNA Damage Repair Buffer	•	10 X	5.0 μL	1 X		
NAD+	•	100 X	0.5 μL	1 X		
ATP high	•	10 mM	5.0 µL	1 mM		
dNTP	•	10 mM	0.5 μL	0.1 mM		
DNA Damage Repair Mix	•		2.0 µL			
H ₂ O			μL to adjust to 50.0* μL			
Total Volume			50.0 μL			

 $^{^{\}star}$ To determine the correct amount of $\mathrm{H}_{2}\mathrm{O}$ to add, use your actual DNA amount noted in the Notes column.

- 2. Mix the reaction well by gentle mixing.
- 3. Spin down contents of LoBind tube with a quick spin in a microfuge.
- 4. Incubate at 37°C for 20 minutes, then return the reaction to 4°C for 1 minute.

Repair Ends

Use the following table to prepare your reaction then purify the DNA.

Reagent	Tube Cap Color	Stock Conc.	Volume	Final Conc.	✓	Notes
cDNA (Damage Repaired)			50 μL			
End Repair Mix		20 X	2.5 μL	1X		
Total Volume			52.5 μL			

^{1.} Mix the reaction well by gentle mixing.

^{2.} Spin down contents of LoBind tube with a quick spin in a microfuge.

^{3.} Incubate at 25°C for 5 minutes, return the reaction to 4°C.

STEP	✓	Purify DNA	Notes
1		Add 1X volume of AMPure PB beads to the End-Repair reaction.	
2		Mix the bead/DNA solution by tapping the tube.	
3		Allow the DNA to bind by letting it sit at room temperature for 10 minutes.	
4		Spin down the LoBind tube (for 1 second) to collect beads.	
5		Place the LoBind tube in a magnetic bead rack to collect the beads to the side of the tube.	
6		Slowly pipette off cleared supernatant and save (in another tube). Avoid disturbing the bead pellet.	
7		Wash beads with freshly prepared 70% ethanol.	
8		Repeat step 8 above.	
9		 Remove residual 70% ethanol and dry the bead pellet. Remove the LoBind tube from the magnetic bead rack and spin to pellet beads. Both the beads and any residual 70% ethanol will be at the bottom of the tube. Place the LoBind tube back on magnetic bead rack. Pipette off any remaining 70% ethanol. 	
10		Check for any remaining droplets in the tube. If droplets are present, repeat step 10.	
11		Remove the LoBind tube from the magnetic bead rack and allow beads to air-dry (with LoBind tube caps open) for 30 to 60 seconds.	
12		Elute the DNA off the beads in 30 µL Elution Buffer. Mix by gently tapping the LoBind tube until homogenous, then let stand at room temperature for 2 minutes.	
13		Optional: Verify your DNA amount and concentration using a Nanodrop or Qubit quantitation platform, as appropriate.	
14		Optional: Perform qualitative and quantitative analysis using a Bioanalyzer system instrument with the DNA 12000 Kit. Note that typical yield at this point of the process (following End-Repair and one 1X AMPure PB bead purification) is approximately between 80-100% of the total starting material.	
15		The End-Repaired DNA can be stored overnight at 4°C or at -20°C for longer duration.	

Prepare Blunt-Ligation Reaction

Use the following table to prepare your blunt-ligation reaction:

1. In a LoBind microcentrifuge LoBind tube (on ice), add the following reagents in the order shown. Note that you can add water to achieve the desired DNA volume. If preparing a Master Mix, ensure that the adapter is NOT mixed with the ligase prior to introduction of the inserts. Add the adapter to the well with the DNA. All other components, including the ligase, should be added to the Master Mix.

Reagent	Tube Cap Color	Stock Conc.	Volume	Final Conc.	✓	Notes
DNA (End Repaired)			29.0 μL to 30.0 μL			
Blunt Adapter (20 µM)	<u> </u>	20 μΜ	1.0 µL	0.5 µM		
	,	Mix befor	e proceeding			
Template Prep Buffer		10 X	4.0 μL	1X		
ATP low	<u> </u>	1 mM	2.0 µL	0.05 mM		
		Mix befor	e proceeding			
Ligase	<u> </u>	30 U/μL	1.0 µL	0.75 U/μL		
H ₂ O			μL to adjust to 40.0 μL			
Total Volume			40.0 μL			

- 2. Mix the reaction well by gentle mixing.
- 3. Spin down contents of LoBind tube with a quick spin in a microfuge.
- 4. Incubate at 25°C for 15 minutes. At this point, the ligation can be extended up to 24 hours or cooled to 4°C (for storage of up to 24 hours).
- 5. Incubate at 65°C for 10 minutes to inactivate the ligase, then return the reaction to 4°C. You must proceed with adding exonucleases after this step.

Exo III and Exo VII Treatment

Reagent	Tube Cap Color	Stock Conc.	/	Volume
Ligated DNA				40 μL
	Mix reaction	on well by pipetting		
ExoIII	•	100.0 U/μL		1.0 µL
ExoVII	•	10.0 U/μL		1.0 μL
Total Volume				42 μL

- 1. Spin down contents of LoBind tube with a quick spin in a microfuge.
- 2. Incubate at 37°C for 1 hour, then return the reaction to 4°C. You must proceed with purification after this step.

Purify SMRTbell Templates

STEP	Purify SMRTbell Templates	Notes
1	Add 1X volume of AMPure PB beads to the exonuclease-treated reaction.	
2	Mix the bead/DNA solution by tapping the tube.	
4	Allow the DNA to bind to beads by letting the sample sit at room temperature for 10 minutes.	
4	Spin down the LoBind tube (for 1 second) to collect beads.	
5	Place the LoBind tube in a magnetic bead rack to collect the beads to the side of the tube.	
6	Slowly pipette off cleared supernatant and save (in another tube). Avoid disturbing the bead pellet.	
7	Wash beads with freshly prepared 70% ethanol.	
8	Repeat step 8 above.	
9	 Remove residual 70% ethanol and dry the bead pellet. Remove the LoBind tube from the magnetic bead rack and spin to pellet beads. Both the beads and any residual 70% ethanol will be at the bottom of the tube. Place the LoBind tube back on the magnetic bead rack. Pipette off any remaining 70% ethanol. 	
10	Check for any remaining droplets in the tube. If droplets are present, repeat step 10.	
11	Remove the LoBind tube from the magnetic bead rack and allow beads to air-dry (with LoBind tube caps open) for 60 seconds.	
12	Elute the DNA off the beads in 50 µL of Elution Buffer. Mix thoroughly by gently tapping the LoBind tube and let sit at room temperature for 2 minutes.	
13	The eluted DNA in $50~\mu L$ Elution Buffer should be taken into the second $1X$ AMPure PB bead purification step.	

STEP	Second Purification	Notes
1	Add 1X volume of AMPure PB beads to the 50 µL of eluted DNA.	
2	Mix the bead/DNA solution by tapping the tube.	
4	Allow the DNA to bind to beads by letting the sample sit at room temperature for 10 minutes.	
4	Spin down the LoBind tube (for 1 second) to collect beads.	
5	Place the LoBind tube in a magnetic bead rack to collect the beads to the side of the tube.	
6	Slowly pipette off cleared supernatant and save (in another tube). Avoid disturbing the bead pellet.	
7	Wash beads with freshly prepared 70% ethanol.	
8	Repeat step 8 above.	
9	 Remove residual 70% ethanol and dry the bead pellet. Remove the LoBind tube from the magnetic bead rack and spin to pellet beads. Both the beads and any residual 70% ethanol will be at the bottom of the tube. Place the LoBind tube back on the magnetic bead rack. Pipette off any remaining 70% ethanol. 	
10	Check for any remaining droplets in the tube. If droplets are present, repeat step 10.	
11	Remove the LoBind tube from the magnetic bead rack and allow beads to air-dry (with LoBind tube caps open) for 60 seconds.	
12	Elute the DNA off the beads in 50 µL of Elution Buffer. Mix thoroughly by gently tapping the LoBind tube and let sit at room temperature for 2 minutes.	
13	Elute the off the beads in 10 μL. Mix thoroughly.	
14	Determine concentration using a Qubit device or similar quantification assay.	
15	Run 1 µL of sample on Agilent DNA 12000 chip according to manufacturer's instructions. Perform qualitative analysis using a Bioanalyzer instrument with the DNA 12000 kit. Refer to Agilent Technologies' guides for specific information.	

DNA Control Complex Dilution

You must have the PacBio Control Complex for this step. Dilute the Control Complex according to the volumes and instructions specified in Sample Setup.

Anneal and Bind SMRTbell Templates

Follow the instructions in Sample Setup to anneal and bind your library.

Revision History (Description)	Version	Date
Initial release. Converted from "Unsupported Protocol" with updates to the cDNA preparation section.	01	June 2018

For Research Use Only. Not for use in diagnostic procedures. © Copyright 2013 - 2018, Pacific Biosciences of California, Inc. All rights reserved. Information in this document is subject to change without notice. Pacific Biosciences assumes no responsibility for any errors or omissions in this document. Certain notices, terms, conditions and/or use restrictions may pertain to your use of Pacific Biosciences products and/or third party products. Please refer to the applicable Pacific Biosciences Terms and Conditions of Sale and to the applicable license terms at https://www.pacb.com/legal-and-trademarks/terms-and-conditions-of-sale/ Pacific Biosciences, the Pacific Biosciences logo, PacBio, SMRT, SMRTbell, Iso-Seq, and Sequel are trademarks of Pacific Biosciences in the United States and/or certain other countries. All other trademarks are the sole property of their respective owners.