

# Hi-C library preparation for the *Lateolabrax maculatus* genome

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## Abstract

This protocol is used to clarify the process of Hi-C library preparation for *L. maculatus* genome.

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## Protocol

### Sample preparation

#### Step 1.

- 1) Blood sample was centrifuged at 2500g for 5min at 4°C, collect precipitated cell;
- 2) Add 1ml PBS to resuspended blood cell.



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PBS by [Invitrogen - Thermo Fisher](#)

### Formaldehyde fixation

#### Step 2.

- 1) 37% formaldehyde was added with the final concentration of 1 % and the reaction was stopped with glycine after standing for 10 min at room temperature.
- 2) The formaldehyde fixed powder was then re-suspended in nuclei isolation buffer (10 mM Tris-HCl pH 8.0 , 10 mM NaCl, 1× PMSF).



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37% formaldehyde by [Sigma](#)



glycine by [Sigma](#)



10 mM Tris-HCl by [Sigma](#)



10 mM NaCl by [Beyotime](#)



1× PMSF by [Sigma](#)

### Enzyme digestion

#### Step 3.

Add enzyme (Mbo I) to digest the DNA.



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 enzyme (Mbo I) by [New England Biolabs](#)

### DNA fragment end reparation

#### Step 4.


Add 10 mM dCTP, 10 mM dGTP, 10 mM dTTP, 5U/  $\mu$ l DNA Polymerase I, Large (Klenow) Fragment using a biotinylated residue (0.4 mM biotin-14-dATP).



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 10 mM dCTP, 10 mM dGTP, 10 mM dTTP by [Invitrogen - Thermo Fisher](#)

 5U/  $\mu$ l DNA Polymerase I, Large (Klenow) Fragment by [New England Biolabs](#)

 biotinylated residue (0.4 mM biotin-14-dATP) by [Invitrogen - Thermo Fisher](#)

### In situ ligation


#### Step 5.


Add 10X NEB T4 DNA ligase buffer, 10% Triton X-100, 10 mg/ml BSA, T4 DNA ligase.




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 10X NEB T4 DNA ligase buffer by [New England Biolabs](#)

 10% Triton X-100 by [Sigma](#)

 10 mg/ml BSA by [New England Biolabs](#)

 T4 DNA ligase by [New England Biolabs](#)

### Reverse-crosslinked


#### Step 6.

Add 10 mg/ml proteinase K and 1% SDS to the tube and incubate at 56°C for overnight and purified, put the Reverse-crosslinked DNA liquid into three tube equally, add 1.5 $\times$  volumes of AMPure XP mixture to each tube, vortex and spin down briefly, incubate for 10 min at room temperature, place on the MPS for 5 min at room temperature, discard supernatant, wash the beads twice with 1 ml of freshly made 70% ethanol, air-dry the beads completely and re-suspend the beads in 30  $\mu$ l of ddH<sub>2</sub>O.



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 10 mg/ml proteinase K by [New England Biolabs](#)

 1% SDS by [Ambion](#)

 MPS by [Invitrogen - Thermo Fisher](#)

 70% ethanol by [Sinopharm Chemical Reagent Co.](#)

### Biotin-containing fragments capture

#### Step 7.

Shear 20 mg of DNA and capturing the biotin-containing fragments on streptavidin-coated beads using Dynabeads MyOne Streptavidin T1.



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 Dynabeads MyOne Streptavidin T1 by [Invitrogen - Thermo Fisher](#)

## DNA fragment end repairation

### Step 8.

Add 10X NEB T4 DNA ligase buffer with 10 mM ATP, 25 mM dNTP mix, 10 U/μl NEB T4 PNK, 3 U/μl NEB T4 DNA polymerase I, 5 U/μl NEB DNA polymerase I, Large (Klenow) Fragment.



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10X NEB T4 DNA ligase buffer by [New England Biolabs](#)



10 mM ATP by [New England Biolabs](#)



25 mM dNTP mix by [Enzymatics](#)



10 U/μl NEB T4 PNK, 3 U/μl NEB T4 DNA polymerase I, 5 U/μl NEB DNA polymerase I, Large (Klenow) Fragment by [New England Biolabs](#)

## DNA fragment adenylation

### Step 9.

Add 10X NEBuffer 2, 10 mM dATP, 5 U/μl NEB Klenow exo minus.



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10X NEBuffer 2 by [New England Biolabs](#)



10 mM dATP by [Invitrogen - Thermo Fisher](#)



5 U/μl NEB Klenow exo minus by [New England Biolabs](#)

## Adaptor ligation

### Step 10.

Add 10X T4 PNK Reaction Buffer, 100mM ATP, 600 U/ul T4 DNA Ligase, 50% PEG8000, 50 uM Ad153 barcode oligo\_2B mix (BGI, Shenzhen, China), and followed by PCR (95°C 3 min.; [98°C 20 sec., 60°C 15 sec., 72°C 15 sec.] (8 cycles); 72°C 10 min.



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10X T4 PNK Reaction Buffer by [New England Biolabs](#)



100mM ATP by [FERMENTAS Inc.](#)



600 U/ul T4 DNA Ligase by [New England Biolabs](#)



50% PEG8000 by [RIGAKU](#)