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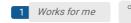


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# Ultra-Rapid Sequencing (PCR)

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#### ABSTRACT

This protocol accompanies the paper "Ultra-Rapid Somatic Variant Detection via Real-Time Threshold Sequencing." This protocol was followed to initiate a sequencing run that resulted in a somatic variant call from known tumor tissue in  $\sim$ 52 minutes. The protocol outlines DNA extraction, PCR, library preparation for Oxford Nanopore Sequencing, and sequencer preparation and loading as executed for the 52 minute demonstration.

#### PROTOCOL CITATION

Jack Wadden 2021. Ultra-Rapid Sequencing (PCR). **protocols.io** https://protocols.io/view/ultra-rapid-sequencing-pcr-bs7bnhin

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## Preparation

- 1 Prepare the PCR reaction tube by mixing the following components in a **Q0.2 mL** PCR tube
  - 12.5 µl

፟ 🛛 🖎 Q5 Hot Start High-Fidelity 2X Master Mix - 500 rxns **New England** 

Biolabs Catalog #M0494L

- □1.25 μI forward primer
- **1.25** μl reverse primer
- **9 µl Nuclease-free Water Contributed by users**

```
Prepare the DNA extraction tube by mixing the following components in a 2 mL Eppendorf tube
                        ⊠ Epicentre QuickExtract™ DNA Extraction
        ■ 500 μl of Solution Epicentre Catalog #QE09050
                                                                                                          11m 18s
       Pre-heat thermocycler and pause for combined PCR and fragmentation protocol
       Pre-heat two heat blocks ( § 65 °C and § 98 °C )
       DNA Extraction (in heat blocks)
          8 65 °C for © 00:06:00
          8 98 °C for © 00:02:00
       PCR (in thermocycler)
          8 98 °C for © 00:00:30
         28 cycles of
            § 98 °C for © 00:00:05
            8 64 °C for © 00:00:05
            § 72 °C for ७00:00:08
          A 72 °C for © 00:00:30
         hold at § 20 °C
       Fragmentation (in thermocycler)
          § 30 °C for ७ 00:01:00
          8 80 °C for © 00:01:00
         hold at § 20 °C
DNA Extraction
       When acquired, place 20 µg tumor tissue into 2 mL DNA extraction tube
                                                                                                              10s
       Vortex on high for © 00:00:10
                                                                                                               8m
       Place in pre-heated heat block at 8 65 °C
        • briefly vortex after 3 minutes of incubation
       Place in pre-heated heat block at § 98 °C
       Remove tube, vortex briefly, spin down
                                                                                                              30s
   9
       Incubate on ice for © 00:00:30
```

12 Let thermocycler run through the PCR protocol 26m

Either let thermocycler draw DNA down to room temperature or quench in ice for **© 00:00:10**. Pause thermocycler program to pre-heat for ONT rapid library preparation tagmentation.

Library Preparation		
14	Add 7.5 µl PCR product to tagmentation mix tube, flick to mix, and spin down	
15	Place in thermocycler and unpause combined protocol	
16	Let thermocycler run through fragmentation protocol	
17	Either let thermocycler draw DNA down to room temperature or quench in ice for © 00:00:10	10s
18	Add 11 µl ONT Rapid Adapter to tagmentation tube, flick to mix, and spin down	
19	Incubate sequencing library for ③ 00:05:00 at room temperature, flicking to mix occasionally and spinnin	5m g down
	During the final 1m of incubation, re-prime the ONT flow-cell with SpotOn port open, using \( \text{\omega} \) FB (Flush Buffer) \( \text{Oxford Nanopore} \)	⊒200 µl
	of Technologies Catalog #EXP-FLP002 buffer	loading
Sequen	cing	
20	Add the entire <b>11 μl</b> library to the sequencing mix tube	
21	Pipette entire $\  \  \  \  \  \  \  \  \  \  \  \  \ $	ort

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Unpause sequencing run in MinKNOW software