

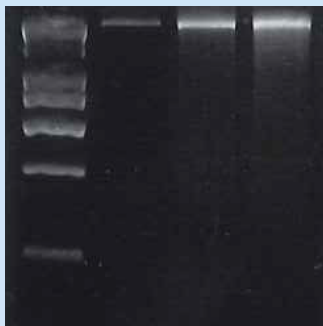
# DNA Digestion with DNase I

Steven J. Burgess

## Abstract

Protocol for DNase I digestion of nuclei for 'double hit' DNase-SEQ analysis adapted from doi:10.1038/nmeth.2762

As a general guide DNA should show moderate to light smearing after completing the protocol as seen the gel image in Figure 1.



**Figure 1:** A representative 2% (w/v) LMP agarose gel of DNase I treated nuclei ( $\sim 2 \times 10^8$ ) from *Sorghum bicolor*. To each sample the following amounts of DNase I were added: none, 7.5U, 12.5U.

Over-digested samples will result in a high background signal from unspecific cutting of DNA and should be avoided. An example of an over-digested sample is provided in lane 5 of Figure 2.



**Figure 2:** A representative 2% (w/v) LMP agarose gel of DNase I treated nuclei ( $\sim 2 \times 10^8$ ) from *Sorghum bicolor*. To each sample the following amounts of DNase I were added: none, 5,U 7.5U,10U, 12.5U.

Gel fragments from ~50-600bp should be excised and DNA purified. From experience taking only smaller fragments (50-200 bp) caused difficulties in library preparation.

**Citation:** Steven J. Burgess DNA Digestion with DNase I. [protocols.io](https://doi.org/10.17504/protocols.io.hdfb23n)

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## Materials

cOmplete™, Mini, EDTA-free (Protease Inhibitor) [#11836170001](#) by [Roche](#)

Qiaquick gel extraction kit 28704 by [Qiagen](#)

DNase I, RNase free [EN0525](#) by [Thermo Fisher Scientific](#)

RNase A [R4642-10MG](#) by [Sigma-aldrich](#)

GlycoBlue Coprecipitant [AM9515](#) by [Thermo Fisher Scientific](#)

UltraPure™ Phenol:Chloroform:Isoamyl Alcohol (25:24:1, v/v) [15593031](#) by [Thermo Fisher Scientific](#)

UltraPure™ Low Melting Point Agarose [16520050](#) by [Thermo Fisher Scientific](#)

## Protocol

### Step 1.

Before the experiment make up stocks and store at 4°C.

Buffer A (1L)

Component	[Stock]/M	Volume/mL	[Final]/mM
TRIS-HCl pH 8.0	1	15	15
NaCl	5	3	15
KCl	1	60	60
EDTA pH 8.0	0.5	2	1
dH <sub>2</sub> O		up to 1L	

10x Buffer D (10 mL)

Component	[Stock]/M	Volume	[Final]/mM
MgCl <sub>2</sub>	1	0.6	60

NaCl	5	0.2	100
CaCl <sub>2</sub>	1	0.1	10
TRIS-HCl pH 8.0	1	4	400
dH <sub>2</sub> O	up to 10 mL		

## Stop Buffer (1L)

Component	[Stock]	Volume/mL	[Final]
TRIS-HCl pH8.0	1M	50	50mM
NaCl	5M	20	100mM
SDS	20% (w/v)	5	0.1% (w/v)
EDTA pH8.0	0.5M	200	100mM
dH <sub>2</sub> O	up to 1L		

## Buffer Preparation

### Step 2.

On the day of the experiment make the following buffers:

### Buffer A+

Component	[Stock]	Volume/μL
Spermine	0.1M	30
Spermidine	0.1M	100
DTT	1M	10
PIC*	100x	200
NP-40		40
Buffer A		20

\*PIC=Protease inhibitor cocktail (1 tablet of Roche EDTA-free complete protease inhibitor is resuspended in 500 μl)

### 1x Buffer D

Component	Volume/mL
10x Buffer D	1.5
Buffer A+	13.5

## Stop Buffer Stock

Component	[Stock]	Volume
Spermine	0.1M	30
Spermidine	0.1M	100
RNAse A	10mg/mL	20
Stop Buffer Stock		Up to 10mL

## Prepare nuclei

### Step 3.

Spin down nuclei at 1500g for 10 mins. Wash once with buffer A and re-suspend in 1x Buffer D to a concentration of  $4 \times 10^8$  nuclei/mL

#### 📌 NOTES

**Steven Burgess** 16 Mar 2017

Note: The amount of nuclei required for each reaction may well vary between species, a concentration of  $\sim 4 \times 10^8$  worked well for *Sorghum bicolor*, *Setaria italica*, *Zea mays* and *Brachypodium distachyon*.

## Preparing Reaction

### Step 4.

Prepare enzyme mixtures in 2 mL centrifuge tubes:

Final Enzyme Concentration/U	Amount of Buffer D/ $\mu$ L;	Amount of DNaseI [1U/ $\mu$ L]/ $\mu$ L
0	0	500
2.5	2.5	497.5
7.5	7.5	492.5
12.5	12.5	487.5

#### 📌 NOTES

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Note: From experience a range of enzyme concentrations should be used due to variability in digestion outcome. Manual inspection of the resulting digestions dictates which samples to choose

for further processing.

#### Pre-Incubation

##### Step 5.

Incubate nuclei and enzyme mixtures at 37°C

 DURATION

00:01:00

#### Digestion

##### Step 6.

Add 500 µL of nuclei to 500 µL of enzyme mix. Mix by inverting and incubate at 37°C.

#### Stopping the reaction

##### Step 7.

Stop the reaction by addition of 1 mL of stop buffer and incubate tubes at 55 °C.

 DURATION

00:15:00

#### Proteinase K digestion

##### Step 8.

Add proteinase K [20mg/mL] to each tube and incubate at 55 °C.

 AMOUNT

2 µl Additional info:

 DURATION

01:00:00

#### DNA extraction part I

##### Step 9.

Add 1:1 (v/v) phenol:chloroform:isoamylalcohol (25:24:1) to each reaction, hand shake vigorously. spin down 14,000rpm in microcentrifuge tube.

 DURATION

00:15:00

##### Step 10.

 DURATION

01:00:00

##### Step 11.

Spin down 14,000 rpm in a microcentrifuge

 DURATION

00:15:00

##### Step 12.

Dispose of ethanol solution and wash pellet in 1 mL 70% ethanol (v/v). Invert tube 3-5 times and spin down 14,000 rpm in microcentrifuge.

**Step 13.**

**Step 14.**

DNA extraction part V

**Step 15.**

Perform gel extraction using QIAGEN QIAquick gel extraction kit, quantify the amount of DNA using a Qubit prior to library construction.