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# One-pot native barcoding of amplicons v2

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1 more workspace



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# OPEN ACCESS



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External link: <a href="http://lab.loman.net/protocols/">http://lab.loman.net/protocols/</a>

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We use this protocol and it's working

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# **Abstract**

This one-pot native barcoding protocol was developed in conjunction with Oxford Nanopore Technologies, New England Biolabs and BCCDC.

# **Attachments**



One-pot native barco...

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# Safety warnings



See SDS (Safety Data Sheet) for safety warnings and hazards.



1 Set up the following reaction for each sample:

# Component PCR dilution from previous step Nuclease-free water Ultra II End Prep Reaction Buffer Ultra II End Prep Enzyme Mix Total Volume Δ 5 μL Δ 7.5 μL Δ 1.75 μL Δ 1.75 μL

- Incubate at room temperature for 00:10:00

  Incubate at 65 °C for 00:10:00

  Incubate on ice for 00:01:00
- In a new 1.5mL Eppendorf tube set up the following reaction:

# Component Volume

Previous reaction mixture

Nuclease-free water

NBXX barcode

Ultra II Ligation Master Mix

Ligation Enhancer

Total

1.5  $\mu$ L

2.5  $\mu$ L

3.0  $\mu$ L

## Note

Use one native barcode from the EXP-NBD104 (1-12) or EXP-NBD114 (13-24) per sample. Use from 6 to 24 barcodes in a library, any fewer and there will be insufficient total material to achieve good yields.

Incubate at room temperature for 00:20:00

Incubate at 65 °C for 00:10:00

Incubate on ice for 00:01:00



### Note

The 65°C incubation is to inactivate the DNA ligase to prevent barcode cross-ligation when reactions are pooled in the next step.

- In a new 1.5 ml Eppendorf tube pool all  $\stackrel{\triangle}{=}$  20  $\stackrel{\triangle}{=}$  one-pot barcoding reactions together.
- Add 0.4x volume of SPRI beads to the sample tube and mix gently by either flicking or pipetting. For example add  $496 \, \mu L$  SPRI beads to  $4240 \, \mu L$  12-plex pooled one-pot native barcoding reactions.

### Note

0.4x volume of SPRI will only bind 400 bp amplicons in the presence of ligation buffer as in a one-pot reaction, do not use 1x as this will result in excessive native barcode carryover.

- 7 Pulse centrifuge to collect all liquid at the bottom of the tube.
- 8 Incubate for 00:05:00 at room temperature.
- Place on magnetic rack and incubate for 00:02:00 or until the beads have pelleted and the supernatant is completely clear.
- 10 Carefully remove and discard the supernatant, being careful not to touch the bead pellet.
- 11 Add 700 µl SFB and resuspend beads completely by pipette mixing.

### Note

SFB will remove excess adapter without damaging the adapter-protein complexes. Do not use 70% ethanol as in early clean-ups.



- 12 Pulse centrifuge to collect all liquid at the bottom of the tube.
- 13 Remove supernatant and discard.
- 14 Repeat steps 11-13 to perform a second SFB wash.
- 15 Pulse centrifuge and remove any residual SFB.

Note

You do not need to allow to air dry with SFB washes.

- 16 Add 200  $\mu$ l of room-temperature 70 % volume ethanol to bathe the pellet.
- 17 Carefully remove and discard ethanol, being careful not to touch the bead pellet.

Note

Only perform 1x 70% ethanol wash

- 18 Pulse centrifuge to collect all liquid at the bottom of the tube and carefully remove as much residual ethanol as possible using a P10 pipette.
- 19 With the tube lid open incubate for 00:01:00 or until the pellet loses it's shine (if the pellet dries completely it will crack and become difficult to resuspend).
- 20 Resuspend pellet in 4 30 µL Elution Buffer (EB), mix gently by either flicking or pipetting and incubate for 00:02:00 .
- 21 Place on magnet and transfer sample to a clean 1.5mL Eppendorf tube ensuring no beads are transferred into this tube.