





EpiQuik™ Global Histone H3-K27 Methylation Assay Kit

Base Catalog # P-3020

PLEASE READ THIS ENTIRE USER GUIDE BEFORE USE

The *EpiQuik™* Global Histone H3-K27 Methylation Assay Kit is suitable for specifically measuring global histone H3-K27 methylation using variety of mammalian cells including fresh and frozen tissues, and cultured adherent and suspension cells.

Suitable lab coat, disposable gloves and eye protection are required when working with the kit.











KIT CONTENTS

Components	48 assays P-3020-48	96 assays P-3020-96
GG1 (10X Lysis Buffer)	5 ml	10 ml
GG2 (Extraction Buffer)	8 ml	16 ml
GG3 (10X Wash Buffer)	14 ml	28 ml
GG4 (Histone Buffer)	0.5 ml	1 ml
GG5 (Blocking Buffer)	10 ml	20 ml
GG6 (Antibody Buffer)	6 ml	12 ml
GG7 (Capture Antibody, 100 μ g/ml)*	25 <i>μ</i> Ι	$50~\mu$ l
GG8 (Detection Antibody, 400 μg/ml)*	10 <i>μ</i> Ι	20 <i>μ</i> Ι
GG9 (Developing Solution)	5 ml	10 ml
GG10 (Stop Solution)	3 ml	6 ml
Methylated H3-K27 Control (60 μg/ml)*	10 <i>µ</i> l	$20~\mu$ l
8-Well Assay Strips (with frame)	6	12
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^{*} For maximum recovery of the products, centrifuge the original vial prior to opening the cap.

SHIPPING & STORAGE

Upon receipt: (1) Store GG8 and Methylated H3-K27 Control at -20° C away from light; (2) Store GG3, GG5, GG7, GG9, and 8-Well Assay Strips at 4° C away from light; (3) Store all other components at room temperature. The kit is stable for up to 6 months from the shipment date, when stored properly.

Note: Check if wash buffer, GG3, contains salt precipitates before using. If so, warm (at room temperature or $37^{\circ}C$) and shake the buffer until the salts are re-dissolved.

MATERIALS REQUIRED BUT NOT SUPPLIED

Orbital	shaker

☐ Pipettes and pipette tips

☐ Microplate reader

□ 1.5 ml microcentrifuge tubes

☐ 60 or 100 mm plate

□ Dounce homogener

□ 100% TCA solution

☐ Glycerol

☐ Acetone











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5%	П	10.1

■ Distilled water

GENERAL PRODUCT INFORMATION

Usage Limitation: The $EpiQuik^{\mathsf{TM}}$ Global Histone H3-K27 Methylation Assay Kit is for research use only and is not intended for diagnostic or therapeutic application.

Quality Control: Epigentek guarantees the performance of all products in the manner described in our product instructions.

Product Updates: Epigentek reserves the right to change or modify any product to enhance its performance and design.

Intellectual Property: The $EpiQuik^{\mathsf{TM}}$ Global Histone H3-K27 Methylation Assay Kit and methods of use contain proprietary technologies by Epigentek. $EpiQuik^{\mathsf{TM}}$ is a trademark of Epigentek, Inc.

A BRIEF OVERVIEW

Epigenetic activation or inactivation of genes play a critical role in many important human diseases, especially in cancer. A major mechanism for epigenetic inactivation of the genes is methylation of CpG islands in genome DNA caused by DNA methyltransferases. Histone methyltransferases (HMTs) control or regulate DNA methylation through chromatin-dependent transcription repression or activation. HMTs transfer 1-3 methyl groups from S-adenosyl-L-methionine to the lysine and arginine residues of histone proteins. G9a and polycomb group enzyme such as EZH2 are histone methyltransferases that catalyze Methylation of histone H3 at lysine 27 (H3-K27) in mammalian cells. H3-K27 methylation mediates heterochromatin formation by forming a binding site for polycomb and also participates in silencing gene expression at euchromatic sites. Increased global H3-K27 methylation is also found to be involved in some pathological processes such as cancer progress. There are only a couple of methods such as western blot used for measuring histone H3-K27 methylation. The methods available so far are time consuming and labor intensive, or have low throughput. The EpiQuik™ Global H3-K27 Methylation Assay Kit addresses these problems by using a unique procedure to measure global methylation of histone H3-K27. The kit has the following features:

- Quick and efficient procedure, which can be finished within 5 hours.
- Innovative colorimetric assay without the need for radioactivity, electrophoresis, or chromatography.
- Strip microplate format makes the assay flexible: manual or high throughput.
- Simple, reliable, and consistent assay conditions.

PRINCIPLE & PROCEDURE

The EpiQuik™ Global H3-K27 Methylation Assay Kit is designed for measuring global histone H3-K27 methylation. In an assay with this kit, the histone proteins are stably spotted on the strip wells.









The methylated histone H3-K27 can be recognized with a high-affinity antibody. The ratio or amount of methylated H3-K27 can be quantified through HRP conjugated secondary antibody-color development system and is proportional to the intensity of color development.



Schematic Procedure for Using the EpiQuik™ Global Histone H3-K27 Methylation Assay Kit

PROTOCOL

Nuclear Extraction Preparation

For Tissue Samples:

- 1. Place the tissue sample into a 60 or 100 mm plate. Remove unwanted tissue such as fat and necrotic material from the sample. Weigh the sample and cut it into small pieces (1-2 mm³) with a scalpel or scissors.
- Transfer tissue pieces to a Dounce homogener. Dilute GG1 with distilled water at a 1:10 ratio.
 Add 1 ml of diluted GG1 per every 200 mg of tissue and disaggregate tissue pieces by 10-30 strokes.
- 3. Transfer homogenized mixture to a 15 ml conical tube and centrifuge at 3000 rpm for 5 minutes at 4°C. If total mixture volume is less than 2 ml, transfer mixture to a 2 ml vial and centrifuge at 10,000 rpm for 1 minute at 4°C. Remove supernatant.

For Adherent Cells:

1. Cells (treated or untreated) are grown to 70-80% confluency, then trypsinized and collected into a 15 ml conical tube. Count cells in a hemacytometer.



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- 2. Centrifuge the cells at 1000 rpm for 5 minutes and discard the supernatant. Wash cells with 10 ml of PBS once by centrifugation at 1000 rpm for 5 minutes. Discard the supernatant.
- 3. Add **diluted GG1** to re-suspend cell pellet (200 μ l/1 x 10⁶ cells). Transfer cell suspension to a 1.5 ml vial and incubate on ice for 5 minutes and vortex occasionally.
- 4. Pellet cell debris by centrifuging at 12,000 rpm for 30 seconds.

For Suspension Cells:

- 1. Collect cells (treated or untreated) into a 15 ml conical tube. $(1-2 \times 10^6 \text{ cells are required for each reaction})$. Count cells in a hemacytometer.
- 2. Centrifuge the cells at 1000 rpm for 5 minutes and discard the supernatant. Wash cells with 10 ml of PBS once by centrifugation at 1000 rpm for 5 minutes. Discard the supernatant.
- 3. Add **diluted GG1** to re-suspend cell pellet (100 μ l/1 x 10⁶ cells). Transfer cell suspension to a 1.5 ml vial and incubate on ice for 5 minutes, and vortex occasionally.
- 4. Pellet cell debris by centrifuging at 12,000 rpm for 30 seconds.

Histone Extraction

- 1. Add glycerol to GG2 at a 1:10 ratio (ex: add 100 μ l of glycerol to 900 μ l of GG2) to prepare the GG2/Glycerol solution. Add diluted GG1 to cell debris (10 μ l/1 x 10⁶ cells or 40 mg of tissue), followed by adding 3 volumes of the GG2/glycerol solution. Mix by vortex and incubate on ice for 5 minutes.
- 2. Pellet nucleic debris by centrifuging at 12,000 rpm for 5 minutes at 4°C. Transfer the supernatant to a 1.5 ml vial.
- 3. Add 100% TCA to the supernatant at a 1:4 ratio (ex: add 100 μ l of TCA to 300 μ l of supernatant; final concentration of TCA should be 25%). Incubate on ice for 30 minutes.
- 4. Collect the precipitate by centrifuging at 12,000 rpm for 10 minutes at 4°C.
- 5. Remove supernatant and add 1 ml of acetone containing 0.1% HCl to precipitate. Mix and incubate on ice for 1 minute.
- 6. Collect the pellet by centrifuging at 12,000 rpm for 2 minutes at 4°C. Wash the pellet with 1 ml of acetone. Allow 1 minute on ice for wash.
- 7. Collect the pellet by centrifuging at 12,000 rpm for 2 minutes at 4°C. Remove the supernatant as much as possible and air dry the pellet for 5 minutes.
- 8. Add distilled water to dissolve pellet (10 μ l of water per amount of pellet extracted from 1 x 10⁶ cells or 40 mg of tissue) and measure histone protein concentration. The histone extract can be used immediately or stored at -20° C.

Histone H3-K27 Methylation Detection

- 1. Determine the number of strip wells required. Leave these strips in the plate frame (remaining unused strips can be placed back in the bag. Seal the bag tightly and store at 4°C). Dilute **GG3** with distilled water (pH 7.2-7.5) at a 1:10 ratio.
- 2. Adjust protein concentration to 200 ng/ μ l or 400 ng/ μ l with GG4 and add 5 μ l (1-2 μ g) of the protein solution into the central area of each well. Spread out the solution over the bottom of the strip well by pipetting the solution up and down several times, and incubate at 37°C (with no humidity) for 60-90 minutes to evaporate the solution and dry the wells. For the blank, add 5 μ l of GG4 to the wells. For the positive control, dilute Methylated H3-K27 Control to 2-30 ng/ μ l with











GG4, and then add 5μ l (10-150 ng) of the **diluted Methylated H3-K27 Control solution** to the wells.

- 3. Add 150 μ l of **GG5** to the dried wells and incubate at 37°C for 30-45 minutes.
- 4. Aspirate and wash the wells with 150 μ l of **diluted GG3** three times.
- 5. Dilute **GG7** (at a 1:100 ratio) to 1 μ g/ml with **GG6**. Add 50 μ l of the **diluted GG7** to the wells and incubate at room temperature for 60 minutes on an orbital shaker (50-100 rpm).
- 6. Aspirate and wash the wells with 150 μ l of **diluted GG3** four times.
- 7. Dilute **GG8** (at a 1:1000 ratio) to 0.4 μ g/ml with **GG6**. Add 50 μ l of the **diluted GG8** to the wells and incubate at room temperature for 30 min.
- 8. Aspirate and wash the wells with 150 μ l of **diluted GG3** five times.
- 9. Add 100 μ l of **GG9** to the wells and incubate at room temperature for 2-10 minutes away from light. Monitor the color development in the sample and standard wells (blue).
- 10. Add 50 μ l of **GG10** to the wells and read absorbance on microplate reader at 450 nm.
- 11. Calculate % H3-K27 methylation:

$$\mbox{Methylation \% = } \frac{\mbox{OD (sample - blank)}}{\mbox{OD (untreated control - blank)}} \times 100\%$$

For accurate calculation, plot OD value versus amount of **Methylated H3-K9 Control** and determine the slope as delta OD/ng.

Calculate the amount of methylated H3-K27 using the following formula:

Amount (ng/mg protein) =
$$\frac{\text{OD (sample - blank)}}{\text{slope}} \times 1000$$

TROUBLESHOOTING

No Signal for Both the Positive Control and the Samples

Reagents are added incorrectly. Check if reagents are added in order and if

any steps of the procedure may have been

omitted by mistake.

Incubation time and Ensure the incubation time and temperature temperature are incorrect.

Ensure the incubation time and temperature described in the protocol are followed correctly.

No Signal or Very Weak Signal for Only the Positive Control

The positive control is added insufficiently.

Ensure a sufficient amount of control is added to the well.



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The positive control is degraded due to incorrect storage.

Follow the guidance in the protocol for storage of positive control.

No Signal for Only the Sample

The protein sample is not properly extracted.

Follow the protocol instructions for the histone protein extraction.

The protein amount is added into well insufficiently.

Ensure extract contains a sufficient amount of protein.

Protein extracts are incorrectly stored or were stored for a long time. Ensure the protein extracts are stored at -80°C for no more than 8 weeks.

High Background Present for the Blank

The well is not washed sufficiently.

Check if wash at each step is performed according to the protocol.

Contaminated by the positive control.

Ensure the well is not contaminated from adding the control protein or from using control protein contaminated tips.

Overdevelopment.

Decrease the development time at step 9 of "Histone H3-K27 Methylation Detection."

RELATED PRODUCTS

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P-3015	EpiQuik™ In Situ Histone H3-K4 Methylation assay Kit
P-3016	EpiQuik™ In Situ Histone H3-K9 Methylation Assay Kit
P-3017	EpiQuik™ Global Histone H3-K4 Methylation Assay Kit
P-3018	FniQuik™ Global Histone H3-K9 Methylation Assay Kit



