

**Northwest**  
Life Science Specialties, LLC

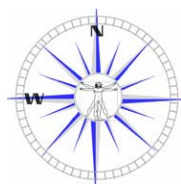
Premier Products for Superior Life Science Research

*NWLSSTM*  
*Malondialdehyde Assay*

Product NWK-MDA01  
*For Research Use Only*



Simple assay kit for quantitative measurement of lipid peroxidation as Thiobarbituric Acid Reactive Substances (TBARS) or more specifically, as malondialdehyde (MDA) when employing extraction or advanced data reduction techniques.



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## Table of Contents

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Section	Page
Introduction	3
Intended Use	3
Test Principle	3
General Specifications	4
Kit Contents	4
Required Materials Not Provided	4
Required Instrumentation	4
Warnings & Precautions	4
Storage Instructions	5
Assay Preparation	5
Reagent Preparation	5
Sample Handling/Preparation	5
Assay Protocol	6
Data Analysis	6
Performance Details	8
Things To Note	10
References	10
Procedure Checklist	11
Statement of Limited Warranty	

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### Procedure Checklist:

\_\_\_Process the sample and place on ice

\_\_\_Preheat heat block or water bath to 60 °C

\_\_\_Reconstitute TBA Reagent with 10.5 mL deionized water

\_\_\_Add 10 µL BHT Reagent to microcentrifuge vial

\_\_\_Add 250 µL Calibrator or sample to microcentrifuge vial

\_\_\_Add 250 µL Phosphoric Acid Reagent to microcentrifuge vial

\_\_\_Add 250 µL TBA reagent to microcentrifuge vial

\_\_\_Cap and vortex (to 5 count)

\_\_\_Place at 60 °C for 60 minutes

\_\_\_Setup spectrophotometer

\_\_\_Centrifuge at 10,000 xg for 2-3 minutes

\_\_\_Scan each reaction mixture; 400-700 nm

\_\_\_Analyze data

### Statement of Limited Warranty:

Northwest Life Science Specialties, LLC (NWLSS) makes no guarantee of any kind, expressed or implied, that extends beyond the description of the material in this kit, except that they will meet our specifications at the time of delivery. Customer's remedy and NWLSS' sole liability is limited to, at NWLSS' option, refund of the purchase price, or the replacement of material not meeting our specification. By acceptance of our product, customer assumes all liability and will indemnify and hold NWLSS harmless for the consequence of this product's use or misuse by the customer, its employees, or others. Refund or replacement is conditioned of customer notifying NWLSS within twenty-one (21) days of the receipt of product. Failure to give notice within 21 days shall constitute a waiver by the customer of all claims hereunder with respect to said product.

**Performance Details (continued)****Stability**

All unopened reagents are stable until the expiration date stated on the package label when stored at 2-8 °C.

**Things to Notes:**

**Extraction:** Samples that contain a significant amount of particles and will have an increased baseline caused by light scattering. This baseline can be significantly reduced by extraction of the reaction mixture into an organic solvent. The TBA2-MDA adduct partitions into the aqueous (upper) phase of the CHCl<sub>3</sub> extract. Butanol has also been used to clarify the reaction mixture.<sup>6</sup> In this case, the TBA2-MDA adduct is extracted into the butanol; again the upper phase. The advantages of CHCl<sub>3</sub> is the results of the extraction is not dependent on pipetting precision and is not flammable. Butanol, produces a slightly cleaner extract and can be used to concentrate the TBA2-MDA adduct but butanol pipetting impression will affect the result.

**Derivative Primer:** Derivative spectroscopy can appear intimidating. However, most spectrophotometric software includes derivative analysis which greatly simplifies data reduction. An excellent primer is available from Agilent Technologies, Publication 5963-3940E at [www.chem.agilent.com/scripts/LiteraturePDF.asp?iWHID=15886](http://www.chem.agilent.com/scripts/LiteraturePDF.asp?iWHID=15886). Please contact NWLSS for additional assistance if necessary.

**Derivative Analysis Peak Shift:** The 3<sup>rd</sup> derivative of a biological sample may show a slight peak shift relative to the calibrators; 556 nm and 550 nm respectively in the example. Because of the large wavelength filter (29 nm), the broad derivative peak minimizes the error.

**Alternative Analysis:** The reaction mixture can also be analyzed using difference absorption spectroscopy, fluorescence spectroscopy or HPLC.<sup>8,9,10</sup>

**TBA Reagent Precipitation:** If the TBA Reagent is refrigerated, the TBA will precipitate. One can re-dissolve the TBA with gentle warming. If the precipitate does not re-dissolve, discard the reagent.

**References:**

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**Introduction:**

Polyunsaturated fatty acids (PUFA) that contain 2 or more double bonds are particularly susceptible to oxidation by free radicals and other highly reactive species. In brief, an allylic hydrogen is abstracted by a reactive species, such as the hydroxyl radical (HO<sup>•</sup>), resulting in the formation of lipid peroxy radicals (LOO<sup>•</sup>). This radical can then react with a second PUFA, forming a lipid hydroperoxide (LOOH) and a second LOO<sup>•</sup>, resulting in the propagation of the lipid oxidation. Alternatively, LOO<sup>•</sup> can attack an intramolecular double bond and form a cyclic endoperoxide which decomposes to malondialdehyde.<sup>1,2</sup>

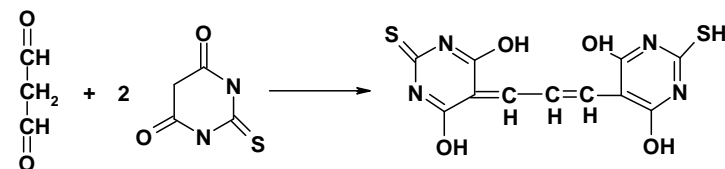
Malondialdehyde (MDA) is one of many low molecular weight end-products of lipid hydroperoxide decomposition and is the most often measured as an index of lipid peroxidation.<sup>3</sup> However, the use of MDA as a marker for lipid peroxidation is controversial. MDA can be formed during eicosanoid metabolism and the analytical methods for measuring MDA are prone to artifactual errors.<sup>4</sup> The NWLSS™ MDA assay conditions have been designed to minimize errors in the determination of MDA concentration.

**Intended Use:**

The NWLSS™ Malondialdehyde Assay is used to measure the concentration of MDA in a variety of animal and plant samples.

**Test Principle:**

The NWK-MDA01 assay is based on the reaction of MDA with thiobarbituric acid (TBA); forming an MDA-TBA<sub>2</sub> adduct that absorbs strongly at 532 nm.



**Figure 1.**  
Reaction between MDA and TBA.

Butylated hydroxytoluene (BHT) and EDTA are added to the sample and reaction mixture to minimize oxidation of lipids that contribute artifactually during sample processing and the TBA reaction.<sup>5,6</sup> The temperature of the reaction mixture has also been reduced to minimize the decomposition of lipid hydroperoxides. Because much of the MDA is protein bound, mostly as a Schiff base, the pH of the reaction has been optimized to facilitate hydrolysis of the MDA.<sup>7</sup> Additionally, the reaction mixture is subjected to derivative spectrophotometric analysis that resolves the problem of the variable and nonlinear baseline observed when attempting to measure the A<sub>532</sub> absorbance in various biological samples (see Data Analysis).

**General Specifications:**

Format: Spectrophotometric  
 Number of tests: 200  
 Specificity: Malondialdehyde  
 Sensitivity: Processed Sample = 0.08  $\mu\text{M}$   
 Reaction Mixture = 0.03  $\mu\text{M}$

**Kit Contents:**

TBA Reagent 2-Thiobarbituric acid, dry powder, 5 bottles  
 BHT Reagent Butylated hydroxytoluene in ethanol, 2.0 mL vial  
 Acid Reagent 1M Phosphoric acid, 5 x 10 mL bottles  
 Assay Buffer Phosphate buffer, pH 7.0 with EDTA, 125 mL bottle  
 Calibrators: Tetramethoxypropane in a stabilizing buffer at 4,3,2,1 and 0  $\mu\text{M}$  MDA equivalents; 2.5 mL each

**Required Materials Not Provided:**

Adjustable pipettes capable of 10 and 250  $\mu\text{L}$  volumes  
 Pipette tips for 10 and 250  $\mu\text{L}$  volumes  
 Stir bar, 1/8 inch (0.3 cm) or similar  
 Polypropylene microcentrifuge tubes  
 Semi-micro cuvettes (plastic glass or quartz)  
 Deionized water

**Required Instrumentation:**

Spectrophotometer  
 Vortex mixer  
 Magnetic stirrer (recommended)  
 Dry heating block or water bath set to 60°C  
 Microcentrifuge

**Warnings & Precautions:**

2-Thiobarbituric Acid (TBA, CAS 504-17-6) has a strong mercaptan odor. Do not breathe dust. Avoid contact with skin and eyes.

Phosphoric Acid (CAS 7664-38-2) is corrosive and can cause burns. In case of contact with eyes, rinse immediately with copious amounts of water and seek medical advice. Wear suitable protective clothing.

Butylated hydroxytoluene (2,6-di-tert-butyl-4-methylphenol, BHT, CAS 128-37-0) is harmful if swallowed and irritating to the eyes, respiratory system and skin. In case of contact with eyes, rinse immediately with copious amounts of water and seek medical advice. Wear protective clothing.

**Performance Details (continued)****Accuracy**

Recovery: The recovery of MDA (hydrolyzed TMOP) and TMOP added to a muscle homogenate is shown in Table 2.

		Initial	Observed	Added	Recovery
Buffer	TMOP	0	1.91	2.00	95%
	MDA	0	1.83	1.99	92%
Muscle	TMOP	0.50	1.87	1.91	72%
	MDA	0.50	1.81	1.83	71%

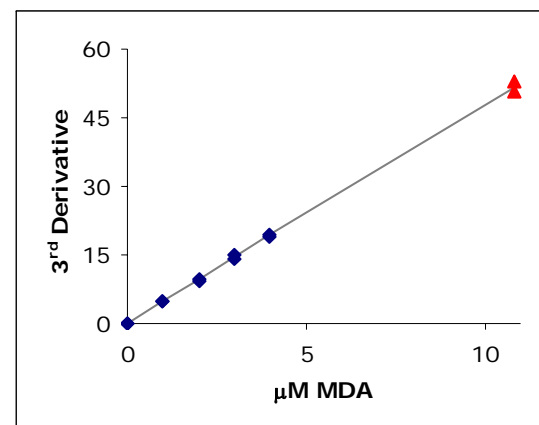
**Table 2.**

Recovery of MDA added to muscle homogenates and Assay Buffer.

The recovery of less than 100% can be attributed to the tendency of MDA to undergo several reactions in a complex sample that are not reversible by acid hydrolysis.<sup>1</sup>

**Dynamic Range:**

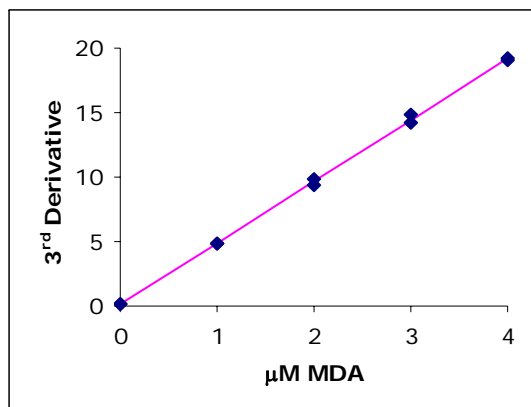
The linearity and sensitivity of the calibration curve defines the useful range of the assay. As can be seen in Figure 5, the calibration curve is linear to 11  $\mu\text{M}$  MDA. The standard method can measure up to 50  $\mu\text{M}$  MDA but those concentrations will not be found in most biological samples. Therefore, the useful range of the NWLSS MDA assay is 0.1 to 10  $\mu\text{M}$ .

**Figure 5.**

The 3<sup>rd</sup> derivative of a 10.8  $\mu\text{M}$  MDA sample (▲) was determined using the standard method and plotted against the calibration curve (◆).

**Data Analysis (continued):**

Plotting the third derivative peak at 514 nm as a function of MDA concentration produces a linear calibration curve as shown in Figure 4.



**Figure 4.**  
Example calibration curve.

Simple linear regression of the 3<sup>rd</sup>-derivative vs. μM MDA yields the calibration equation:

$$\frac{d^3A}{d\lambda^3} = 4.777[\text{MDA}] + 0.1167$$

where

$$\begin{aligned} \text{Slope} &= 4.777 \\ \text{Intercept} &= 0.1167 \\ r^2 &= 0.9993 \\ \text{Syx} &= 0.2280 \end{aligned}$$

Continuing the example, the 3<sup>rd</sup>-derivative for the muscle homogenate sample was 1.831. Solving for [MDA]...

$$[\text{MDA}] = \frac{1.831 - 0.117}{4.777} = 0.359 \mu\text{M}$$

**Performance Details:****Sensitivity**

The sensitivity of the NWK-MDA01 assay is defined as the MDA concentration obtained from 2-times the standard error of the calibration curve or 0.08 μM in the prepared sample or 0.03 μM in the reaction mixture.

**Storage Instructions:**

Store all reagents at 2-8 °C.

Reconstituted TBA Reagent is stable for at least 1 week when stored at room temperature. **Do not refrigerate.**

**Assay Preparation**

Spectrophotometer

Wavelength: 400-700 nm

Spectral bandwidth: ≤ 2 nm

Resolution: ≤ 1 nm

Heat block or water bath preheated to 60 °C

**Reagent Preparation:****TBA Reagent**

Add 10.5 mL deionized water to the TBA bottle. Insert magnetic stir bar and mix until TBA has dissolved; approximately 10 minutes. Store at room temperature, do not refrigerate.

**Other Reagents**

The Acid Reagent, BHT Reagent, Assay Buffer and Calibrators are supplied ready-to-use.

**Sample Handling/Preparation:**

The multi-disciplinary interest in measuring MDA has resulted in a myriad of sample types and experimental conditions and is beyond the scope of this product insert to describe sample processing in detail. However, general guidelines are provided below for representative sample types. Please contact NWLSS to discuss the particular sample under investigation.

**Tissue**

Hemoglobin may interfere with the assay; therefore, if practical, blood should be removed by perfusion with an appropriate buffer, such as phosphate buffered saline containing heparin. Prepare a 10% w/v homogenate in cold Assay Buffer or other buffer. Clarify the homogenate by centrifugation and store the supernatant on ice. The MDA concentration can be normalized to the wet weight of the tissue sample or to the protein concentration of the homogenate.

**Plasma**

Blood, using EDTA, heparin or other anticoagulant, should be kept cold and the plasma separated as soon as possible after being drawn. It is important to minimize hemolysis. Plasma can be assayed without further processing. Note that the MDA concentration in normal plasma is at the limits of sensitivity for the assay.

**Sample Handling/Preparation (continued):****Urine**

Urine samples should be stored at -20 °C (-70 °C long term storage). Upon thawing, the sample must be clarified by centrifugation prior to assay. The MDA concentration can be normalized to the creatinine concentration or other appropriate marker. NWLSS™ NWK-CTNO1 can be used to measure creatinine.

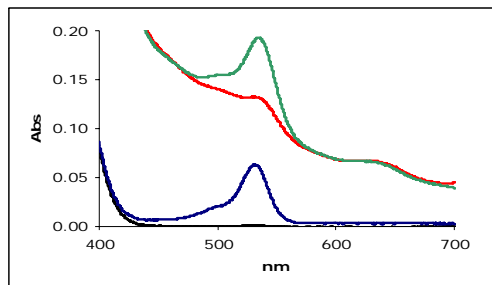
**Assay Protocol:**

1. Add 10 µL BHT Reagent to microcentrifuge vial
2. Add 250 µL Calibrator or sample to vial
3. Add 250 µL Acid Reagent to vial
4. Add 250 µL TBA Reagent to vial
5. Vortex vigorously (5-count)
6. Incubate 60 minutes at 60 °C
7. Centrifuge at 10,000 xg for 2-3 minutes
8. Transfer reaction mixture to cuvette
9. Record spectra from 400-700 nm
10. Perform 3<sup>rd</sup> derivative analysis (recommended) or other analytical method.

**Data Analysis:**

The following example shows the expected behavior and results of MDA determinations on a biological sample using the standard method of the NWLSS™ MDA assay.

A sample of bovine muscle was homogenized in Assay Buffer. To enhance the signal for this example, a known amount of MDA (hydrolyzed TMOP) was added to the muscle homogenate and then assayed using the standard method. **Figure 2** shows the zero order plots of the reaction mixtures including the 1 µM MDA Calibrator and blank for reference (the other calibrators and replicates are not shown for clarity).

**Figure 2.**

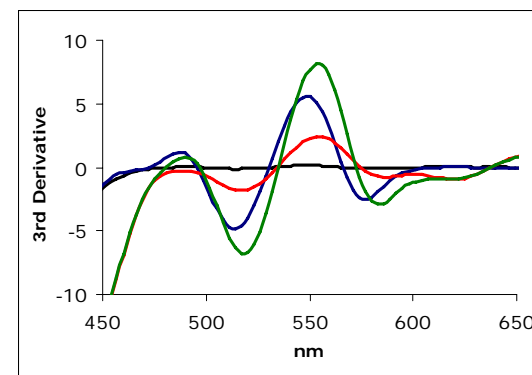
Zero order plot of MDA Calibrator (■), muscle homogenate (■), muscle homogenate with added MDA (■) and blank (■).

**Data Analysis (continued):**

As can be seen in the figure 2, the muscle homogenate shows a severe non-linear baseline making it difficult to estimate the A<sub>532</sub> absorbance for the homogenate. This is typical of TBA reactions with biological samples.

Therefore the preferred method to calculate the MDA concentration in biological samples is by derivative analysis of the spectra for the reaction mixture.<sup>5</sup> Derivative spectroscopy applications are provided by most spectrophotometer software. These data were analyzed using the Math Application supplied in the Cary Win UV version 3.0 software. The Filter Size was set to 29 nm and the Interval set to 3 nm. For convenience, the derivative is multiplied by -1x10<sup>6</sup> to provide a more convenient number.

Figure 3 shows the plot of the 3<sup>rd</sup>-derivative as a function of wavelength. As can be seen, the derivative easily resolves the MDA-TBA<sub>2</sub> peak.

**Figure 3.**

Plot of the 3<sup>rd</sup> derivatives of scans from Figure 2

Calibration Curve		
[MDA]	$\frac{d^3A}{dx^3}$	
	1	2
0	0.133	0.136
1	4.807	4.894
2	9.449	9.893
3	14.800	14.291
4	19.264	19.227

**Table 1.**

Calibrator data of 3<sup>rd</sup> derivative peaks at 514 nm.