



APR 15, 2024

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**Protocol Citation:** Stephen Douglas Russell, Harte Singer 2024. FUNDIS ONT V14 Nanopore Adapter Ligation for Fungal DNA Barcoding Flongle 10.4.1. **protocols.io**  
<https://protocols.io/view/fundis-ont-v14-nanopore-adapter-ligation-for-funga-db4w2qxe>


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**Protocol status:** Working  
We use this protocol and it's working

**Created:** Apr 14, 2024

**Last Modified:** Apr 15, 2024

## FUNDIS ONT V14 Nanopore Adapter Ligation for Fungal DNA Barcoding Flongle 10.4.1

 Forked from [ONT V14 Nanopore Adapter Ligation for Fungal DNA Barcoding](#)

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### ABSTRACT

This process will take your A-tailed library and add the nanopore adapters. Simply combine several chemicals for a single reaction and do a bead cleanup.

#### Tested with:

Flowcells: Flongle 10.4.1

Ligation Kit: V14 - LSK114

Time required: ~45 minutes

Adapted from [dx.doi.org/10.17504/protocols.io.dm6gpb5zdlzp/v5](https://doi.org/10.17504/protocols.io.dm6gpb5zdlzp/v5)

PROTOCOL integer ID: 98166

**Keywords:** nanopore, fungi, flongle, fmol, library preparation

## MATERIALS

### Reagents



Ligation Sequencing Kit V14 **Oxford Nanopore Technologies Catalog #SQK-LSK114**

: \$694.43 per 6 reactions (\$115.74 per MinION run; \$57.87 per Flongle run)



NEBNext Quick Ligation Module **New England Biolabs Catalog #E6056S** :

\$361.00 per 20 reactions (\$18.05 per MinION run; \$9.03 per Flongle run)

\*note: This kit has two components. We use one - NEBNext Quick T4 DNA Ligase. NEB checking on whether the single one is available for purchase. Samples of this kit should be available from NEB.



HighPrep™ PCR Clean-up System **MagBio Genomics Inc. Catalog #AC-60005** :

\$117.88 per 50 mL. \$0.047 per rxn.

Note: Most magnetic beads from most vendors can be used with the same protocol.

Total per Flongle run (1/2 rxns): \$66.95

Total per MinION run: \$133.84

Total per sample (Flongle: 480 samples): \$0.139

Total per sample (Flongle: 960 samples): \$0.07

### Consumables

Eppendorf DNA LoBind 1.5mL tubes

10uL pipette tips

100-200uL pipette tips

### Equipment

PCR tube rack

Vortex mixer

Mini centrifuge

PCR cleanup magnet

10uL Pipette

100uL Pipette


Hula mixer (Ebay): \$200.00 (optional)

Quantus or Qubit Fluorometer (optional)

PROTOCOL MATERIALS

 Ligation Sequencing Kit V14 **Oxford Nanopore Technologies Catalog #SQK-LSK114**

In 2 steps


 NEBNext Quick Ligation Module **New England Biolabs Catalog #E6056S** Step 1

## Adapter Ligation Flongle 10.4.1

- 1 You should already have your reagents on ice from the dA tailing step, but if not you will need the NEBNext Quick Ligation Module enzyme and from the ONT kit you will need 1 tube of SFB, and LA, LNB, EB, and AXP. Spin down the NEB enzyme and immediately place on ice. Thaw AXP and allow to come up to room temperature, do not place on ice. Thaw other reagents at room temperature, flick-mix, spin down, and keep on ice.

LNB, LA, AXP, SFB, EB -


 Ligation Sequencing Kit V14 **Oxford Nanopore Technologies Catalog #SQK-LSK114**

Quick T4 Ligase -  NEBNext Quick Ligation Module **New England Biolabs Catalog #E6056S**


- 2 In a 1.5 mL DNA LoBind tube, mix in the following order:

Between each addition, pipette mix 10-20 times.

| Reagent                     | Volume  |
|-----------------------------|---------|
| dA Tailed DNA               | 30 µl   |
| Ligation Buffer (LNB)       | 12.5 µl |
| NEBNext Quick T4 DNA Ligase | 5 µl    |
| Ligation Adapter (LA)       | 2.5 µl  |
| Total                       | 50 µl   |


- 3 Spin down with a mini centrifuge for  00:00:02 .


2s



- 4 Incubate the reaction for  00:10:00 at room temperature. This is a good time to do a flow cell check and get your computer set up.

10m




5 Resuspend AMPure XP (AXP) magnetic bead stock by vortexing.

6 Add  20  $\mu\text{L}$  of resuspended beads to the reaction and mix by flicking the tube.


7 Incubate for  00:05:00 at room temperature, ONT suggests using a Hula mixer, you can just gently flick the tube a few times. 5m


8 Spin down the sample for  00:00:02 and pellet on a magnet for  00:02:00 . 2m 2s





Keep the tube on the magnet, and pipette off the supernatant.

9 Wash the beads by adding  125  $\mu\text{L}$  of Short Fragment Buffer (SFB). Flick the beads vigorously to resuspend (they will only partially resuspend), spin down for  00:00:02 , then return the tube to the magnetic rack for  00:02:00 and allow the beads to pellet. Remove the supernatant using a pipette and discard. 2m 2s

SFB -  Ligation Sequencing Kit V14 **Oxford Nanopore Technologies Catalog #SQK-LSK114**


10 Repeat the previous step. 

11 Spin down for  00:00:02 and place the tube back on the magnet. Pipette off any residual supernatant. 2s  
Allow to dry for ~30 seconds, but do not dry the pellet to the point of cracking.

- 12 Remove the tube from the magnetic rack and resuspend the pellet in  7  $\mu$ L Elution Buffer (EB). Incubate for  00:10:00 at room temperature. Use this time to put away all of the reagents from the previous steps and set up your reagents for the final library mixing step. Use the glass vials that come with the Flongle kit, you will need FCF (Flow Cell Flush) SB (Sequencing Buffer) LIB (Library Beads). You will also need the FLT (Flow Cell Tether) From the main Ligation Sequencing kit. Make sure you have your Qubit reagents ready for quantification. 10m
- 13 Pellet the beads on a magnet until the eluate is clear and colorless, for at least  00:01:00 . 1m
- 14 Remove and retain  7  $\mu$ L of eluate containing the DNA library into a clean 1.5 ml Eppendorf DNA LoBind tube.

Store on ice until you are ready to load in your flowcell.

## Quantification

- 15 Quantify  2  $\mu$ L of your ligated library using the Qubit fluorimeter

It is recommend loading 5 fmol to 10 fmol of this final prepared library onto your flow cells. Loading more than 20 fmol of DNA can reduce the rate of duplex read capture. ONT recommends that you dilute the library in Elution Buffer if required, however we use molecular grade water instead. We dilute the entire 5uL of library, however you may dilute a smaller amount if you choose.

<https://www.promega.com/resources/tools/biomath/>

For 900bp length DNA (what our ITS1F-4 rxns appear to average, with adapters), we are looking for:  
10 fmol - 20 fmol = 6ng - 12ng of DNA in 5uL of final dilute library.

For a 22 ng/uL sample

22ng/uL \* 5uL = 110ng DNA in sample of 5uL elution buffer.

How much additional molecular water to have 5uL needed for the next step give us correct amount of DNA?

$C1V1 = C2V2$ , rearrange  $C1V1/C2 = V2$

$(110\text{ng} * 5\text{uL}) / 10\text{ng}(17 \text{ fmol DNA}) = 55\text{uL total volume}$

Bring initial volume (5uL) up to 55uL = add 50uL molecular water

So at 22ng/uL quantification, add an additional 50uL of molecular water to have right concentration to use 5uL for the next step with Flongle.