

Reducing Sugar Content Assay Kit

Note: Take two or three different samples for prediction before test.

Operation Equipment: Spectrophotometer/microplate reader

Catalog Number: BC0235

Size: 100T/48S

Product Composition: Before use, please carefully check whether the volume of the reagent is consistent with the volume in the bottle. If you have any questions, please contact Solarbio staff in time.

Reagent name	Size	Preservation Condition
Reagent I	Liquid 50 mL×1	2-8°C
Reagent II	Liquid 10 mL×1	2-8°C
Standard	Powder×1	2-8°C

Solution Preparation:

1. **Standard:** Containing 10 mg of anhydrous glucose (loss on drying < 0.2%). Add 1 mL of distilled water to dissolve it for standby before use. It can be stored for two weeks at 2-8°C, or it can be stored for a longer time with saturated benzoic acid solution.

Product Description

Reducing sugar is widely found in animals, plants, microorganisms and cultured cells. Reducing sugars in plants mainly include glucose, fructose and maltose, which are the most common monosaccharides and disaccharides. Glucose and fructose are not only the main substrates of respiration, but also the substrates for further synthesis of sucrose, starch and cellulose.

Heating can promote the formation of brownish red amino compound from 3,5-Dinitrosalicylic acid solution and reducing sugar in alkaline solution. This brownish red amino compound has a characteristic absorption peak at 540 nm. Within a certain concentration range, the content of reducing sugar has a linear relationship with the absorbance at 540 nm. According to the standard curve, the amount of reducing sugar in the sample can be calculated.

Reagents and Equipment Required but Not Provided.

Spectrophotometer/microplate reader, water-bath, table centrifuge, sonicator transferpettor, micro glass cuvette/96 well flat-bottom plate, mortar and distilled water.

Procedure

I. Extraction of reducing sugar

a. Bacteria or cell treatment:

Collect the bacteria or cells into the centrifuge tube, discard the supernatant after centrifugation; The bacteria or cells (10^4) : the volume(mL) of Reagent I is 500~1000: 1 (It is suggest to add 1 mL of

Reagent I to 5 million of bacteria or cells), ultrasonic broke bacteria or cells (ice bath, power of 200 W, ultrasound for 3 s, interval of 10 s, repeat 30 times). Transfer to the covered centrifuge tube (to prevent water loss during heating), water bath at 80°C for 40 minutes and during which shake for 8-10 times. Centrifuge at 8000 ×g for 10 minutes at 25°C, take the supernatant for determination.

b. Tissue:

The proportion of tissue mass (g): the volume(mL) of Reagent I is 1:5-10 (it is recommended to weigh about 0.1 g of tissue and add 1 mL of Reagent I), ice bath homogenate. Transfer to a covered centrifuge tube (to prevent water loss during heating), water bath at 80°C for 40 minutes and during which shake for 8-10 times. Centrifuge at 8000 ×g for 10 minutes at 25°C, take the supernatant for determination.

c. Treatment of serum (slurry):

The proportion of serum (slurry) volume (mL): Reagent I volume (mL) is 1:5 ~ 10 (it is recommended to take 0.1 mL of serum (slurry) and add 0.9 mL of Reagent I), ice bath homogenate. Transfer to covered centrifuge tube (to prevent water loss during heating), water bath at 80°C for 40 minutes and during which shake for 8-10 times. Centrifuge at 8000 ×g for 10 minutes at 25°C, take the supernatant for determination.

II. Determination procedure:

a. Preheat the spectrophotometer/microplate reader for 30 minutes, adjust the wavelength to 540 nm and

the spectrophotometer adjust zero with distilled water.

b. Standard preparation: Dilute the standard with distilled water to 0.6、0.5、0.4、0.3、0.2、0.1 mg/mL.

c. Add the following reagents successively into the EP tube:

Reagent (μL)	Contrast Tube (C)	Test Tube (T)	Standard Tube (S)	Blank Tube (B)
Sample	175	175	-	-
Standard solution	-	-	175	-
Reagent II	-	125	125	125
Distilled water	125	-	-	175

Mix, heat in boiling water bath for 5 minutes (cover tightly to prevent water loss), cool to room temperature immediately after taking out, mix well. Take 200 μL of the reaction solution to a micro glass cuvette or 96 well plate, and read the absorbance values of standard tube, contrast tube, test tube and blank tube at 540 nm. Calculate $\Delta A = A_T - A_C$. Blank tubes and standard curves only need to be done 1-2 times.

III. Calculation of reducing sugar content:

1. Standard curve:

According to the concentration and absorbance of the standard tube ($A_S - A_B$), establish the standard curve, x is the absorbance value, y is the concentration of the standard (mg/mL). Calculate the content of reducing sugar in the sample according to the standard curve. Take $\Delta A (A_T - A_C)$ into x

to obtain y value by calculate.

2. Calculate by Sample fresh weight:

$$\text{Reducing sugar content } (\mu\text{g /g fresh weight}) = 1000 \times y \times V1 \div W = 1000 \times y \div W$$

3. Calculate by the protein concentration:

$$\text{Reducing sugar content } (\mu\text{g /mg prot}) = (1000 \times y \times V1) \div (V1 \times Cpr) = 1000 \times y \div Cpr$$

4. Calculate by the number of bacteria or cells

$$\text{Reducing sugar content } (\mu\text{g /}10^4 \text{ cell}) = 1000 \times y \times V1 \div 500 = 2 \times y$$

5. Calculate by the volume of serum (plasma):

$$\text{Reducing sugar content } (\mu\text{g /mL}) = 1000 \times y \times V2 \div V3 = 10000 \times y$$

1000: Unit conversion coefficient, 1 mg/mL = 1000 μ g/mL;

V1: Add the volume of Reagent I, 1 mL;

V2: Add the total volume of serum and Reagent I, 1 mL;

V3: Add the volume of serum (plasma), 0.1 mL;

Cpr: Sample protein concentration, mg/mL;

W: Sample fresh weight, g;

500: Total number of bacteria or cells, 5 million.

Notes:

1. Each test tube shall be provided with a contrast tube.
2. If the absorbance value exceeds the linear range, the sample size can be increased or diluted before determination

Recent Product Citations:

- [1] Kang L, Wu Y, Jia Y, Chen Z, Kang D, Zhang L, Pan C. Nano-selenium enhances melon resistance to *Podospaera xanthii* by enhancing the antioxidant capacity and promoting alterations in the polyamine, phenylpropanoid and hormone signaling pathways. *J Nanobiotechnology*. 2023 Oct 16;21(1):377. doi: 10.1186/s12951-023-02148-y. PMID: 37845678; PMCID: PMC10577987.
- [2] Xu Y, Zhang Y, Zhu J, Sun Y, Guo B, Liu F, Huang J, Wang H, Dong S, Wang Y, Wang Y. *Phytophthora sojae* apoplastic effector AEP1 mediates sugar uptake by mutarotation of extracellular aldose and is recognized as a MAMP. *Plant Physiol*. 2021 Sep 4;187(1):321-335. doi: 10.1093/plphys/kiab239. PMID: 34618132; PMCID: PMC8418418.
- [3] Chen J, Lan M, Zhang X, Jiao W, Chen Z, Li L, Li B. Effects of Simulated In Vitro Digestion on the Structural Characteristics, Inhibitory Activity on α -Glucosidase, and Fermentation Behaviours of a Polysaccharide from *Anemarrhena asphodeloides* Bunge. *Nutrients*. 2023 Apr 19;15(8):1965. doi: 10.3390/nu15081965. PMID: 37111183; PMCID: PMC10145594.

- [4] Ye Z, Xu YJ, Liu Y. Influence of different dietary oil consumption on nutrient malabsorption: An animal trial using Sprague Dawley rats. *J Food Biochem.* 2021 Apr;45(4):e13695. doi: 10.1111/jfbc.13695. Epub 2021 Mar 11. PMID: 33694208.

References:

[1] Lindsay H. A colorimetric estimation of reducing sugars in potatoes with 3, 5-dinitrosalicylic acid[J]. *Potato Research*, 1973, 16(3): 176-179.

[2] Breuil C, Saddler J N. Comparison of the 3, 5-dinitrosalicylic acid and Nelson-Somogyi methods of assaying for reducing sugars and determining cellulase activity[J]. *Enzyme and microbial technology*, 1985, 7(7): 327-332.

[3] Brunton N P, Gormley T R, Murray B. Use of the alditol acetate derivatisation for the analysis of reducing sugars in potato tubers[J]. *Food chemistry*, 2007, 104(1): 398-402.

Related Products:

- BC0330/BC0335 Trehalose Content Assay Kit
BC0340/BC0345 Glucogen Content Assay Kit
BC2530/BC2535 Sorbitol Dehydrogenase(SDH) Activity Assay Kit
BC0030/BC0035 Plant Soluble Sugar Content Assay Kit

Technical Specifications:

Minimum Detection Limit: 0.0616 mg/mL

Linear Range: 0.1-0.6 mg/mL