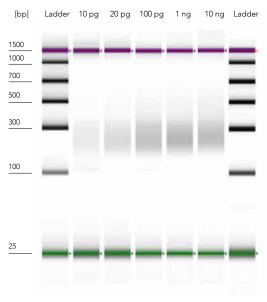


# Reliable Bisulfite Conversion for Next-Gen Sequencing

## EZ DNA Methylation-Lightning® Kits

- **Streamlined Process:** Ready-to-use conversion reagent is added directly to DNA. Purified bisulfite converted DNA in < 1.5 hours.
- **High-Quality:** Bisulfite-converted DNA has > 99.5% conversion efficiency with reduced fragmentation.
- **NGS-Ready:** Low DNA input requirement makes it ideal for preparing whole genome or targeted enrichment bisulfite libraries for methylation analysis.

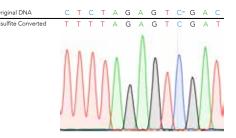
#### Compatible with Low DNA Inputs



Whole-genome bisulfite sequencing libraries were successfully prepared from DNA inputs as low as 10 pg using the EZ DNA Methylation – Lightning® Kit. Libraries were generated using the Zymo Research Pico Methyl-Seq™ Library Prep Kit, then analyzed using the Agilent 2200 Tapestation® system.

Référence	Désignation	Conditionnement
ZD5030T	EZ DNA Methylation-Lightning Kit	10 rxns
ZD5030	EZ DNA Methylation-Lightning Kit	50 rxns
ZD5031	EZ DNA Methylation-Lightning Kit	200 rxns
ZD5032	EZ-96 DNA Methylation-Lightning Kit (Shallow-Well)	2 x 96 rxns
ZD5033	EZ-96 DNA Methylation-Lightning Kit (Deep-Well)	2 x 96 rxns
ZD5046	EZ-96 DNA Methylation-Lightning MagPrep	4 x 96 rxns
ZD5047	EZ-96 DNA Methylation-Lightning MagPrep	8 x 96 rxns
Produits associés : les standards d'ADN methylé / non méthylé		
ZD5011	Universal Methylated Human DNA Standard	20 rxns
ZD5012	Universal Methylated Mouse DNA Standard	20 rxns
ZD5013	Human Methylated & Non-Methylated (WGA) DNA Set (DNA w/ primers)	1 kit
ZD5014	Human Methylated & Non-Methylated DNA Set (DNA w/ primers)	1 kit
ZD5015	Bisulfite-Converted Universal Methylated Human DNA Standard	50 rxns
ZD5016	E. coli Non-Methylated Genomic DNA	5 µg, 20µl
ZD5017	Methylated & Non-Methylated pUC19 DNA Set	1 set
ZD5101-2	Methylated / Non-Methylated Control DNA & Primer Set	1 set
ZD5018	Human Matched DNA Set	1 set

### High Conversion Efficiency



DNA with a methylated CpG (at nucleotide position 10) was processed using the EZ DNA Methylation-Lightning® Kit. The recovered DNA was amplified by PCR then sequenced. The methylated cytosine at position #10 remained intact, while the unmethylated cytosines (i.e. postiions #1, 3, and 13) were converted into uracil following bisulfite treatment and detected as thymine following PCR.

Specifications			
Input	100 pg - 2 µg purified DNA		
Conversion Efficiency	> 99.5%		
Elution Volume	Spin Column: ≥ 10 µl 96-well: ≥ 15 µl Magprep: ≥ 25 µl		
DNA Recovery	Spin Column: > 80% 96-well: > 70% Magprep: > 70%		
DI vi	Roche NimbleGen SeqCap™ Epi Enrichment System		
Platform Compatibility	Illumina TruSeq® DNA Methylation Kit Illumina Truseq® Methyl Capture EPIC Library Prep Kit		

#### **Nous contacter**



**Service client - commande** : commande@ozyme.fr **Service technique :** 

Réactifs: 01 34 60 60 24 - tech@ozyme.fr

Instrumentation: 01 30 85 92 88 - instrum@ozyme.fr