

1 **Blood collection procedure**

2 The site of blood collection was disinfected with alcohol pads. The plastic cap
3 from the needle tip was lifted and the needle was injected into the vein. The
4 plunger was softly pulled back and let the blood flow into the tube until the
5 syringe was filled. The animal was not anesthetized and blood collection was
6 carried out according to the 'Regulations for Animal Experiments in Chittagong
7 Veterinary and Animal Sciences University'

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9 **DNA Isolation methodology**

10 At first 200 µl of whole blood sample of BBG (healthy and disease free) was
11 transferred into the microcentrifuge tube with 20 µl of each Proteinase K and
12 RNase A enzyme. Then the Binding solution (200 µl) was added to the lysate
13 and mixed well by pulse-vortexing for 15 sec. the Lysate was then incubated at
14 56°C for 10 min and further 200 µl of absolute ethanol was added. Thereafter,
15 pulse vortexing for 15 sec, the lysate was carefully transferred into the spin
16 column and centrifuged at 13,000 rpm for 1 min. Subsequently, the flow
17 through was discarded and 500 µl of Washing buffer 1 was added in the spin
18 column and centrifuged (13,000 rpm for 1 min). This step was also repeated for
19 Washing solution 2. Again the flow-through was discarded and the spin column
20 was dried by centrifugation (13,000 rpm, 1 min). Then, the spin column was
21 transferred into a microcentrifuge tube and 100 µl elution buffer was added
22 and let it stand for 1 min. Finally, to elute the genomic DNA centrifugation was
23 done at 13,000 rpm for 1 min. The collected genomic DNA was then stored at -
24 20°C before going to further downstream processing.

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26 **Library preparation**

27 At first, 1µg genomic DNA was randomly fragmented by Covaris. The
28 fragmented DNA material was examined by Gel-Electrophotometric and then
29 purified by AxyPrep Mag PCR clean up Kit. The fragmented DNA material was
30 combined with End Repair Mix, incubated at 20°C for 30 min. The purified end-
31 repaired DNA was combined with A-Tailing Mix and incubated at 37°C for 30
32 min. Then, Illumina adaptors were ligated to the Adenylate 3'Ends DNA and
33 incubated at 16°C for 16h. The purified adapter-ligated DNA fragments were
34 selected as based on the insert size. Several rounds of PCR amplification with
35 PCR Primer Cocktail and PCR Master Mix were performed to enrich the Adapter-
36 ligated DNA fragments. Subsequently, the PCR products were purified and the
37 libraries were qualified by the Agilent Technologies 2100 bioanalyzer and ABI
38 StepOnePlus Real-Time PCR System. Finally, the qualified libraries were
39 sequenced pair-end using Hiseq 2500 System (Illumina).

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