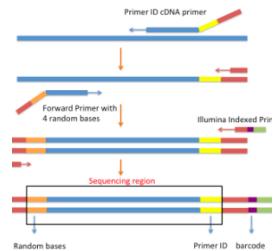


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Primer ID MiSeq Library Prep V.3

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Protocol status: Working

We use this protocol and it's working

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Abstract

This is the protocol to prepare Primer ID MiSeq sequencing library. Viral RNA was first extracted using QIAamp viral RNA extraction kit. The block of random nucleotides (Ns) in the cDNA primers served as the Primer ID. The Superscript III kit was used for the cDNA synthesis. We used two rounds of PCR to amplify the cDNA and incorporate Illumina indexed adapters with KAPA2G Robust and KAPA HiFi PCR kits, respectively.

Attachments

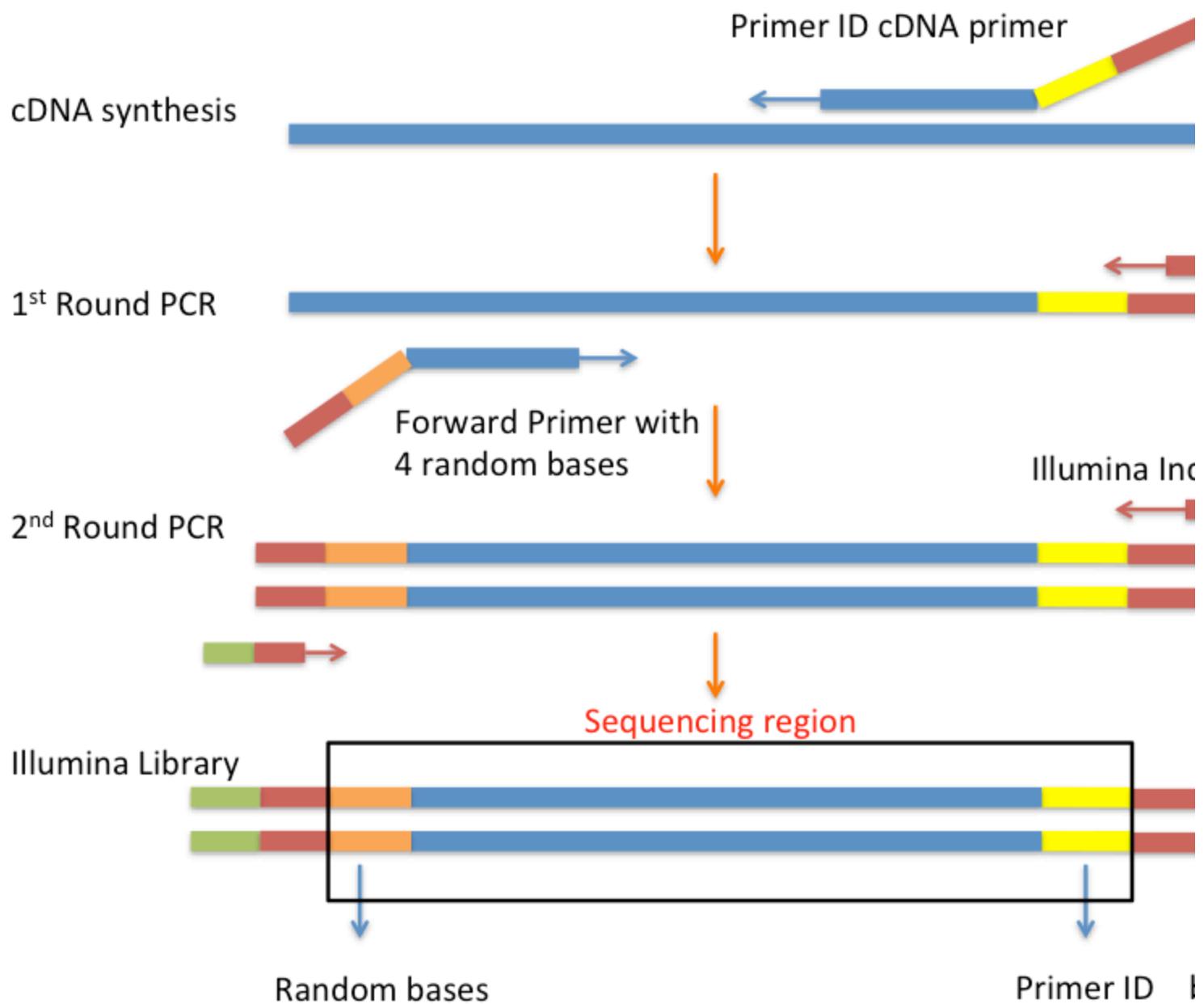


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Guidelines

Viral RNA extraction using QIAamp vRNA mini kit.



Primers (NOTE: HIV-1 ENV REGION AS AN EXAMPLE, SUBSTITUTE 1 in the forward primer and cDNA primer)

V1F(forward) 5'-3'	GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAG AAAGCCTAAAGCCATGTGTA
BV3R Uni (cDNA Primer) 5'-3'	GTGACTGGAGTTCAGACGTGTGCTTCCGATCTNNNN TGCTCTACTAATGTTACAATGTGC
Universal Adapter	AATGATACGGCGACCACCGAGATCTACACGCCTCCCTC G
Indexed Adapter	CAAGCAGAACGGCATACGAGATNNNNNNGTGACT GCTC
ADPT_2a	GTGACTGGAGTTCAGACGTGTGCTC

Note: Primer ID primer and forward primer use random bases. Indexed primers have 24 fixed barcodes.

Table of Indexed Primers

Indexed Primer	Index	Index Sequence	Sequence
PCR Primer, Index 1	1	ATCACGA	CAAGCAGAAGACGGCATACGAGAT CGTGATGT
PCR Primer, Index 2	2	CGATGTA	CAAGCAGAAGACGGCATACGAGAT ACATCGGT
PCR Primer, Index 3	3	TTAGGCA	CAAGCAGAAGACGGCATACGAGAT GCCTAAGT
PCR Primer, Index 4	4	TGACCAA	CAAGCAGAAGACGGCATACGAGAT TGGTCAGT
PCR Primer, Index 5	5	ACAGTGA	CAAGCAGAAGACGGCATACGAGAT CACTGTGT
PCR Primer, Index 6	6	GCCAATA	CAAGCAGAAGACGGCATACGAGAT ATTGGCGT
PCR Primer, Index 7	7	CAGATCA	CAAGCAGAAGACGGCATACGAGAT GATCTGGT
PCR Primer, Index 8	8	ACTTGAA	CAAGCAGAAGACGGCATACGAGAT TCAAGTGT
PCR Primer, Index 9	9	GATCAGA	CAAGCAGAAGACGGCATACGAGAT CTGATCGT
PCR Primer, Index 10	10	TAGCTTA	CAAGCAGAAGACGGCATACGAGAT AAGCTAGT
PCR Primer, Index 11	11	GGCTACA	CAAGCAGAAGACGGCATACGAGAT GTAGCCGT
PCR Primer, Index 12	12	CTTGTAA	CAAGCAGAAGACGGCATACGAGAT TACAAGGT
PCR Primer, Index 13	13	TCCATAA	CAAGCAGAAGACGGCATACGAGAT TATGGA GT
PCR Primer, Index 14	14	GTACTAA	CAAGCAGAAGACGGCATACGAGAT TAGTACGT
PCR Primer, Index 15	15	ACAGTAA	CAAGCAGAAGACGGCATACGAGAT TACTGTGT
PCR Primer, Index 16	16	CTCATGA	CAAGCAGAAGACGGCATACGAGAT CATGAGGT
PCR Primer, Index 17	17	ACGATAA	CAAGCAGAAGACGGCATACGAGAT TATCGTGT
PCR Primer, Index 18	18	TGCAGAA	CAAGCAGAAGACGGCATACGAGAT TCTGCA GT
PCR Primer, Index 19	19	TTCATAA	CAAGCAGAAGACGGCATACGAGAT TATGAA GT
PCR Primer, Index 20	20	TGCTGTA	CAAGCAGAAGACGGCATACGAGAT ACAGCA GT
PCR Primer, Index 21	21	TATCACA	CAAGCAGAAGACGGCATACGAGAT GTGATA GT
PCR Primer, Index 22	22	TGGATAA	CAAGCAGAAGACGGCATACGAGAT TATCCA GT
PCR Primer, Index 23	23	CGCATT	CAAGCAGAAGACGGCATACGAGAT AATGCG GT
PCR Primer, Index 24	24	GCCTTAA	CAAGCAGAAGACGGCATACGAGAT TAAGGC GT

Materials

MATERIALS

☒ QIAamp vRNA mini kit Qiagen Catalog #51304

☒ Agencourt RNAClean XP Beckman Coulter Catalog #A63987

☒ 70% ethanol

☒ DNase-free water

☒ AmpureXP PCR cleanup kits Beckman Coulter Catalog #A63880

☒ QIAquick gel extraction kit Qiagen Catalog #28704

☒ Qubit dsDNA BR Assay kit Invitrogen - Thermo Fisher Catalog #Q32850

Prepare Primer Mix (Optional, only for multiplexed Primer ID library prep)

1 For multiplexing sequencing, first, prepare Primer Mix.

Example (For HIV drug resistance pipeline).

	Regions	DR cDNA primer	DR F primer
	PR	R2614_PID	F2163AD
	RT	R3284_PID11	F2620_AD
	INT	R4752_PID11	F4383_AD
	V3	V3R_Buni_11	V1F_AD

Make 10 µM primer mix: mix 10 µL of each primer in one set and 60 µL of dH₂O.

Primer Tables

	primer	sequence (5'-3')
	R2614_P_ID	GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNNNNCAGTTAA CTTTGGGCCATCCATTCC
	R3284_P_ID11	GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNNNNNCAGTC CTATAAGGCTGTACTGTCCATTATC
	R4752_P_ID11	GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNNNNNATCGAA TACTGCCATTGTACTGC
	R7209_P_ID11	GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNNNNNCAGTC CATTGCTYAYTRABVTTACAATRTGC
	F2163AD	GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNNNTCAGAGCAG ACCAGAGCCAACAGCCCCA
	F2620_A_D	GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNNNGGCCATTGA CAGAAGAAAAAAATAAAAGC
	F4383_A_D	GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNNNAAAAGGAGA AGCCATGCATG
	V1F_AD	GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNNNTTATGGGATC AAAGCCTAAAGCCATGTGA

"N" in this Primer table is a random nucleotide.

P1	AATGATACGGCGACCACCGAGATCTACACGCCCTCCCTCGCGCCATCA GAGATGTG
Indexed Adapter	CAAGCAGAACAGGGCATACGAGATNNNNNNGTGACTGGAGTTAGAC GTGTGCTC
ADPT_2a	GTGACTGGAGTTCAGACGTGTGCTC

The 6 "N"s are not random nucleotides. They are a set of 24 pre-designed indexed sequences.

2 Mix well.

cDNA synthesis

3 Pipette the following components into a 0.5 ml RNase-free tube:

	$\mu\text{l}/\text{tube}$		[stock]	[final]	[mastermix]
	3.0	dNTP Mix	10 mM each	0.5	
	1.5	cDNA primer	10 μM	0.25 μM	
	34.5	RNA template			
	39.00	Total volume			

pipette 3 μL dNTPMix

pipette 1.5 μL cDNA primer

pipette 34.5 μL RNA template

4 Place tube in 65°C heat block for 3-5'.

timer 00:05:00 65°C heat block

5 Place the tube on ice for 1'.

timer 00:01:00 on ice

6 Add the following components:

	$\mu\text{l}/\text{tube}$		[stock]	[final]	[mastermix]
	12.0	5x buffer	5x	1x	
	3.0	DTT	100 mM	5	
	3.0	RNaseOUT	40 μL	2	
	3.0	SSIII RT	200 μL	10	
	21.0	Per tube			

pipette 12 μL 5x buffer

pipette 3 μL DTT

pipette 3 μL RNaseOUT

pipette 3 μL SSII RT

7 Mix and incubate at 50°C for 1 hr.

timer 01:00:00 Incubation at 50°C

- 8 Increase to 55°C and incubate for 1 hr.
 01:00:00 Incubation at 55°C
- 9 Inactivate SSIII RT by heating at 70°C for 15'.
 00:15:00 Heating at 70°C
- 10 To each tube, add 1 µL RNase H, incubate at 37°C for 20'.
 1 µL RNase H
 00:20:00 Incubation at 37°C

Purification

- 11 Purify cDNA using Agencourt RNAClean XP.
Resuspend the beads and take an aliquot out. Keep at room temperature for at least **30 minutes** before use. (Should be in 1ml aliquots)
 00:30:00 Room temperature
- 12 Transfer the cDNA reactions into 1.7 mL RNase-free tubes.
- 13 Resuspend the beads (Vortex). Add **42 µL of beads to 60 µL cDNA (Ratio: 0.6 – 0.8)** Agencourt RNAClean XP beads to each cDNA reaction.
 42 µL beads
 60 µL cDNA
- 14 Mix the Agencourt RNAClean XP and sample thoroughly by pipette mixing 15 times. No **vortexing**. Let the tube incubate at room temperature for **20 minutes** before proceeding to the next step.
 00:20:00 Room temperature
- 15 Place the tube onto the magnetic tube rack for **5 minutes** to separate the beads from solution.
 00:05:00 Magnetic tube rack
- 16 Slowly aspirate the cleared solution from the tube and discard. This step should be performed while the tube is situated on the rack. Do not disturb the magnetic beads, which have formed a spot on the side of the tube.
- 17 Dispense 500 µL of 70% ethanol into the tube and incubate for **30 seconds** at room temperature. Aspirate out the ethanol and discard. (1/3)

It is important to perform this step with the tube situated **on the rack. Do not disturb the separated magnetic beads.** Be sure to remove all of the ethanol from the bottom of the well as it may contain residual contaminants.

00:00:30 Incubation at room temperature

- 18 Dispense 500 µL of 70% ethanol into the tube and incubate for **30 seconds** at room temperature. Aspirate out the ethanol and discard. (2/3)

It is important to perform this step with the tube situated **on the rack. Do not disturb the separated magnetic beads.** Be sure to remove all of the ethanol from the bottom of the well as it may contain residual contaminants.

00:00:30 Incubation at room temperature

- 19 Let the reaction tube air-dry **10 minutes** on the rack with the cap open. The tube(s) should air-dry until the last visible traces of ethanol evaporate. Over drying the sample may result in a lower recovery.

00:10:00 Air-drying

- 20 **(Optional for clinical samples)** Elute in 60 µl water. Repeat step 12 to 19.

go to step #13 Optional for clinical samples

- 21 Remove the tube from the rack and resuspend beads in 24 µl DNase-free water by pipetting up and down. Place tube back on the rack and leave for **3 minutes**.

24 µL DNase-free water

00:03:00 Magnetic rack

- 22 Pipette the eluant from the tube while it is situated **on the magnetic tube rack.**

PCR 1

- 23 Complete thaw and vortex KAPA reagents (except for enzyme) before use.

24

	µl/tube		[stock]	[final]	[mastermix]
	10.0	5x Buffer A	5x	1x	
	10.0	Enhancer	5x	1x	
	1.0	dNTPs	10 mM	0.2 mM	
	2.5	Forward primer	10 µM	0.5 µM	
	2.5	ADPT_2a	10 µM	0.5 µM	
	0.5	KAPA Robust polymerase	5 U/µl	2.5 U	

	23.5	Template cDNA			
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Prepare mastermix in cold box and use repeater pipette to add to each tube in the cold box. Add template cDNA to each tube and pipette up and down to mix.

- pipette 10 µL 5x Buffer A
- pipette 10 µL Enhancer
- pipette 1 µL dNTPs
- pipette 2.5 µL Forward primer
- pipette 2.5 µL ADPT_2a
- pipette 0.5 µL KAPA Robust polymerase
- pipette 23.5 µL Template cDNA

25 Cycle

95°C	1 min
95°C	15 s
58°C	1 min
72°C	30 s
15-25 cycles	
72°C	3 min
4°C	On hold

Purification PCR products

- 26 Purify PCR products using AmpureXP PCR cleanup kits.
Vortex the 1 ml aliquot and remove the needed volume. Keep at room temperature for at least **30 minutes** before use.
 00:30:00 Room temperature
- 27 Transfer the PCR1 reactions into 1.7 mL RNase-free tubes.
- 28 Resuspend the beads. Add **40 µl (Ratio: 0.6 – 0.8: 1, 36µl – 48µl)** Ampure XP beads to each cDNA reaction.
 40 µL Ampure XP beads
- 29 Mix the Ampure XP and sample thoroughly by **vortexing**. Let the tube incubate at room temperature for **5 minutes** before proceeding to the next step (incubate off the rack).
 00:05:00 Incubation at room temperature

- 30 Place the tube onto the magnetic tube rack for **5 minutes** to separate the beads from solution.
 00:05:00 Incubation magnetic rack
- 31 Slowly aspirate the cleared solution from the tube and discard. This step should be performed while the tube is situated on the rack. Do not disturb the magnetic beads, which have formed a spot on the side of the tube.
- 32 Dispense 500 µL of 70% ethanol into the tube and incubate for **30 seconds** at room temperature. Aspirate out the ethanol and discard. (1/2)

Note

It is important to perform this step with the tube situated **on the rack**. **Do not disturb the separated magnetic beads**. Be sure to remove all of the ethanol from the bottom of the well as it may contain residual contaminants.



- 33 Dispense 500 µL of 70% ethanol into the tube and incubate for **30 seconds** at room temperature. Aspirate out the ethanol and discard. (2/2)

Note

It is important to perform this step with the tube situated **on the rack**. **Do not disturb the separated magnetic beads**. Be sure to remove all of the ethanol from the bottom of the well as it may contain residual contaminants.



- 34 Let the reaction tube air-dry **10 minutes** on the rack with the cap open. The tube(s) should air-dry until the last visible traces of ethanol evaporate. Over drying the sample may result in a lower recovery.
 00:10:00 Air-drying
- 35 Remove the tube from the rack and resuspend beads in 50 µL DNase-free water by pipetting up and down. Place tube back on the rack and leave for **3 minutes**.
 50 µL DNase-free water
 00:03:00 Magnetic rack
- 36 Pipette the 45 µL eluant from the tube while it is situated **on the magnetic tube rack**.

PCR 2

37 Complete thaw and vortex KAPA reagents (except for enzyme) before use.

38

µl/tube		[stock]	[final]	[mastermix]
5.0	5x KAPA HiFi Fidelity Buffer	5x	1x	
1.0	dNTP Mix	10 mM	0.4 mM	
1.0	Uni Adapter (ADPT_P1)	10 µM	0.4 µM	
0.5	KAPA HiFi polymerase	1 U/µl	0.5 U	
1.0	Indexed Adapter	10 µM	0.4 µM	
2.0	Purified template DNA			
14.5	Water			

Prepare mastermix in cold box and use repeater pipette to add to each tube in the cold box. Add Indexed Adapter to each tube.

 5 µL 5x KAPA HiFi Fidelity Buffer

 1 µL dNTP Mix

 1 µL Uni Adapter (ADPT_P1)

 0.5 µL KAPA HiFi polymerase

 1 µL Indexed Adapter

 2 µL Purified template DNA

 14.5 µL Water

39 **Cycle** (PCR machine #5 SZ → ILM2):

95°C	2 min
98°C	20 s
63°C	15 s
72°C	30 s
25 - 35 cycles	
72°C	3 min
4°C	On hold

Gel Purification and quantification

- 40 Before gel purification, run 2 µl products on 1% agarose gel to check the bands.
- 41 **Gel purification.** (Qiagen QIAquick gel extraction kit)
Run 2nd round PCR products on 1.2% agarose gel. E = 4 V/cm, T = 60 min.
- 42 Excise DNA fragment.
- 43 Weight the gel; add 3 volume of Buffer QG to 1 volume of gel.
- 44 Incubate at 50 °C for **10 minutes** to completing dissolve. Vortex every **2-3 minutes** to help dissolve.
 00:10:00 Incubation at 50 °C
 00:02:30 Vortex
- 45 Check the color of gel solution (should be yellow, otherwise add 10 µl 3M sodium acetate).
- 46 Place **MinElute** column, apply the sample to the column and centrifuge for **1 minute**.
 00:01:00 Centrifugation
- 47 Add 500 µl buffer QG and centrifuge for **1 minute**.
 500 µL Buffer QG
 00:01:00 Centrifugation
- 48 Add 0.75 ml buffer PE, **incubate for 5 minutes** at room temperature, centrifuge for **1 minute**.
 00:05:00 Incubation at room temperature
 00:01:00 Centrifugation
- 49 Discard the fluid, centrifuge for additional **3 minutes**.
 00:03:00 Centrifugation
- 50 Put the column in a new 1.7 ml tube, add **10 µl** buffer EB. Stand for **4 minutes**, centrifuge for **2 minutes**.
 00:04:00 Stand

- 51 Quantification using Invitrogen Qubit dsDNA BR Assay kit. See Qubit dsDNA BR assay protocol. **Don't use Nanodrop to quantify!**
- 52 After quantification, pool libraries in equal amount. Use AMPure XP beads to purify pooled libraries (2:3, two washes), elute in 20 to 30 µL Elution Buffer.
- 53 Quantify the pooled library using Qubit dsDNA BR assay kit. Check the quality of the library by Agilent Bioanalyzer or Bio-rad Experion. Repeat purification if primer dimmer is present (~200bp).