

Bradford Protein Assay Kit

Product #: Bi2M-BPA-250

Quantity: 250 mL

Product Description

- The Bradford assay is widely used for determining protein concentration.
 - Based on the binding of Coomassie Brilliant Blue G250 to proteins.
 - Free dye has maximum absorption at 488 nm.
 - When bound to protein, maximum absorption shifts to 595 nm.
 - Absorbance at 595 nm is proportional to protein concentration.
 - Effective for most samples, tolerating up to 1 mM β -mercaptoethanol and 5 mM dithiothreitol.
 - Sensitive to descaling agents: ensure SDS < 0.01%, Triton X-100 < 0.05%, and Tween-20/60/80 < 0.015%.
 - Use BCA Protein Quantitative Detection Kit (G2026) for samples with descaling agents.
- **Advantages of Bradford Method:**
 - Quick and easy to perform.
 - High sensitivity.
 - Fast detection, suitable for protein concentrations of 25-1000 μ g/mL.

Storage

- Store BSA standard ampules at 4°C.
- Store protein standard solution (prepared with BSA standard ampules and standard ampules dilution) at -20°C.

Components Provided

Coomassie G-250 dye	250 mL
Bovine Serum Albumin (BSA) Standard Ampules	25 mg
Standard Ampules Dilution	1.5 mL

Important Notes (PLEASE READ CAREFULLY)**1. Coomassie G-250 Dye:**

- Bring to room temperature before use.
- Mix by inverting to maintain assay sensitivity.

2. Protein Standard Preparation:

- Ensure the protein standard stock solution is fully dissolved.
- Dilute the working solution in 10-fold gradients.
- Avoid 50-fold dilutions to minimize errors.

3. Safety Precautions:

- Wear safety glasses.
- Use gloves.
- Wear protective clothing.

Protocol**1. Preparation of Protein Standard Stock Solution:**

1. Dissolve 25 mg Bovine Serum Albumin (BSA) in 1 mL Standard Ampules Dilution.
2. Final concentration: 25 mg/mL.
3. Store at -20°C.

2. Preparation of Protein Standard Working Solution:

1. Dilute the 25 mg/mL Protein Standard Stock Solution 50 times with PBS or normal saline.
2. Final concentration: 0.5 mg/mL.
3. Use a 10-fold gradient dilution method for accuracy.

3. Preparation of Standard Curve (Microplate Reader Method):

1. Add protein standard working solution to 96-well plate: 0, 1, 2, 4, 8, 12, 16, 20 μ L.
2. Add PBS or normal saline: 20, 19, 18, 16, 12, 8, 4, 0 μ L to each well.

3. Final concentrations: 0, 25, 50, 100, 200, 300, 400, 500 $\mu\text{g/mL}$.

4. Preparation of Samples:

1. Dilute samples appropriately to fall within the standard curve range.
2. Add 20 μL of each sample to a 96-well plate.
3. Use the same diluent as the protein standard.

5. Microplate Procedure:

1. Add 200 μL of Coomassie G-250 dye to each well.
2. Shake plate for 30 seconds.
3. Incubate at room temperature for 3-5 minutes.
4. Measure absorbance at 595 nm using a microplate reader.
5. Plot the standard curve and calculate sample protein concentration from the absorbance values.

6. Spectrophotometer Procedure:

1. Adjust the gradient standard reaction volume to 1 mL.
2. Add 0, 0.05, 0.1, 0.2, 0.4, 0.6, 0.8, 1.0 mL of 0.5 mg/mL protein standard working solution to glass test tubes.
3. Add 1.0, 0.95, 0.9, 0.8, 0.6, 0.4, 0.2, 0 mL of PBS or saline to make up to 1 mL.
4. Mix thoroughly to obtain gradient concentrations of 0, 25, 50, 100, 200, 300, 400, 500 $\mu\text{g/mL}$.

7. Sample Preparation for Spectrophotometer:

1. Dilute protein samples as needed.
2. Add 1 mL of diluted sample to glass test tubes.

8. Colorimetric Detection (Spectrophotometer):

1. Add 3 mL of Coomassie G-250 dye to each standard curve and sample tube.

2. Mix thoroughly and let stand for 3-5 minutes.
3. Measure absorbance at 595 nm.
4. Use standard curve to determine protein concentration in samples.

9. Calculation:

1. Plot gradient protein content ($\mu\text{g/mL}$) against absorbance.
2. Determine sample protein concentration from the standard curve.
3. Multiply by the dilution factor for actual protein concentration.

NOTES:

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