

Human Low Methylated Genomic DNA Control



Product Highlights

- ▶ Human Low Methylated Genomic DNA is derived from male liver or whole blood and chemically and enzymatically methylated resulting in less than 5% in all regions of the genome.
- ▶ Ideal for use as control in bisulfite methylation analysis procedures including Pyrosequencing, targeted NGS, and Methylation Sensitive High-Resolution Melting (MS-HRM)
- ▶ Tested on gene specific and global methylation (*LINE-1*) assays for consistent performance
- ▶ Requires bisulfite modification prior to use

Product Contents

1 vial Human Low Methylated Genomic DNA (5 µg at 100 ng/µL, < 20% Methylation)

Ordering Information

CATALOG NUMBER	PRODUCT	PRICE
80-8062-HGUM5	Human low methylated genomic DNA (5 µg at 100 ng/ µL)	\$259.97

Example Protocol

- ▶ Bisulfite modification of controls and sample(s) of interest.
 - Zymo Research EZ Methylation kit (Cat.#D5002 or D5004) with 200 - 500 ng of input DNA following manufacturers recommended protocol.
- ▶ PCR amplification Protocol:

Component	Per 30µl reaction
10X PCR buffer (Contains 15mM MgCl ₂)	3 µL (1x)
25 mM MgCl ₂	1.8 µL (3.0 mM final conc.)
10 mM dNTPs	0.6 µL (200 µM of each)
10 µM Forward primer	0.6 µL (6 pmol)
10 µM Reverse primer	0.6 µL (6 pmol)
HotStart Taq Polymerase (5 U/µl)	0.15 µL (0.75 U)
DNA	1 µL of bisulfite treated DNA
Water	Adjust to 30 µL

Contact our office by email or by phone to place an order

©2024 EpigenDx. All Rights

EpigenDx, Inc
96 South Street, Hopkinton, MA 01740

Email: info@epigenDX.com

Tel: (508)-497-9400

Toll Free: (800)-941-6884

Fax: (508)-497-9450

Website: Labs.epigenDX.com

- HotStart Taq Polymerase Qiagen (Cat. #203205) recommended with the following PCR cycling conditions:

- 95°C 15 min; 45 x (95°C 30 s; T_a°C 30 s; 72°C 30 s); 72°C 5 min; 4°C ∞

- Additional optimization is needed if different PCR system is used in analysis.

- ▶ Sequencing Analysis: Pyrosequencing, NGS, or MS-HRM.

Technical Specifications

- ▶ 5µg DNA in TE buffer (10mM Tris-HCl, 1mM EDTA, pH 8.0)
- ▶ Store at -20°C, in aliquots, for 2 years. For best results, do not freeze/thaw an individual aliquot more than three times. For longer term storage -70°C is recommended.

Intended Use: FOR RESEARCH USE ONLY

Example Quality Control Results

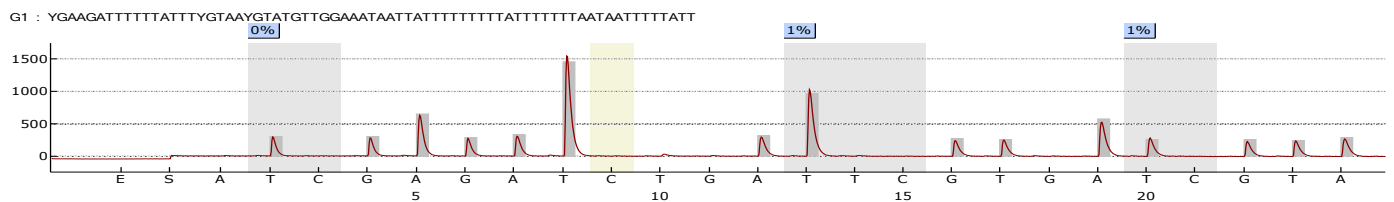


Figure: Low Methylated Control DNA Tested on a Human *BRCA1* Promoter Methylation Assay showing 0-1% methylation at all CpG sites