

Technical Bulletin

## Lipid Peroxidation (MDA) Assay Kit

#### **Catalog Number MAK085**

## **Product Description**

Lipid peroxidation is the degradation of lipids that occurs as a result of oxidative damage and is a useful marker for oxidative stress. Polyunsaturated lipids are susceptible to an oxidative attack, typically by reactive oxygen species, resulting in a well-defined chain reaction with the production of end products such as malondialdehyde (MDA). Lipid peroxidation may contribute to the pathology of many diseases including atherosclerosis, diabetes, and Alzheimer's.

In this kit, lipid peroxidation is determined by the reaction of MDA with thiobarbituric acid (TBA) to form a colorimetric (532 nm) / fluorometric ( $\lambda_{Ex} = 532/\lambda_{Em} = 553$  nm) product, proportional to the MDA present.

## Components

The kit is sufficient for 100 colorimetric or fluorometric assays in 96-well plates.

•	MDA Lysis Buffer Catalog Number MAK085A	25 mL
•	Phosphotungstic Acid Solution Catalog Number MAK085B	12.5 mL
•	BHT, 100× Catalog Number MAK085C	1 mL
•	TBA Catalog Number MAK085D	4 bottles
•	MDA Standard, 4.17 M Catalog Number MAK085E	0.1 mL

# Reagents and Equipment Required but Not Provided

- Pipetting devices and accessories (e.g., multichannel pipettor)
- 96-well flat-bottom plate It is recommended to use black plates with clear bottoms for fluorescence assays and clear plates for colorimetric assays. Cell culture or tissue culture treated plates are **not** recommended.
- Fluorescence or spectrophotometric multiwell plate reader
- Glacial acetic acid (Catalog Number A6283 or equivalent)
- Perchloric acid (Catalog Number 244252 or equivalent)
- Sulfuric acid (Catalog Number 258105 or equivalent)
- 1-Butanol (Catalog Number 360465 or equivalent)

#### Precautions and Disclaimer

For R&D Use Only. Not for drug, household, or other uses. Please consult the Safety Data Sheet for information regarding hazards and safe handling practices.

## Storage/Stability

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The kit is shipped on wet ice. Store components at -20 °C, protected from light.



## **Preparation Instructions**

Briefly centrifuge vials before opening. To maintain reagent integrity, avoid repeated freeze/thaw cycles. Use purified water for the preparation of all reagents. Allow all components to come to room temperature before starting.

TBA Solution: Reconstitute a bottle with 7.5 mL of Glacial Acetic Acid, then adjust the final volume to 25 mL with water. Sonication can be used to assist dissolution if necessary. Store at 4 °C and use within 1 week of preparation.

#### Procedure

All samples and standards should be run in duplicate.

Use purified water for the preparation of all standards and samples.

#### MDA Standards for Colorimetric Detection

- 1. A new standard curve must be set up each time the assay is run.
- 2. Dilute 10  $\mu$ L of the 4.17 M MDA Standard Solution with 407  $\mu$ L of purified water to prepare a 0.1 M MDA Standard Solution.
- 3. Further dilute 20  $\mu$ L of the 0.1 M MDA Standard Solution with 980  $\mu$ L of purified water to prepare a 2 mM MDA Standard.
- 4. Prepare MDA standards in 1.5 mL microcentrifuge tubes according to Table 1.

**Table 1.**Preparation of MDA Standards for Colorimetric Assay

Well	2 mM MDA Standard	Purified Water	MDA (nmole/well)
1	10 μL	190 µL	20
2	8 µL	192 μL	16
3	6 µL	194 µL	12
4	4 µL	196 µL	8
5	2 μL	198 µL	4
6	-	200 μL	0

#### MDA Standards for Fluorometric Detection

- 1. A new standard curve must be set up each time the assay is run.
- 2. Prepare a 2 mM Standard Solution as for the Colorimetric assay Steps 1 and 2.
- 3. Dilute 100  $\mu$ L of the 2 mM MDA Standard Solution with 900  $\mu$ L of water to make a 0.2 mM MDA standard solution.
- 4. Prepare MDA standards in 1.5 mL microcentrifuge tubes according to Table 2.

**Table 2.**Preparation of MDA Standards for Fluorometric Assay

Well	0.2 mM MDA Standard	Purified Water	MDA (nmole/well)
1	10 μL	190 µL	2.0
2	8 µL	192 µL	1.6
3	6 µL	194 µL	1.2
4	4 µL	196 µL	0.8
5	2 µL	198 µL	0.4
6	_	200 μL	0

## Sample Preparation

#### Serum or Plasma

- 1. Gently mix 20  $\mu$ L of serum or plasma samples with 500  $\mu$ L of 42 mM sulfuric acid in a microcentrifuge tube.
- 2. Add 125  $\mu$ L of Phosphotungstic Acid Solution and mix by vortexing.
- 3. Incubate at room temperature for 5 minutes and then centrifuge the samples at  $13,000 \times g$  for 3 minutes. Remove supernatant and retain pellet for assay.
- 4. In a separate tube, add 2  $\mu L$  of BHT (100×) to 100  $\mu L$  of purified water.
- 5. Resuspend the pellet on ice with the water/BHT solution. Adjust the volume to 200 µL with water.

#### Tissue or Cells

- 1. Homogenize tissue (10 mg) or cells (2  $\times$  10<sup>6</sup>) on ice in 300  $\mu$ L of the MDA Lysis Buffer containing 3  $\mu$ L of BHT (100 $\times$ ).
- 2. Centrifuge the samples at  $13,000 \times g$  for 10 minutes to remove insoluble material.
  - Alternatively, protein can be precipitated by homogenizing 10 mg of sample in 150  $\mu L$  of purified water containing 3  $\mu L$  of BHT (100×) and adding 1 volume of 2 N perchloric acid, vortexing, and centrifuging to remove precipitated protein.
- 3. Place 200 µL of the supernatant from each homogenized Sample into a microcentrifuge tube.

#### **Assay Reaction**

- 1. To form the MDA-TBA adduct, add 600  $\mu L$  of the TBA solution into each vial containing Standard and Sample.
- 2. Incubate at 95 °C for 60 minutes.
- 3. Cool to room temperature in an ice bath for 10 minutes.
- Pipette 200 μL from each Standard and Sample reaction mixture, except for Serum or Plasma Samples (see Step 5), into a 96-well plate for analysis.

#### Note:

- a. To enhance sensitivity, add 300  $\mu$ L of 1-butanol to extract the MDA-TBA adduct from the 800  $\mu$ L reaction mixture. If separation does not occur, add 100  $\mu$ L of 5 M NaCl and vortex vigorously. Centrifuge at 16,000  $\times$  g for 3 minutes at room temperature to separate the layers. Transfer the 1-butanol layer (the top layer) to another tube and evaporate the 1-butanol. The 1-butanol can be removed either by freeze-drying or heating on a hot block at 55 °C. Dissolve the residue containing the MDA-TBA adduct in 200  $\mu$ L of purified water, and then transfer to a 96-well plate for analysis.
- b. Occasionally samples will exhibit turbidity, which can be eliminated by filtering through a 0.2  $\mu m$  filter.
- TBA can react with other compounds in Samples to produce other colored products.
  These should not generally interfere with quantitation of the TBA-MDA adduct.

5. For Serum or Plasma Samples, mix each reaction mixture with 300  $\mu$ L of 1-butanol and 100  $\mu$ L of 5 M NaCl. Vortex and then centrifuge for 3 minutes at 16,000  $\times$  g at room temperature. Transfer the 1-butanol layer (the top layer) to a new centrifuge tube and remove the 1-butanol. The 1-butanol can be removed either by freezedrying, or heating on a hot block at 55 °C. Resuspend the remaining material in 200  $\mu$ L of ultrapure water. Mix well and add 200  $\mu$ L into a 96-well plate.

#### Measurement

For colorimetric assays, measure the absorbance (A) at 532 nm. For fluorometric assays, measure relative fluorescence units (RFU) at  $\lambda_{Ex} = 532$  nm  $/\lambda_{Em} = 553$  nm).

## Results

- 1. Calculate  $\Delta A$  /  $\Delta RFU$  by subtracting the A / RFU reading of Standard #6 (Blank) from the remaining Standard reading values. Background values can be significant and must be subtracted from all readings. The amount of MDA present in the samples may be determined from the standard curve.
- 2. Plot the  $\Delta A$  /  $\Delta RFU$  values against Standard concentrations and determine the slope of the standard curve.
- 3. Using the standard curve determine the quantity of MDA in nmole in the Sample.
  - a. For Samples without the 1-butanol concentration step, calculate the MDA concentration of the Sample:

MDA (nmol/mL) =

$$(S_A/S_V) \times DF = C$$

where:

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- S<sub>A</sub> = Amount of MDA in Sample (nmole) as determined from the standard curve
- S<sub>V</sub> = Sample volume (mL) or amount (mg) added into the wells
- DF = Sample dilution factor (DF = 1 for undiluted Samples)
- C = Concentration of MDA in sample

#### **Example Calculation**

Amount of MDA  $(S_A) = 5.84$  nmole

Sample volume  $(S_V) = 0.020 \text{ mL}$ 

Concentration of MDA in sample =

 $(5.84 \text{ nmole}/0.020 \text{ mL}) \times 1 = 292 \text{ nmole/mL}$ 

b. For Samples with the 1-butanol concentration step, calculate the MDA concentration of the Sample:

MDA (nmol/mL) =

$$(S_A/S_V) \times 4 \times DF = C$$

#### where:

- S<sub>A</sub> = Amount of MDA in Sample (nmole) as determined from the standard curve
- $S_V =$  Sample volume (mL) or amount (mg) added into the wells
- 4 = Correction factor for using 200  $\mu$ L of the 800  $\mu$ L reaction
- DF = Sample dilution factor (DF = 1 for undiluted Samples)
- C = Concentration of MDA in sample

#### **Example Calculation**

Amount of MDA  $(S_A) = 5.84$  nmole

Sample volume  $(S_V) = 0.020 \text{ mL}$ 

Concentration of MDA in sample

 $(5.84 \text{ nmole}/0.020 \text{ mL}) \times 4 \times 1 = 1,168 \text{ nmole}/\text{mL}$ 

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