

\* Based on Kristina Terpis training, who uses a modified Emma strand protocol (github lab notebook May 31, 2019)  
 First time using kit: Add 96  $\mu$ L to Wash Buffer, reconstitute Pro K & DNase as directed & store aliquots in freezer ( $-20^{\circ}\text{C}$ )

Preparation: This protocol is for clippings in Zymo DNA/RNA shield (vol  $\sim 1\text{ mL}$ ). These were stored @  $-80^{\circ}\text{C}$ . Thaw samples to RT on benchtop.

Clean bench w/ 10% Bleach  $\rightarrow$  DI water  $\rightarrow$  70% EtOH  $\rightarrow$  RNase Zap  
 Clean pipettes, tip boxes, & centrifuge/heating block controls w/ 70% EtOH  $\rightarrow$  RNase Zap (w/ kimwipe, don't spray)

Notes: This protocol will start with taking 300  $\mu$ L of the DNA/RNA shield from the sample and performing a ProK digestion.

\* If the liquid looks very dark (ex: P. aeruginosa), you can take 150  $\mu$ L of the shield and add 150  $\mu$ L of clean DNA/RNA shield. proceed as is.

\* If you want to add tissue from the clipping into the extraction, example protocol from Kristina: (but see github for other bead beating ext)

- $\rightarrow$  100  $\mu$ L of clean DNA/RNA shield
- $\rightarrow$  300  $\mu$ L of sample DNA/RNA shield
- $\rightarrow$  some tissue + beads
- $\rightarrow$  bead beat for 1-2 minutes
- $\rightarrow$  transfer 300  $\mu$ L to a new tube and proceed.

① On a clean work surface, stand up the tubes in numerical order & photograph (for github post).

② Into clean 1.5 mL tubes, transfer 300  $\mu$ L of the DNA/RNA shield from the sample tube after inverting the tube to mix.

Add: 30  $\mu$ L ProK digestion buffer  
 15  $\mu$ L of ProK (freezer)

Invert 3x + incubate 15 minutes

③ Put samples back in  $-80^{\circ}\text{C}$  freezer

④ Turn on heating block to  $70^{\circ}\text{C}$  and put a 1.5 mL tube w/ Tris-EDTA for DNA elution into the heating block to be warmed.

To Page No. \_\_\_\_\_

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From Page No. \_\_\_\_\_

- ⑤ Invert / flick tubes & centrifuge @ 9000 rcf for 3 minutes to pellet
- ⑥ Transfer all supernatant to a new 1.5 mL tube and add 345 (1:1)  $\mu$ L of DNA/RNA lysis buffer  
 ↳ pipette up and down and add ~~add~~ whole volume (700  $\mu$ L max) to labelled yellow DNA spin - away column in collection tube.
- ⑦ Spin 30s @ 15,000 rcf and SAVE FLOW THROUGH (transfer to a new 1.5 mL tube for RNA extraction. Discard FT.)
- ⑧ Place column in new collection tube. Add 400  $\mu$ L of prep buffer & spin.
- ⑨ Add 700  $\mu$ L of ~~DNA~~ wash buffer & spin. (30s, 15000 rcf). Discard flow through.  
 ↳ repeat this step.
- ⑩ Spin columns (dry) for 2 min @ 15,000 rcf to dry column.
- ⑪ Discard collection tube & transfer column to final DNA tube [sample #, gDNA, initials, project info, date]  
 ↳ Add 50  $\mu$ L of warmed Tris directly to column. & incubate for 5 min @ RT.  
 ↳ Spin @ 15000 rcf for 30 seconds.  
 ↳ Repeat elution step into same tube.
- ⑫ Aliquot 10  $\mu$ L of DNA into labelled strip tubes & place DNA tubes in  $-20^{\circ}\text{C}$ .  
 ↳ keep strip tubes on cold block until QC (same day)
- ⑬ Add 700  $\mu$ L (1:1 ratio) of 100% Ethanol to the RNA tube (flow through step 7)
- ⑭ Pipette up & down to mix & add 700  $\mu$ L of mixture to labelled green RNA column in collection tube  
 ↳ spin for 30s @ 15000 rcf & discard flow through  
 ↳ repeat step w/ rest of mixture.
- ⑮ Add 400  $\mu$ L of wash buffer & spin. discard flow through
- ⑯ DNase digestion:  
 Make sol<sup>n</sup>: - per sample, mix 5  $\mu$ L DNase I ( $-20^{\circ}\text{C}$ ) with 75  $\mu$ L of DNA digestion buffer in a 1.5 mL (or larger) tube for all samples. Mix gently (Vortex)  
 ↳ add 80  $\mu$ L of this mixture directly into each column.  
 ↳ incubate @ RT for 15 min
- ⑰ Add 400  $\mu$ L of prep buffer. Spin & discard FT.
- ⑱ Add 700  $\mu$ L of wash buffer. Spin & discard FT.  
 ↳ Repeat this step.
- ⑲ Dry membrane by spinning @ 15000 rcf for 2 min. (not warmed Tris)
- ⑳ Same as for DNA, elute twice w/ 50  $\mu$ L of DNase/RNase free  $\text{H}_2\text{O}$   
 ↳ before spinning, incubate ~ 2 minutes instead of the whole 5 for DNAs
- ㉑ Aliquot 10  $\mu$ L for QC & put final tubes into  $-80^{\circ}\text{C}$ .

To Page No. \_\_\_\_\_

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TITLE ENCORE DNA Extractions

Project No. ①

Book No. ①

From Page No. \_\_\_\_\_

Qubit Results 10/3/22

	DNA 1	DNA 2	RNA 1	RNA 2
DLAB D1	4.20	4.08	12.4	12.2
DLAB F6	2.20	2.14	nd	nd
MCAV B1	3.70	3.70	14.8	14.8
MCAV C4	6.86	6.78	12.2	12.2
MDEC A7	15.0	14.9	18.4	18.2
MDEC C6	6.24	6.08	16.8	16.6
PAST D7	2.34	2.24	nd	nd
PAST F4	3.26	3.14	14.4	14.4

DNA

S1 201.76

S2 23899.02

RNA

S1 415.70

S2 9890.95

DLAB D1 ① DD  
DLAB F6 ② DF  
MCAV B1 ③ CB  
MCAV C4 ④ CC  
MDEC A7 ⑤ MA  
MDEC C6 ⑥ MC  
PAST D7 ⑦ PD  
PAST F4 ⑧ PF

10/3/22

Day 2 10/5/22

DLAB A4	DA	1
DLAB C7	DC	2
MCAV E6	CE	3
MCAV F7	CF	4
MDEC D4	MD	5
MDEC E1	ME	6
PAST A1	PA	7
PAST B6	PB	8

Extraction Notes: PAST-F4 looks bad @ lysis buffer step. Mucus-y & not wanting to mix w/ the lysis buffer. Filter pigmented throughout RNA extraction & RNA @ end pale brown.

Notes on Qubit & Gels (Kristine training):

- Qubit: for each DNA & RNA, make enough reactions for  $\# \text{ samples} + 2 \text{ stds} + 1 = (n)$ 
  - standards are in basket in fridge, reagent & buffer in drawer HD11 (RT)
  - standard tubes get 190  $\mu\text{L}$  of master mix & sample tubes get 199  $\mu\text{L}$
  - Master mix:  $(199 \times n) \mu\text{L buffer} + (1 \times n) \mu\text{L reagent}$  (make 1 DNA mM & 1 RNA mM)
  - add 10  $\mu\text{L}$  of standards & 1  $\mu\text{L}$  of samples. Vortex, spin, & sit 2 min.
- Qubit: read standards once then read all samples 2x.

- Gel: 1.5% thin gel (50  $\mu\text{L}$  TAE (NEW) + 0.75 g agarose), Microwave 45s & cool
  - add 1  $\mu\text{L}$  of gel green. Swirl & pour. pop bubbles & add 2 combos.
  - sit for 30 mins.

Loading: add 1  $\mu\text{L}$  of loading dye to remaining DNA/RNA (in  $9 \mu\text{L}$ ) in strip tube.  
 spin & load whole volume into gel. Load 3  $\mu\text{L}$  of Gene Ruler 1kb ladder (500)

Run: 400 amp, 60 V, 60 minutes. Image using phone & orange filter.

To Page No. \_\_\_\_\_

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