Technical support: support@abbkine.com

Website: https://www.abbkine.com

# CheKine™ Micro Superoxide Anion Assay Kit

Cat #: KTB1210 Size: 48 T/96 T

[ <u>=</u> Q	Micro Superoxide Anion Assay Kit		
REF	Cat #: KTB1210	LOT	Lot #: Refer to product label
	Applicable samples: Serum, Plasma, Animal and Plant Tissues, Cells, Cell Supernatant		
Å	Storage: Stored at 4°C for 6 months, protected from light		

## **Assay Principle**

Reactive oxygen species such as superoxide anions in organisms have immune and signal transduction effects, but when they accumulate too much, they will destroy cell membranes and biological macromolecules, leading to abnormal metabolism of cells and tissues in the body, causing many diseases. The superoxide anion in plant, animal tissues, serum and other samples reacts with hydroxylamine hydrochloride to produce  $NO_2^-$ , and  $NO_2^-$  reacts with Gris reagent. The mechanism of Gris analysis is summarized as the azo coupling between diazoniums, which is It is produced by sulfonamides and  $NO_2^-$  and N-(1-naphthyl) ethylenediamine dihydrochloride to generate a red azo compound with a characteristic absorption peak at 540 nm. The  $O_2^-$  content in the sample can be calculated based on the A540 value. The kit can detect samples of plants, animal tissues, serum and cells.

#### **Materials Supplied and Storage Conditions**

Kit sama ananta	s	ize	Storage conditions	
Kit components	48 T	96 T		
Extraction Buffer	50 mL	100 mL	4°C	
Reagent I	4 mL	8 mL	4°C	
Reagent II	3 mL	6 mL	4°C, protected from light	
ReagentIII	3 mL	6 mL	4°C, protected from light	
NaNO <sub>2</sub> Standard (10 mmoL/L)	0.5 mL	0.5 mL	4°C	

#### **Materials Required but Not Supplied**

- Microplate reader or visible spectrophotometer capable of measuring absorbance at 540 nm
- 96-well plate or microglass cuvette, precision pipettes, disposable pipette tips
- · Refrigerated centrifuge, ice maker, water bath
- · Trichloromethane, deionized water



Version 20230704

## **Reagent Preparation**

Extraction Buffer: Ready to use as supplied. Equilibrate to room temperature before use. Store at 4°C.

Reagent I: Ready to use as supplied. Equilibrate to room temperature before use. Store at 4°C.

Reagent II: Ready to use as supplied. Equilibrate to room temperature before use. Store at 4°C, protected from light.

Reagent III: Ready to use as supplied. Equilibrate to room temperature before use. Store at 4°C, protected from light.

Setting of standard curves: Dilute 20  $\mu$ L of NaNO<sub>2</sub> Standard (10 mmoL/L) to 200  $\mu$ mol/L with 980  $\mu$ L Extraction Buffer. And dilute the standard furtherly with Extraction Buffer as shown in the table below:

Num.	Volume of 200 μmol/L NaNO₂ Standard	Volume of Extraction Buffer	Concentration
Std.1	200 μL	0	200 μmoL/L
Std.2	100 μL	100 µL	100 μmoL/L
Std.3	50 μL	150 µL	50 μmoL/L
Std.4	20 μL	180 µL	20 μmoL/L
Std.5	10 μL	190 μL	10 μmoL/L
Std.6	5 μL	195 µL	5 μmoL/L
Std.7	2 μL	198 µL	2 μmoL/L
Std.8	1 μL	199 µL	1 μmoL/L

### **Sample Preparation**

- 1. Animal Tissues: Weigh 0.1 g tissues, add 1 mL Extraction Buffer and homogenize on ice. Centrifuge at 10,000 g for 10 min at 4°C. Use supernatant for assay, and place it on ice to be tested.
- 2. Plant Tissues: Weigh 0.1 g tissues, add 1 mL Extraction Buffer and mash. Ultrasonic break in ice bath 5 min (power 20% or 200 W, ultrasonic 3 s, interval 7 s, repeat 30 times). Centrifuge at 10,000 g for 10 min at 4°C. Use supernatant for assay, and place it on ice to be tested.
- 3. Cells: Collect 5×10<sup>6</sup> cells into the centrifuge tube, wash cells with cold PBS, discard the supernatant after centrifugation; add 1 mL Extraction Buffer to ultrasonically disrupt the cells 5 min (power 20% or 200 W, ultrasonic 3 s, interval 7 s, repeat 30 times). Centrifuge at 10,000 g for 10 min at 4°C. Use supernatant for assay, and place it on ice to be tested.
- 4. Serum, Plasma, Cell Supernatant or other liquid samples: Tested directly.

Note: It will be better to quantify the total protein with Protein Quantification Kit (BCA Assay), Cat #: KTD3001, if the content is calculated by protein concentration.

### **Assay Procedure**

- 1. Preheat the microplate reader or visible spectrophotometer for more than 30 min, and adjust the wavelength to 540 nm, visible spectrophotometer was returned to zero with deionized water.
- 2. Operation table:

Reagent	Control Tube (µL)	Blank Tube(µL)	Test Tube (µL)	Standard Tube (µL)
Different Concentration of Std.	0	0	0	40
Sample	40	0	40	0
Extraction Buffer	140	100	60	60



Version 20230704

Reagent I	0	80	80	80
Mix well, incubate in 37°C water bath for 20 min				
Reagent II	60	60	60	60
ReagentIII	60	60	60	60
Mix well, incubate in 37°C water bath for 20 min				
Trichloromethane	100	100	100	100

Mix well, Centrifuge at 8,000 g for 5 min at 25°C, Add 200  $\mu$ L into 96-well plate or microglass cuvette, measure the absorbance value at 540 nm and record it as A.  $\Delta$ A<sub>Test</sub>=A<sub>Test</sub>-A<sub>Control</sub>,  $\Delta$ A<sub>Standard</sub>=A<sub>Standard</sub>-A<sub>Blank</sub>.

Note: Each sample needs to set up a control tube to eliminate the influence of  $NO_2$  existing in the sample itself, so 96 T can only measure 48 samples. In order to guarantee the accuracy of experimental results, need to do a pre-experiment with 2-3 samples. If  $\Delta A_{Test}$  is less than 0.005, increase the sample quantity appropriately. If  $\Delta A_{Test}$  is greater than 1.0, the sample can be appropriately diluted with Extraction Buffer, the calculated result multiplied by the dilution factor, or decrease the sample quantity appropriately.

#### **Data Analysis**

Note: We provide you with calculation formulae, including the derivation process and final formula. The two are exactly equal. It is suggested that the concise calculation formula in bold is final formula.

1. Drawing of standard curve

With the concentration of the standard solution as the y-axis and the  $\Delta A_{Standard}$  as the x-axis, draw the standard curve y=kx+b. Bring the  $\Delta A_{Test}$  of the sample into the equation to get the y value ( $\mu$ mol/L).

- 2. Calculation of Superoxide Anion content
- (1) Calculated by protein concentration

Superoxide Anion content (µmol/mg prot)=y×V<sub>Sample</sub>÷(V<sub>Sample</sub>×Cpr)×10<sup>-3</sup>=y÷Cpr

Superoxide Anion production rate (µmol/ min/mg prot) =y×V<sub>Sample</sub>÷(V<sub>Sample</sub>×Cpr)÷T×10<sup>-3</sup>=0.05y÷Cpr

(2) Calculated by fresh weight of samples

Superoxide Anion content (µmol/g fresh weight)=y×V<sub>Sample</sub>÷(V<sub>Sample</sub>÷V<sub>Sample</sub> Total×W)×10<sup>-3</sup>=y÷W

Superoxide Anion production rate (µmol/min/g fresh weight)=y×V<sub>Sample</sub>÷(V<sub>Sample</sub>÷V<sub>Sample</sub>Total×W)÷T×10<sup>-3</sup>=0.05y÷W

(3) Calculated by volumet of liquid samples

Superoxide Anion content (µmol/mL)=y×10<sup>-3</sup>=y

Superoxide Anion production rate (µmol/min/mL)=y÷T×10<sup>-3</sup>=0.05y

Where:  $V_{Sample}$ : sample volume added, 0.04 mL; Cpr: sample protein concentration, mg/mL; T: reaction time, 20 min;  $V_{Sample Total}$ : extract buffer added to samples, 1 mL; W: sample weight, g;  $10^{-3}$ : 1 mL= $10^{-3}$  L.

# **Typical Data**

Typical standard curve:



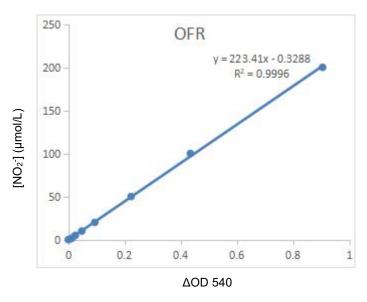


Figure 1. Standard curve for Superoxide Anion.

### **Recommended Products**

Catalog No.	Product Name
KTB1050	CheKine™ Micro Lipid Peroxidation (MDA) Assay Kit
KTB1041	CheKine™ Micro Hydrogen Peroxide (H <sub>2</sub> O <sub>2</sub> ) Assay Kit
KTB1310	CheKine™ Micro Glucose Oxidase Activity (GOD) Assay Kit
KTB1070	CheKine™ Micro Xanthine Oxidase (XO) Activity Assay Kit
KTB1200	CheKine™ Micro Protein Carbonyl Assay Kit

# **Disclaimer**

The reagent is only used in the field of scientific research, not suitable for clinical diagnosis or other purposes.

