



Jun 03, 2021

© DNA Library Prep for BGISEQ-500/MGISEQ-2000/DNBSEQ-G400/MGISEQ-200/DNBSEQ-G50

Hongfang Zhang¹

¹GigaScience Press



ABSTRACT

This DNA library prep protocol is specifically designed for WGS library construction for MGI highthroughput sequencing platforms. It's applicable for samples from all common animals, plants, fungi bacteria, etc., including human, mice, rice, Arabidopsis, yeast, *E. coli*, and Metagenomics samples.

EXTERNAL LINK

https://en.mgi-tech.com/Download/download_file/id/200

DO

dx.doi.org/10.17504/protocols.io.bve2n3ge

EXTERNAL LINK

https://en.mgi-tech.com/Download/download_file/id/200

PROTOCOL CITATION

Hongfang Zhang 2021. DNA Library Prep for BGISEQ-500/MGISEQ-2000/DNBSEQ-G400/MGISEQ-200/DNBSEQ-G50. **protocols.io**

https://dx.doi.org/10.17504/protocols.io.bve2n3ge

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

CREATED

May 31, 2021

LAST MODIFIED

Jun 03, 2021

PROTOCOL INTEGER ID

50362

Fragmentation

1.1 Transfer genomic DNA to a new 0.2 mL PCR tube. The volume should be less than or equal to □45 μl

. If the volume is less than □45 μl , add dilution buffer to bring the final volume to □45 μl

 $\textbf{Citation:} \ \ \text{Hongfang Zhang (06/03/2021).} \ \ DNA \ \ \text{Library Prep for BGISEQ-500/MGISEQ-2000/DNBSEQ-G400/MGISEQ-2000/DNBSEQ-G50.} \\ \underline{\text{https://dx.doi.org/10.17504/protocols.io.bve2n3ge}}$

Components	Volume
DNA	XμL
dilution buffer	45-X μL
Total	45 μL

Table 1.1 Input DNA Dilution

- 1.2 Mix Frag Enzyme II by inverting 10 times then centrifuge briefly and place on ice for use. DO NOT vortex Frag Enzyme II. Vortex Frag Buffer II 3 times (3s each) then centrifuge briefly and place on ice for use.
- 1.3 Prepare the fragmentation mixture on ice. Pipette up and down at least 10 times to mix thoroughly.
 (DO NOT vortex)

Components	Volume
Frag Buffer II	10 μL
Frag Enzyme II	5 μL
Total	15 μL

Table 1.2 Fragmentation Mixture

- 1.4 Transfer \Box 15 μ I of the fragmentation mixture to the 0.2 mL PCR tube from step 1.1. Pipette at least 10 times to mix thoroughly and briefly centrifuge to collect the solution at the bottom of the tube.
- 1.5 Set and run the following program on the thermocycler (see Table 1.3). Make sure the thermocycler has cooled to § 4 °C . Place the 0.2 mL PCR tube from step 1.4 into the thermocycler and skip the § 4 °C Hold step to start the reaction at § 30 °C .

Temperature	Time
Heated lid	On
4°C	Hold
30°C	8 min
65°C	15 min
4°C	Hold

Table 1.3 Fragmentation Reaction Conditions

Briefly centrifuge to collect the solution at the bottom of the tube.

2 Size Selection

30m 2.1 Take DNA Clean Beads out of the refrigerator and allow **© 00:30:00** for the solution to warm to room temperature. Vortex and mix thoroughly before use. 2.2 Transfer \blacksquare 36 μ I of DNA Clean Beads to the 1.5 mL centrifuge tube containing \blacksquare 60 μ I of fragmentation product from step 1.6. Pipette up and down at least 10 times or vortex to mix thoroughly. 5m 2.3 Incubate at room temperature for 00:05:00. 2.4 Centrifuge briefly and place the tube onto the Magnetic Separation Rack for © 00:05:00 until the liquid becomes clear. Then, carefully transfer the supernatant to a new 1.5 mL centrifuge tube. 2.5 Transfer 12 μl of DNA Clean Beads to the centrifuge tube with 96 μl of supernatant. Pipette at least 10 times to mix thoroughly. 5m $2.6 \quad \text{Incubate at room temperature for } \textcircled{00:05:00} \; .$ 2.7 Centrifuge briefly and place the tube onto a Magnetic Separation Rack for © 00:05:00 until the liquid becomes clear. Carefully remove and discard the supernatant with pipette. 2.8 Keep the centrifuge tube on the Magnetic Separation Rack and add ⊒200 µl of freshly prepared 80% ethanol to wash the beads and the walls of the tube. Carefully remove and discard the supernatant. Repeat step 2.8 once. Remove all of the liquid from the tube without disrupting the beads. You may 2.9 centrifuge briefly to collect any remaining liquid at the bottom of the centrifuge tube, separate the beads magnetically, and then remove any remaining liquid using a small volume pipette. Keep the centrifuge tube on the Magnetic Separation Rack with the lid open, and air-dry beads at room 2.10 temperature until no wetness (reflectiveness) is observed but before the pellet begins to crack. 2.11 Remove the centrifuge tube from the Magnetic Separation Rack and add 44 µl of TE Buffer to elute the DNA. Pipette up and down at least 10 times to mix thoroughly.

- $2.12 \quad \text{Incubate at room temperature for } \textcircled{00:05:00} \; .$
- 2.13 Centrifuge briefly and place the tube onto a Magnetic Separation Rack for **© 00:05:00** until the liquid becomes clear. Carefully transfer **41 µl** of supernatant to a new 0.2 mL PCR tube.
- 2.14 Quantify the Size selection products with dsDNA Fluorescence Assay Kits such as Qubit® dsDNA HS Assay Kit or Quant-iT™ PicoGreen® dsDNA Assay Kit.

3 End Repair and A-tailing

- 3.1 Transfer ≤100 ng of fragmentation product to a new 0.2 mL PCR tube. Add TE Buffer for a total volume of \Box 40 μ I .
- 3.2 Prepare the end repair and A-tailing mixture on ice (see Table 3.1).

Components	Volume
ERAT Buffer	7.1 μL
ERAT Enzyme Mix	2.9 μL
Total	10 μL

Table 3.1 End Repair and A-tailing Mixture

- 3.3 Transfer 10 μl of the end repair and A-tailing mixture to the 0.2 mL PCR tube from step 3.1. Vortex 3 times (3s each) and briefly centrifuge to collect the solution at the bottom of the tube.
- 3.4 Place the 0.2 mL PCR tube from step 3.3 into the thermocycler and run the program in Table 3.2.

Temperature	Time
Heated lid	On
37°C	30 min
65°C	15 min
4°C	Hold

Table 3.2 End Repair and A-tailing Reaction Conditions

3.5 Briefly centrifuge to collect the solution at the bottom of the tube.

4 Adapter Ligation

4.1 Add 5 μ L of MGIEasy DNA Adapters (diluent) to the PCR tube from step 3.5. Vortex 3 times (3s each) and briefly centrifuge to collect the solution at the bottom of the tube.

4.2	Prepare the	Adapter	ligation	mixture	on ice.
-----	-------------	---------	----------	---------	---------

Components	Volume
Ligation Buffer	23.4 μL
DNA Ligase	1.6 μL
Total	25 μL

Table 4.1 Adapter Ligation Mixture

4.3 Pipette slowly and transfer **25 μl** of Adapter ligation mixture to the 0.2 mL PCR tube from step 4.1. Vortex 6 times (3s each) and centrifuge briefly to collect the solution at the bottom of the tube.

4.4 Place the PCR tube from step 4.3 into the thermocycler and run the program.

Temperature	Time
Heated lid	On
23°C	30 min
4°C	Hold

Table 4.2 Adapter Ligation Reaction Conditions

- 4.5 Centrifuge briefly to collect the solution at the bottom of the tube.
- 4.6 Add ■20 μl of TE Buffer for a total volume of ■100 μl and transfer all of the solution to a new 1.5 mL centrifuge tube.

5 Adapter-Ligated DNA Cleanup

- Repeat step 5.5 once. Remove all of the liquid from the tube without disrupting the beads. You may centrifuge briefly to collect any remaining liquid at the bottom of the centrifuge tube, separate the beads magnetically and then remove any remaining liquid using a small volume pipette.
- 5.7 Keep the centrifuge tube on the Magnetic Separation Rack with the lid open and air-dry beads at room temperature until no wetness (reflectiveness) is observed, but before the pellet begins to crack.
- 5.8 Remove the centrifuge tube from the Magnetic Separation Rack and add **21 μl** of TE Buffer to elute the DNA. Pipette up and down at least 10 times to mix thoroughly.

5m

5.9 Incubate at room temperature for \bigcirc 00:05:00.

6 PCR Amplification

6.1 Prepare the PCR amplification mixture on ice.

6.2 Transfer 31 μl of PCR amplification mixture to the PCR tube from step 5.10. Vortex 3 times (3s each) and centrifuge briefly to collect the solution at the bottom of the tube.

Components	Volume
PCR Enzyme Mix	25 μL
PCR Primer Mix	6 μL
Total	31 µL

Table 6.1 PCR Amplification Mixture

6.3 Place the PCR tube from step 6.2 into the thermocycler and run the program in Table 6.2. Please refer to Table 6.2 to adjust the number of cycles with different DNA input.

Temperature	Time	Cycles
Heated lid	on	
95°C	3 min	1 cycle
98°C	20 s	
60°C	15 s	3-12 cycles
72°C	30 s	
72°C	10 min	1 cycle
4°C	Hold	

Table 6.2 PCR Amplification Reaction Conditions

- 6.4 Centrifuge briefly to collect the solution at the bottom of the tube.
- 6.5 Transfer all the solution to a new 1.5 mL centrifuge tube.

7 Cleanup of PCR Product

7.1 Take DNA Clean Beads out of the refrigerator and allow **© 00:30:00** for the solution to warm to room temperature. Vortex and mix thoroughly before use.

7.2 Transfer \blacksquare 50 μ I of DNA Clean Beads to the centrifuge tube from step 6.5. Pipette up and down at

protocols.io
7
06/03/2021

7.3 Incubate at room temperature for **© 00:05:00**.

5m

- 7.4 Centrifuge briefly and place the tube onto a Magnetic Separation Rack for © 00:05:00 until the liquid becomes clear. Carefully remove and discard the supernatant with a pipette.
- 7.5 Keep the tube on the Magnetic Separation Rack and add 200 μl of freshly prepared 80% ethanol to wash the beads and the walls of the tube. Incubate for 00:00:30 and carefully remove and discard the supernatant.
- 7.6 Repeat step 7.5 once. Remove all of the liquid from the tube without disrupting the beads. You may centrifuge briefly to collect any remaining liquid at the bottom, separate the beads magnetically, and remove any remaining liquid using a small volume pipette.
- 7.7 Keep the centrifuge tube on the Magnetic Separation Rack with the lid open, and air-dry beads at room temperature until no wetness (reflectiveness) is observed but before the pellet begins to crack.
- 7.8 Remove the centrifuge tube from the Magnetic Separation Rack and add 32 μl of TE Buffer to elute the DNA. Pipette up and down at least 10 times to mix thoroughly.
- 7.9 Incubate at room temperature for © 00:05:00.

5m

7.10 Centrifuge briefly, then place the centrifuge tube back onto the Magnetic Separation Rack for © 00:05:00 until the liquid becomes clear. Transfer 30 μl of supernatant to a new 1.5 mL centrifuge tube. 5m

8 Quality Control of PCR Product

8.1 Quantify the purified PCR products with dsDNA Fluorescence Assay Kits such as Qubit® dsDNA HS Assay Kit or Quant-iT™ PicoGreen® dsDNA Assay Kit. The required yield for PCR products is ≥ 1 pmol. Please refer to Formula 1 to calculate the amount of DNA needed. For example, for 384 bp PCR product the yield should reach 250 ng. For pooled sequencing, please follow instructions provided by MGIEasy DNA Adapters User Manual. Detailed information shows how to plan your sample pooling. Quantify your Adapter-ligated samples before pooling. The total yield after pooling should be 1 pmol, with a total volume ≤ 48 µl.

Formula 1

Conversion between 1 pmol of dsDNA sample and Mass in ng Mass (ng) corresponding to 1 pmol PCR Products= DNA Fragment Size (bp) /1000 bp $\times 660$ ng

8.2 Assess the fragment size distribution of purified PCR products with electrophoresis-based equipment such as Bioanalyzer, Tapestation (Agilent Technologies), LabChip® GX, GXII, GX Touch (PerkinElmer), or Fragment Analyzer (Advanced Analytical).

9 Denaturation

- 9.1 According to the PCR product size, transfer 1 pmol of PCR product to a new 0.2 mL PCR tube. Add TE Buffer for a total volume of **48 μl**.
- 9.2 Place the 0.2 mL PCR tube from step 9.1 into the thermocycler and run the program in Table 9.1.

Temperature	Time
Heated lid	On
95°C	3 min

Table 9.1 Denaturation Reaction Conditions

9.3 After the reaction is completed, immediately place the 0.2 mL PCR tube on ice for **© 00:02:00**, then centrifuge briefly.

10 Single Strand Circularization

10.1 Prepare the single strand circularization mixture on ice (see Table 10.1).

Components	Volume	
Splint Buffer	11.6 μL	
DNA Rapid Ligase	0.5 μL	
Total	12.1 μL	

Table 10.1 Single Strand Circularization Mixture

- Transfer 12.1 μ L single strand circularization mixture to the 0.2 mL PCR tube from step 9.3. Vortex 3 times (3s each) and centrifuge briefly to collect the solution at the bottom of the tube.
- $10.3 \quad \text{Place the PCR tube into the thermocycler and run the program in Table 10.2}.$

Temperature	Time
Heated lid	On
37°C	30 min
4°C	Hold

Table 10.2 Single Strand DNA Circularization Reaction Conditions

10.4 After the reaction is complete, immediately place the tube on ice for the next reaction.

11 Enzymatic Digestion

11.1 Prepare the following enzymatic digestion mixture (see Table 11.1) on ice during the reaction in step 10.3.

Components	Volume
Digestion Buffer	1.4 μL
Digestion Enzyme	2.6 μL
Total	4 μL

Table 11.1 Enzymatic Digestion Mixture

11.2 Transfer **4 μl** of enzymatic digestion mixture into the PCR tube from step 10.4. Vortex 3 times (3s each) and centrifuge briefly to collect the solution at the bottom of the tube.

11.3 Place the PCR tube from step 11.2 into the thermocycler and run the program in Table 11.2.

Temperature	Time
Heated lid	On
37°C	30 min
4°C	Hold

Table 11.2 Enzymatic Digestion Reaction Conditions

 $11.4 \quad \text{Centrifuge briefly to collect the solution at the bottom of the tube.}$

11.5 Add 27.5 μl Digestion Stop Buffer to the 0.2 mL PCR tube. Vortex 3 times (3s each) and centrifuge briefly to collect the solution at the bottom of the tube. Transfer all of the solution into a new 1.5 mL centrifuge tube.

12 Enzymatic Digestion Product Cleanup

30m 12.1 Take DNA Clean Beads out of the refrigerator and allow © 00:30:00 for the solution to warm to room temperature. Vortex and mix thoroughly before use. 12.2 Transfer ■170 µl of DNA Clean Beads to the Enzymatic Digestion product from step 11.5. Gently pipette at least 10 times or vortex to mix thoroughly. 10m 12.3 Incubate at room temperature for **© 00:10:00**. 12.4 Centrifuge briefly and place the 1.5 mL tube on the Magnetic Separation Rack for © 00:05:00 until the liquid becomes clear. Carefully remove and discard the supernatant using a pipette. 30s 12.5 With the 1.5 mL tube on the Magnetic Separation Rack, add 500 µl of freshly prepared 80% ethanol to the tube without disturbing the beads. Incubate for \(\oint 00:00:30 \). Carefully remove and discard the supernatant. 12.6 Repeat step 12.5 once. Remove all of the liquid from the tube without disrupting the beads. You may centrifuge briefly to collect any remaining liquid at the bottom, separate the beads magnetically, then remove any remaining liquid using a small volume pipette. 12.7 Keep the 1.5 mL centrifuge tube on the Magnetic Separation Rack with the lid open, and air-dry beads at room temperature until no wetness (reflectiveness) is observed but before the pellet begins to crack. 12.8 Remove the 1.5 mL centrifuge tube from the Magnetic Separation Rack and add **□32** μI of TE Buffer to elute the DNA. Gently pipette up and down at least 10 times to mix thoroughly or until the beads are fully resuspended. 10m 12.9 Incubate at room temperature for **© 00:10:00**. 5m 12.10 Centrifuge briefly and place the centrifuge tube back onto the Magnetic Separation Rack for © 00:05:00 until the liquid becomes clear. Transfer ■30 µl of supernatant to a new 1.5 mL

centrifuge tube.

13 Quality Control of Enzymatic Digestion Product

Quantitate the purified Enzymatic Digestion product with Qubit® ssDNA Assay Kit. The final Enzymatic Digestion products (ssDNA, ng) / input products of PCR (dsDNA, ng) should be \geq 7%. For example, for 384 bp PCR product the final Enzymatic Digestion products should reach 17.5 ng.