

Jun 12, 2023

Version 6

# © DNA extraction (BOMB) V.6

DOI

dx.doi.org/10.17504/protocols.io.n2bvj6mdnlk5/v6

Yin-Tse Huang<sup>1</sup>, Tsu-Chun Hung<sup>1</sup>

<sup>1</sup>KMU



### Yin-Tse Huang

Kaohsiung Medical University

## Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN ACCESS



DOI: https://dx.doi.org/10.17504/protocols.io.n2bvj6mdnlk5/v6

**Protocol Citation:** Yin-Tse Huang, Tsu-Chun Hung 2023. DNA extraction (BOMB). **protocols.io** <a href="https://dx.doi.org/10.17504/protocols.io.n2bvj6mdnlk5/v6">https://dx.doi.org/10.17504/protocols.io.n2bvj6mdnlk5/v6</a> Version created by <a href="https://dx.doi.org/10.17504/protocols.io.n2bvj6mdnlk5/v6">Yin-Tse Huang</a>

**License:** This is an open access protocol distributed under the terms of the **Creative Commons Attribution License**, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

Protocol status: Working

We use this protocol and it's working

Created: June 11, 2023



Last Modified: September 13, 2023

**Protocol Integer ID: 83214** 

**Keywords:** dna extraction, bomb, dna, extraction

### **Abstract**

DNA extraction (BOMB)

## **Materials**

1. Lysis master mix (870 uL/sample)

	А	В	
	TE buffer	225 uL	
	Lysis buffer	375 uL	
	Ammonium acetate	270 uL	

### 2. TE buffer

А	В
Tris HCI pH8.0	10mM
EDTA	1mM

## 3. Lysis buffer

А	В
GITC	4M
Tris HCI pH8.0	50mM
SDS	0.5g
EDTA	20mM



# Troubleshooting



# **Sample Collection**

3m

Add  $\perp$  200  $\mu$ L of **0.5 mm beads** to 2mL screw tube

30s



2 Add  $\perp$  200  $\mu$ L of **1 mm beads** to 2mL screw tube





3 Add  $\perp$  870  $\mu$ L Lysis master mix to 2mL screw tube. The final look:





#### Note

In 11F, 4°C fridge

Lysis master mix: 225 µL of TE buffer + 375 µL of lysis buffer + 270 µL of 10M ammonium acetate

4 Collect 4 20-50 mg of **sample** to 2mL screw tube

### 1m

#### Note

You can collect up to 100 mg of sample if you can until you bump into the low DNA quality or PCR success rate; by then it means too many inhibitors in the sample and you have to lower the input.

# Sample crush

4m

5 Put the 2mL screw tube in mixmill for sample crush, at 3200 rpm 00:04:00



4m



### Note

Remember to balance if you have odd number of samples

# Centrifugation

3m

6 Put 2mL screw tube in centrifuge for centrifugation, at this condition:

3m

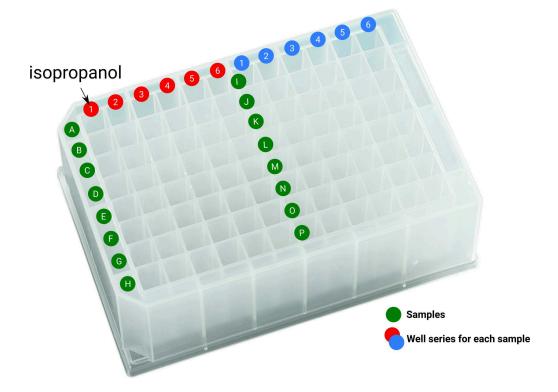
**3** 10 x g, 25°C, 00:03:00

# **DNA** purification

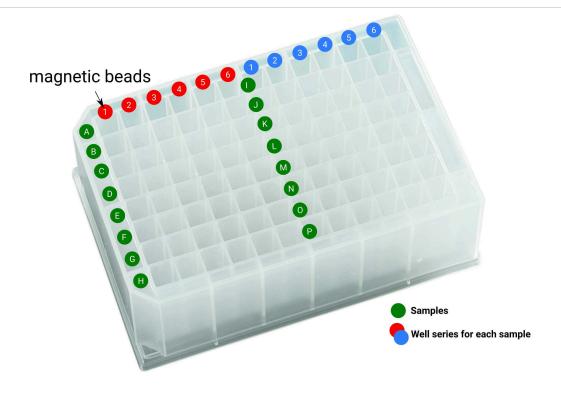
37m 30s

Add  $\Delta 350 \mu L$  of **isopropanol** to the 1st well of 96 well plate

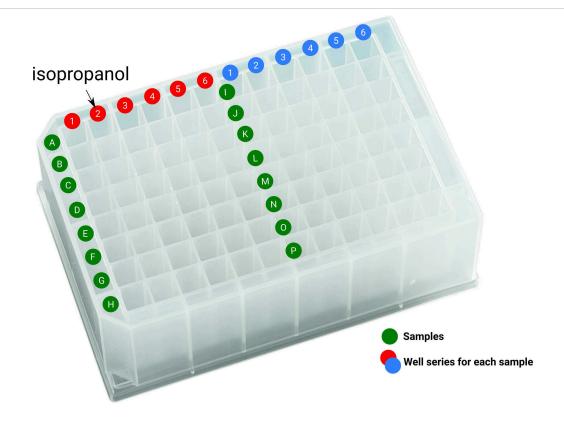
30s



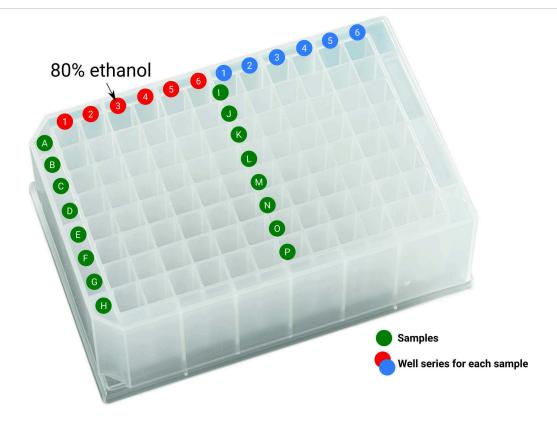
Add  $\perp$  100  $\mu$ L of magnetic beads (10mg/ml) to the 1st well of 96 deep well plate



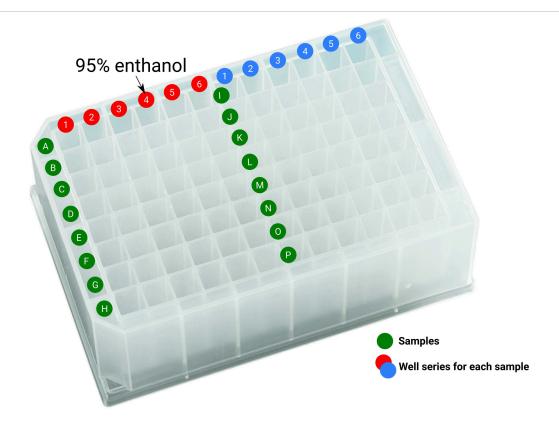
9 Add  $\perp$  400  $\mu$ L of **isopropanol** to the 2nd well of 96 deep well plate



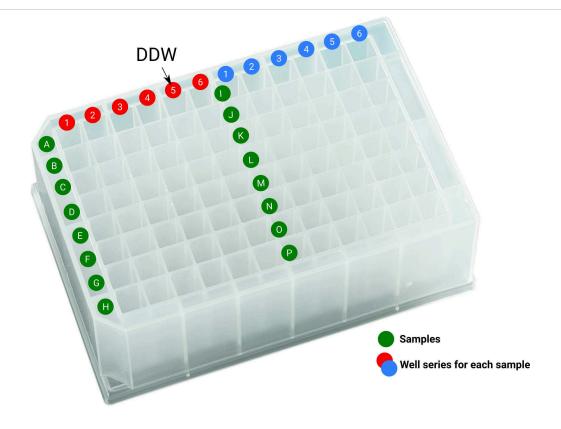
10 



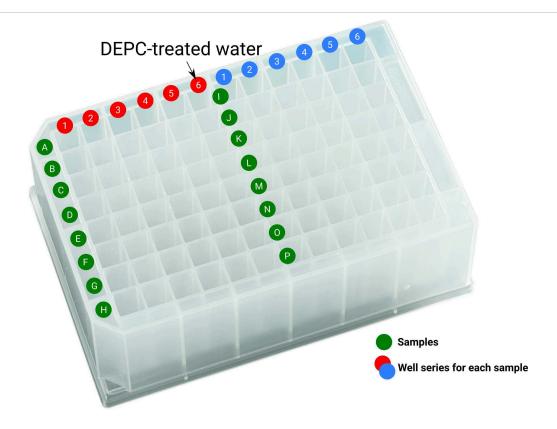
11 



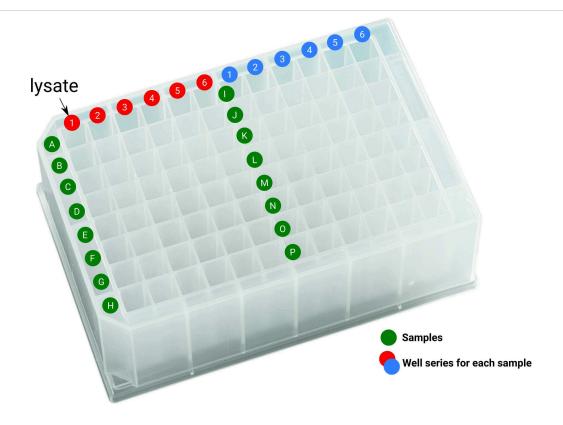
12 



13 Add  $\perp$  100  $\mu$ L of **DEPC-treated water** to the 6th well of 96 deep well plate



14 well of 96 deep well plate



### Note

Pipetting as many lysate as you can, as long as it's free of any cell debris (no solids in your tip)

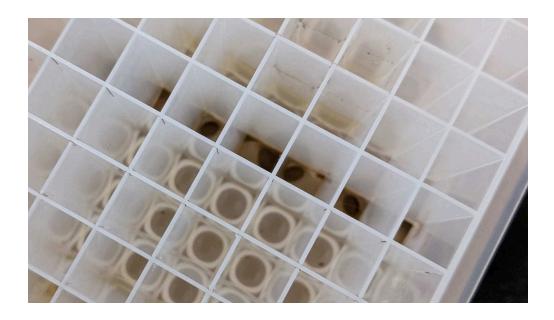
15 Put the prepared 96 deep well plate in the automated DNA extraction machine and select the BOMB protocol

34m

16 After the extraction is done, put on the 96 magnetic plate to pellet the magnetic bead residues.







17 Collect  $\perp$  100  $\mu$ L of the **eluted sample** (avoid getting magnetic bead) as the DNA template for downstream experiments



