

Primer ID MiSeq Library Prep Version 4

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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.

TAGS

Library prep

NGS

Show tags

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PROTOCOL STATUS

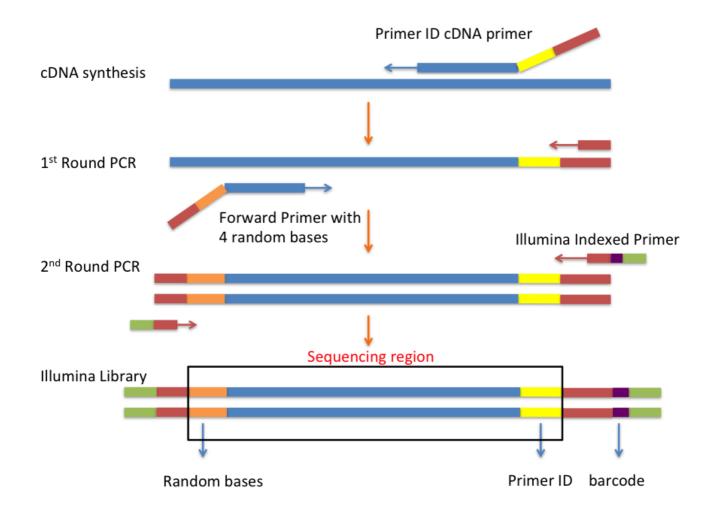
Working

We use this protocol in our group and it is working

GUIDELINES

Viral RNA extraction using QIAamp vRNA mini kit.





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

| | GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNNNNTTATGGGATC |
|--------------------|---|
| V1F(forward) 5'-3' | AAAGCCTAAAGCCATGTGTA |
| BV3R Uni (cDNA | GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNNNNNCAGTCCATTT |
| Primer) 5'-3' | TGCTCTACTAATGTTACAATGTGC |
| | AATGATACGGCGACCACCGAGATCTACACGCCTCCCTCGCGCCATCAGAGATGT |
| Universal Adapter | G |
| | CAAGCAGAAGACGGCATACGAGAT NNNNNNGTGACTGGAGTTCAGACGTGT |
| Indexed Adapter | GCTC |
| ADPT_2a | GTGACTGGAGTTCAGACGTGTGCTC |

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

Table of Indexed Primers

| | | Index | |
|----------------------|-------|----------|--|
| Indexed Primer | Index | Sequence | Sequence |
| PCR Primer, Index 1 | 1 | ATCACGA | CAAGCAGAAGACGCCATACGAGATCGTGATGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 2 | 2 | CGATGTA | CAAGCAGAAGACGCCATACGAGAT <mark>ACATCG</mark> GTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 3 | 3 | TTAGGCA | CAAGCAGAAGACGCCTACGAGATGCCTAAGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 4 | 4 | TGACCAA | CAAGCAGAAGACGCCATACGAGATTGGTCAGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 5 | 5 | ACAGTGA | CAAGCAGAAGACGGCATACGAGATCACTGTGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 6 | 6 | GCCAATA | CAAGCAGAAGACGCCATACGAGATATTGGCCGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 7 | 7 | CAGATCA | CAAGCAGAAGACGCATACGAGATGATCTGGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 8 | 8 | ACTTGAA | CAAGCAGAAGACGCATACGAGATTCAAGTGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 9 | 9 | GATCAGA | CAAGCAGAAGACGCATACGAGATCTGATCGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 10 | 10 | TAGCTTA | CAAGCAGAAGACGGCATACGAGATAAGCTAGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 11 | 11 | GGCTACA | CAAGCAGAAGACGCCATACGAGATGTAGCCCGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 12 | 12 | CTTGTAA | CAAGCAGAAGACGGCATACGAGATTACAAGGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 13 | 13 | TCCATAA | CAAGCAGAAGACGGCATACGAGATTATGGAGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 14 | 14 | GTACTAA | CAAGCAGAAGACGCATACGAGATTAGTACGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 15 | 15 | ACAGTAA | CAAGCAGAAGACGCATACGAGATTACTGTGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 16 | 16 | CTCATGA | CAAGCAGAAGACGCATACGAGATCATGAGGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 17 | 17 | ACGATAA | CAAGCAGAAGACGCATACGAGATTATCGTGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 18 | 18 | TGCAGAA | CAAGCAGAAGACGGCATACGAGATTCTGCAGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 19 | 19 | TTCATAA | CAAGCAGAAGACGGCATACGAGATTATGAAGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 20 | 20 | TGCTGTA | CAAGCAGAAGACGGCATACGAGATACAGCAGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 21 | 21 | TATCACA | CAAGCAGAAGACGCCATACGAGATGTGATAGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 22 | 22 | TGGATAA | CAAGCAGAAGACGCCATACGAGATTATCCAGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 23 | 23 | CGCATTA | CAAGCAGAAGACGGCATACGAGATAATGCGGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 24 | 24 | GCCTTAA | CAAGCAGAAGACGCCATACGAGATTAAGGCGTGACTGGAGTTCAGACGTGTGCTC |

MATERIALS

| NAME ~ | CATALOG # | VENDOR |
|-----------------------------|-----------|-------------------------------|
| QIAamp vRNA mini kit | 51304 | Qiagen |
| Agencourt RNAClean XP | A63987 | Beckman Coulter |
| 70% ethanol | | Contributed by users |
| DNase-free water | | Contributed by users |
| AmpureXP PCR cleanup kits | A63880 | Beckman Coulter |
| QIAquick gel extraction kit | 28704 | Qiagen |
| Qubit dsDNA BR Assay kit | Q32850 | Invitrogen - Thermo Fisher |

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 |
| IN | R4752_PID11 | F4383_AD |
| V3 | R7209_PID11 | 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_PID | GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNN |
| | NNNNCAGTTTAACTTTTGGGCCATCCATTCC |
| R3284_PID11 | GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNN |
| | NNNNNCAGTCACTATAGGCTGTACTGTCCATTTATC |
| R4752_PID11 | GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNN |
| | NNNNNATCGAATACTGCCATTTGTACTGC |
| R7209_PID11 | GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNN |
| | NNNNNCAGTCCATTTTGCTYTAYTRABVTTACAATRTGC |
| F2163AD | GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNN |
| | NNTCAGAGCAGACCAACAGCCCCA |
| F2620_AD | GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNN |
| | NNGGCCATTGACAGAAGAAAAATAAAAGC |
| F4383_AD | GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNN |
| | NNAAAAGGAGAAGCCATGCATG |
| V1F_AD | GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNN |
| | NNTTATGGGAT CAAAGCCT AAAGCCATGTGTA |

[&]quot;N" in this Primer table is a random nucleotide.

| P1 | AAT GAT ACGGCGACCACCGAGAT CT ACACGCCT CCCT C |
|-----------------|--|
| | GCCATCAGAGATGTG |
| Indexed Adapter | CAAGCAGAAGACGGCATACGAGAT NNNNN GTGACTGG |
| | AGTTCAGACGTGTGCTC |
| 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:

| μl/tube | | [stock] | [final] | [mastermix] |
|---------|--------------|------------|---------|-------------|
| 3.0 | dNTP Mix | 10 mM each | 0.5 | |
| 1.5 | cDNA primer | 10 μΜ | 0.25 μΜ | |
| 34.5 | RNA template | | | |
| 39.00 | Total volume | | | |





■34.5 µl RNA template

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

© 00:05:00 65°C heat block

5 Place the tube on ice for 1'.

© 00:01:00 on ice

6 Add the following components:

| μl/tube | | [stock] | [final] | [mastermix] |
|---------|-----------|----------|---------|-------------|
| 12.0 | 5x buffer | 5x | 1x | |
| 3.0 | DTT | 100 mM | 5 | |
| 3.0 | RNaseOUT | 40 u/μl | 2 | |
| 3.0 | SSIII RT | 200 u/μl | 10 | |
| 21.0 | Pertube | | | |



3 μl SSII RT

7 Mix and incubate at 50°C for 1 hr.

© 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 μ l 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.



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.

- 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

Dispense 500 μ L 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

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.

goto can't be rendered without protocol or steps

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.

PCR₁

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 μΜ | 0.5 μΜ | |
| 2.5 | ADPT_2a | 10 μΜ | 0.5 μΜ | |
| 0.5 | KAPA Robust polymerase | 5 U/μΙ | 2.5 U | |
| 23.5 | Template cDNA | | | |

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.



25 Cycle

| 95°C | 1 min |
|--------------|---------|
| | |
| 95°C | 15 s |
| 58°C 72°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
```

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 (incube off the rack).

```
\bigcirc 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
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- 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 μ L 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.

□500 µl 70% ethanol

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

ENOTE

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.

□500 µl 70% ethanol

♦ 00:00:30 Incubation at room temperature

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 34 ethanol evaporate. Over drying the sample may result in a lower recovery.

©00:10:00 Air-drying

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 35 rack and leave for 3 minutes.

□50 µl DNase-free water © 00:03:00 Magnetic rack

Pipette the 45 μ l eluant from the tube while it is situated **on the magnetic tube rack**. 36

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 μΜ | 0.4 μΜ | |
| 0.5 | KAPA HiFI polymerase | 1 U/µl | 0.5 U | |
| 1.0 | Indexed Adapter | 10 μΜ | 0.4 μΜ | |
| 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

8



39 Cycle (PCR machine #5 SZ -> ILM2):

| 95°C | 2 min |
|----------------|---------|
| | |
| 98°C | 20 s |
| 63°C 72°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.
- 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 μ l 3M sodium acetate).
- 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.

७00:03:00 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

- Quantification using Invitrogen Qubit dsDNA BR Assay kit. See Qubit dsDNA BR assay protocol. **Don't use Nanodrop to quantify!**
- 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.
- 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).

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