

DNA Methylation Bisulfite Kit

RK30203



Instruction Manual

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1. Introduction

The DNA Methylation Bisulfite Kit combines DNA denaturation and bisulfite conversion in one step and includes an additional heating step to promote rapid DNA denaturation. This product is a ready-to-use conversion reagent. Simply add this reagent to the DNA sample, and the conversion can be completed within 40 minutes. Subsequently, use the unique low-elution spin columns for desulfonation and purification of the converted DNA. The kit can accommodate conversion of samples ranging from 100 pg to 2 μ g, with a conversion rate of unmethylated cytosine to uracil ≥99%. The high-yield converted DNA is highly suitable for applications such as PCR, qPCR and next-generation sequencing.

| Components | Component | 10 RXN | 50 RXN | 200 RXN |
|-------------------------|-----------|--------|----------|-----------|
| | number | | | |
| Conversion reagent | RM30451 | 1.3 mL | 1.3 mL*5 | 1.3 mL*20 |
| M-Binding buffer | RM30445 | 6 mL | 30 mL | 125 mL |
| M-Wash buffer | RM30446 | 1.2 mL | 6 mL | 24 mL |
| M-Desulphonation buffer | RM30447 | 2 mL | 10 mL | 40 mL |
| M-Elution buffer | RM30448 | 200 µL | 1 mL | 4 mL |
| Spin-column | RM30449 | 10 pk | 50 pk | 200 pk |
| Collection tube | RM30450 | 10 pk | 50 pk | 200 pk |

2. Components

3. Storage and Transport Conditions

Transport and store at room temperature.

4. Procedure

 Add 130 µl of Lightning Conversion Reagent to 20 µl of a DNA sample;



 Purification of Conversion Products;





4.1. Buffer Preparation

4.1.1. Before the first use, ethanol should be added to the M-Wash Buffer according to the instructions on the reagent bottle label. Check the box to mark it to avoid multiple additions. Tighten the bottle to prevent ethanol evaporation.

10 RXN: Add 4.8 mL of ethanol to 1.2 mL of M-Wash Buffer;

50 RXN: Add 24 mL of ethanol to 6 mL of M-Wash Buffer;

200 RXN: Add 96 mL of ethanol to 24 mL of M-Wash Buffer.

4.2. DNA denaturation and bisulfite conversion processes

4.2.1. In a PCR tube, add 130 μ l of Conversion Reagent to a 20 μ l of DNA sample. Mix well, then centrifuge quickly to ensure no droplets in the cap or sides of the tube.

Note: If the volume of DNA is less than 20 µl, compensate with water. Samples >20 µl must be processed using multiple

conversion reactions. Replicate reactions can be cleaned using the same column.

4.2.2. Place the PCR tube in a thermal cycler and perform the following steps:

| Temperature | Time |
|-------------|--------|
| 98°C | 8 min |
| 54°C | 40 min |
| 4°C | Hold |

Note: Store at 4°C for up to 20 hours. The 4°C storage step is optional.

4.3. Purification

4.3.1. Add 600 µl of M-Binding Buffer to a Spin-Column and place the column into a Collection Tube.

Note: The capacity of the collection tube with the column inserted is 800 µl. Empty the collection tube whenever necessary to

prevent contamination of the column contents by the flow-through.

4.3.2. Load the sample (from Step 4.2.2) into the Spin-Column containing the M-Binding Buffer. Close the cap and mix by inverting the column several times.

4.3.3. Centrifuge at full speed (12,000 x g) for 30 seconds. Discard the flow-through.

4.3.4. Add 100 µl of M-Wash Buffer to the column. Centrifuge at full speed (12,000 x g) for 30 seconds.

4.3.5. Add 200 µl of M-Desulphonation Buffer to the column and let stand at room temperature (20-30°C) for 15-20 minutes. After the incubation, centrifuge at full speed (12,000 x g) for 30 seconds.

4.3.6. Add 200 µl of M-Wash Buffer to the column. Centrifuge at full speed (12,000 x g) for 30 seconds. Repeat this wash step.

4.3.7. Repeat step4.3.6.



4.3.8. Discard the flow-through. Place the column into a 1.5 ml microcentrifuge tube and add 10 μ l of M-Elution Buffer directly to the column matrix. Centrifuge for 30 seconds at full speed (12,000 x g) to elute the DNA.

Note: Alternatively, water or TE (pH \ge 6.0) can be used for elution if required for your experiments. The DNA is ready for immediate analysis or store at or below -70°C for long term storage. The elution volume can be > 10 µl depending on the requirements of your experiments, but small elution volumes will yield higher DNA concentrations.







ABclonal Biotechnology Co. Ltd.

E-mail:info@abclonal.com

Web:www.abclonal.com

Headquarters: Building 5, Precision Medicine Industry Base Project I, Gaokeyuan 3rd Road,Donghu New Technology Development Zone, Jiangxia District, Wuhan, Hubei, China