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## Primer ID MiSeq Library Prep for HIV-1 DR and diversity

Forked from [Primer ID MiSeq Library Prep](#)

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1 Works for me [dx.doi.org/10.17504/protocols.io.6hchb2w](https://doi.org/10.17504/protocols.io.6hchb2w)

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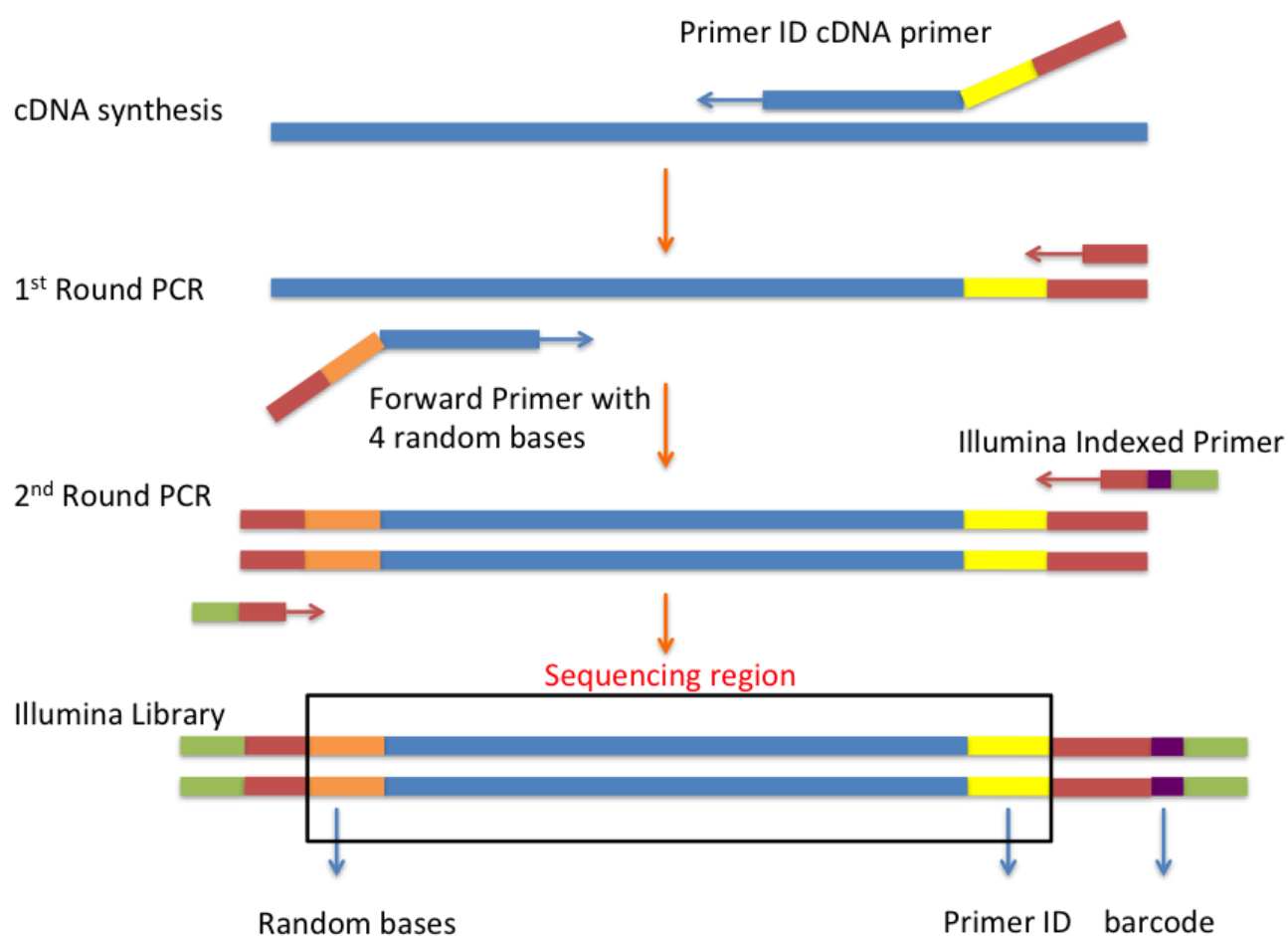
### ABSTRACT

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.

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### 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)**

|                              |   |
|------------------------------|---|
| V1F(forward) 5'-3'           | GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNNNNTTATGGGATC<br>AAAGCCTAAAGCCATGTGTA      |
| BV3R Uni (cDNA Primer) 5'-3' | GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNNNNNNCAGTCCATT<br>TGCTCTACTAATGTTACAATGTGC |
| Universal Adapter            | AATGATACGGCGACCACCGAGATCTACAGCCTCCCTCGCGCCATCAGAGATGT<br>G                        |
| Indexed Adapter              | CAAGCAGAAGACGGCATACGAGATNNNNNNGTGACTGGAGTTCAGACGTGT<br>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        | CAAGCAGAAGACGGCATACGAGATCGTGATGTGACTGGAGTTCAGACGTGTGCTC    |
| PCR Primer, Index 2  | 2     | CGATGTA        | CAAGCAGAAGACGGCATACGAGATACATCGGTGACTGGAGTTCAGACGTGTGCTC    |
| PCR Primer, Index 3  | 3     | TTAGGCA        | CAAGCAGAAGACGGCATACGAGATGCCTAAGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 4  | 4     | TGACCAA        | CAAGCAGAAGACGGCATACGAGATGGGTCAAGTGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 5  | 5     | ACAGTGA        | CAAGCAGAAGACGGCATACGAGATCACTGTTGTGACTGGAGTTCAGACGTGTGCTC   |
| PCR Primer, Index 6  | 6     | GCCAATA        | CAAGCAGAAGACGGCATACGAGATATTGGCGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 7  | 7     | CAGATCA        | CAAGCAGAAGACGGCATACGAGATGATCTGTGTGACTGGAGTTCAGACGTGTGCTC   |
| PCR Primer, Index 8  | 8     | ACTTGAA        | CAAGCAGAAGACGGCATACGAGATCAAGTGTGTGACTGGAGTTCAGACGTGTGCTC   |
| PCR Primer, Index 9  | 9     | GATCAGA        | CAAGCAGAAGACGGCATACGAGATCTGATCGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 10 | 10    | TAGCTTA        | CAAGCAGAAGACGGCATACGAGATAAGCTAGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 11 | 11    | GGCTACA        | CAAGCAGAAGACGGCATACGAGATGTAGCCGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 12 | 12    | CTTGTA         | CAAGCAGAAGACGGCATACGAGATTACAAGGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 13 | 13    | TCCATAA        | CAAGCAGAAGACGGCATACGAGATTATGGAAGTGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 14 | 14    | GTAATAA        | CAAGCAGAAGACGGCATACGAGATTAGTACGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 15 | 15    | ACAGTAA        | CAAGCAGAAGACGGCATACGAGATTACTGTGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 16 | 16    | CTCATGA        | CAAGCAGAAGACGGCATACGAGATCATGAGGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 17 | 17    | ACGATAA        | CAAGCAGAAGACGGCATACGAGATTATCGTGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 18 | 18    | TGCAGAA        | CAAGCAGAAGACGGCATACGAGATTCTGCAAGTGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 19 | 19    | TTCATAA        | CAAGCAGAAGACGGCATACGAGATTATGAAGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 20 | 20    | TGCTGTA        | CAAGCAGAAGACGGCATACGAGATACAGCAAGTGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 21 | 21    | TATCACA        | CAAGCAGAAGACGGCATACGAGATGTGATAAGTGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 22 | 22    | TGGATAA        | CAAGCAGAAGACGGCATACGAGATTATCCAAGTGTGACTGGAGTTCAGACGTGTGCTC |
| PCR Primer, Index 23 | 23    | CGCATTA        | CAAGCAGAAGACGGCATACGAGATAATGCGGTGTGACTGGAGTTCAGACGTGTGCTC  |
| PCR Primer, Index 24 | 24    | GCCTTAA        | CAAGCAGAAGACGGCATACGAGATTAAGGCGTGTGACTGGAGTTCAGACGTGTGCTC  |

MATERIALS

NAME

QIAamp vRNA mini kit

CATALOG #

51304

VENDOR

Qiagen

| NAME ▾                      | CATALOG # ▾ | VENDOR ▾                   |
|-----------------------------|-------------|----------------------------|
| Agencourt RNAClean XP       | A63987      | Beckman Coulter            |
| 70% ethanol                 |             |                            |
| DNase-free water            |             |                            |
| 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  $\mu$ M primer mix: mix 10  $\mu$ L of each primer in one set and 60  $\mu$ L of dH<sub>2</sub>O.

#### Primer Tables

| primer      | sequence (5'-3')   |
|-------------|--|
| R2614_PID   | GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNN<br>NNNCAGTTAACTTTTGGGCCATCCATTCC          |
| R3284_PID11 | GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNN<br>NNNNNCAGTCACTATAGGCTGTACTGTCCATTTATC   |
| R4752_PID11 | GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNN<br>NNNNNATCGAATACTGCCATTTGTACTGC          |
| R7209_PID11 | GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNN<br>NNNNNCAGTCCATTTTCTYTAYTRABVTTACAATRTGC |
| F2163AD     | GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNNN<br>NTCAGAGCAGACCAGAGCCAACAGCCCCA         |
| F2620_AD    | GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNNN<br>NGGCCATTGACAGAAGAAAAAATAAAGC          |
| F4383_AD    | GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNNN<br>NAAAAGGAGAAGCCATGCATG                 |
| V1F_AD      | GCCTCCCTCGCGCCATCAGAGATGTGTATAAGAGACAGNNN<br>NTTATGGGATCAAAGCCTAAAGCCATGTGTA       |

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

|                 |   |
|-----------------|---|
| P1              | AATGATACGGCGACCACCGAGATCTACACGCTCCCTCGCG<br>CCATCAGAGATGTG  |
| Indexed Adapter | CAAGCAGAAGACGGCATACGAGATNNNNNNGTGACTGGA<br>GTTTACAGCTGTGCTC |
| 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] |
|---------|--------------|------------|---------|-------------|
| 2.0     | dNTP Mix     | 10 mM each | 0.5     |             |
| 1.0     | cDNA primer  | 10 µM      | 0.25 µM |             |
| 23.0*   | RNA template |            |         |             |
| 26.00   | Total volume |            |         |             |

\* The input volume of vRNA depends on the viral loads. We recommend 1,000 to 20,000 copies, but it can work with samples as little as 200 copies. If copy number unknown, use the maximum volume of 23 µl.

📄 2 µl dNTP Mix

📄 1 µl cDNA primer

📄 23 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] |
|---------|-----------|----------|---------|-------------|
| 8.0     | 5x buffer | 5x       | 1x      |             |
| 2.0     | DTT       | 100 mM   | 5       |             |
| 2.0     | RNaseOUT  | 40 u/µl  | 2       |             |
| 2.0     | SSIII RT  | 200 u/µl | 10      |             |
| 14.0    | Per tube  |          |         |             |

📄 8 µl 5x buffer

📄 2 µl DTT

📄 2 µl RNaseOUT

📄 2 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 µl RNase H, incubate at 37°C for 20'.

📄 0.5 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 **28 µl of beads to 40 µl cDNA** Agencourt RNAClean XP beads to each cDNA reaction.

📄 28 µl beads

📄 40 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 400 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 400 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 Remove the tube from the rack and resuspend beads in 40 µl DNase-free water by pipetting up and down, Place tube back on the rack and leave for **3 minutes**.

📄 40 µl DNase-free water

🕒 00:03:00 Magnetic rack

- 21 Pipette the eluant from the tube while it is situated **on the magnetic tube rack**. Transfer the eluant into the tube with 28 µl of RNACleanup XP beads.
- 22 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**
- 23 Pipette the eluant from the tube while it is situated **on the magnetic tube rack**.

## PCR 1

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

25

| µl/tube |                        | [stock] | [final] | [mastermix] |
|---------|------------------------|---------|---------|-------------|
| 10.0    | 5x <b>Buffer A</b>     | 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          |         |         |             |

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.

 **10 µl 5x Buffer A**

 **10 µl Enhancer**

 **1 µl dNTPs**

 **2.5 µl Forward primer**

 **2.5 µl ADPT\_2a**

 **0.5 µl KAPA Robust polymerase**

 **23.5 µl Template cDNA**

- 26 **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

- 27 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**
- 28 Transfer the PCR1 reactions into 1.7 mL RNase-free tubes.
- 29 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**
- 30 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).  
🕒 **00:05:00 Incubation at room temperature**
- 31 Place the tube onto the magnetic tube rack for **5 minutes** to separate the beads from solution.  
🕒 **00:05:00 Incubation magnetic rack**
- 32 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.
- 33 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)



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**

- 34 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)



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**

- 35 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**



- 36 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

- 37 Pipette the 45 µl eluant from the tube while it is situated **on the magnetic tube rack**.

## PCR 2

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

39

| 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

- 40 **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

- 41 Before gel purification, run 2 µl products on 1% agarose gel to check the bands.



- 42 **Gel purification.** (Qiagen QIAquick gel extraction kit)  
Run 2<sup>nd</sup> round PCR products on 1.2% agarose gel. E = 4 V/cm, T = 60 min.
- 43 Excise DNA fragment.
- 44 Weight the gel; add 3 volume of Buffer QG to 1 volume of gel.
- 45 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**
- 46 Check the color of gel solution (should be yellow, otherwise add 10 µl 3M sodium acetate).
- 47 Place **MinElute** column, apply the sample to the column and centrifuge for **1 minute**.  
🕒 **00:01:00 Centrifugation**
- 48 Add 500 µl buffer QG and centrifuge for **1 minute**.  
📄 **500 µl Buffer QG**  
🕒 **00:01:00 Centrifugation**
- 49 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**
- 50 Discard the fluid, centrifuge for additional **3 minutes**.  
🕒 **00:03:00 Centrifugation**
- 51 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**
- 52 Quantification using Invitrogen Qubit dsDNA BR Assay kit. See Qubit dsDNA BR assay protocol. **Don't use Nanodrop to quantify!**
- 53 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.
- 54 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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