Bradford(Coomassie) Protein Assay Plus Kit

P7200

Storage

Bradford (Coomassie) Protein Assay Plus Reagent : Store at 4°C and protected from light.

Albumin Standard Stored at RT.

Contents

- Product manual
- Bradford (Coomassie) Protein Assay Plus Reagant 450ml
- Albumin Standard, 2mg/ml, 5x1ml
  contains bovine serum albumin (BSA) at 2.0 mg/ml in 0.9% slaine and 0.05% sodium azide

Shipping Conditon

Ship at ambient - Stable for 3 weeks at room temperature

Introduction

The Coomassie (Bradford) Protein Assay Kit is a quick and ready-to-use modification of the wellknown Bradford coomassie-binding, colorimetric method for total protein quantitation. When coomassie dye binds protein in an acidic medium, an immediate shift in absorption maximum occurs from 465nm to 595nm with a concomitant color change from brown to blue. Performing the assay in either test tube or microplate format is simple: combine a small amount of protein sample with the assay reagent, mix well, incubate briefly and measure the absorbance at 595nm. Protein concentrations are estimated by reference to absorbances obtained for a series of standard protein dilutions, which are assayed alongside the unknown samples. Because the color response with coomassie is non-linear with increasing protein concentration, a standard curve must be completed with each assay.

Procedure

General considerations

There are four protocols for the use of Bradford (Coomassie) Protein Assay Plus Reagent. Depending on the protein concentration, standard or micro assay can be used. Depending on the volume of protein sample, the assay can be performed in test tubes or microplates.

Preparation of Standards and Assay Reagent

A. Preparation of Diluted Albumin (BSA) Standards

Use Table 1 as a guide to prepare a set of protein standards. Dilute the contents of one Albumin Standard (BSA) ampule into several clean vials, preferably in the same diluent as the sample(s). Each 1mL ampule of Albumin Standard is sufficient to prepare a set of diluted standards for either working range suggested in Table 1. There will be sufficient volume for three replications of each diluted standard.

Table 1. **Preparation of Diluted Albumin** (BSA) Standards
Dilution Scheme for Standard Test Tube and Microplate Protocols
(Working Range = 100–1500µg/mL)

Vial	Vol. of Diluent	Vol. of Source of BSA	Final BSA Concentration
A	0	300uL of Stock	2000ug/mL
B	125uL	375uL of Stock	1500ug/mL
C	325uL	325uL of Stock	1000ug/mL
D	175uL	175uL of Vial B dilution	750ug/mL
E	325uL	325uL of Vial B dilution	500ug/mL
F	325uL	325uL of Vial B dilution	250ug/mL
G	325uL	325uL of Vial D dilution	125ug/mL
H	400uL	100uL of Vial E dilution	25ug/mL
I	400uL	0	0 ug/mL = Blank

Dilution Scheme for Micro Test Tube and Microplate Protocols
(Working Range = 1–25µg/mL)

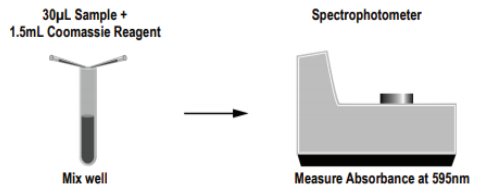
Vial	Vol. of Diluent	Vol. of Source of BSA	Final BSA Concentration
A	2370uL	30uL of Stock	25ug/mL
B	4950uL	50uL of Stock	20ug/mL
C	3970uL	30 uL of Stock	15ug/mL
D	2500uL	2500uL of Vial B dilution	10ug/mL
E	2000uL	2000uL of Vial B dilution	5ug/mL
F	1500uL	1500uL of Vial B dilution	2.5ug/mL
G	5000uL	0	0ug/mL=Blank

B. Equilibrating and Mixing of the Coomassie Reagent

Mix the Coomassie Reagent solution immediately before use by gently inverting the bottle several times (Do not shake the bottle to mix the solution). Remove the amount of reagent needed and equilibrate it to room temperature (RT) before use.

Note: Dye-dye and dye-protein aggregates tend to form in all coomassie-based reagents. If left undisturbed, the aggregates will become large enough over time to be visible. For example, when left overnight in a clear glass tube, the reagent forms dye-dye aggregates that are visible as a dark precipitate in the bottom of the tube with nearly colorless liquid above. Dye-dye aggregates can form over several hours in stored reagent while dye-protein-dye aggregates form more quickly. Fortunately, gentle mixing completely disperses the dye-dye aggregates. Therefore, it is good practice to mix the Coomassie Reagent before pipetting and to mix each tube or plate immediately before measuring absorbances.

Procedure Summary
(Standard Test Tube Protocol):



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Test Tube Procedure

A. Standard Test Tube Protocol (Working Range = 100-1500µg/mL)

1. Pipette 0.03mL (30µL) of each standard or unknown sample into appropriately labeled test tubes.
2. Add 1.5mL of the Coomassie Reagent to each tube and mix well.
3. Optional: For the most consistent results, incubate samples for 10 minutes at room temperature (RT).
4. With the spectrophotometer set to 595nm, zero the instrument on a cuvette filled only with water. Subsequently, measure the absorbance of all the samples.
5. Subtract the average 595nm measurement for the Blank replicates from the 595nm measurements of all other individual standard and unknown sample replicates.
6. Prepare a standard curve by plotting the average Blank-corrected 595nm measurement for each BSA standard vs. its concentration in µg/mL. Use the standard curve to determine the protein concentration of each unknown sample.

B. Micro Test Tube Protocol (Working Range = 1-25µg/mL)

1. Pipette 1.0mL of each standard or unknown sample into appropriately labeled test tubes.
2. Add 1.0mL of the Coomassie Reagent to each tube and mix well.
3. Optional: For the most consistent results, incubate samples for 10 minutes at room temperature (RT).
4. With the spectrophotometer set to 595nm, zero the instrument on a cuvette filled only with water. Subsequently, measurethe absorbance of all the samples.
5. Subtract the average 595nm measurement for the Blank replicates from the 595nm measurements of all other individual standard and unknown sample replicates.
6. Prepare a standard curve by plotting the average Blank-corrected 595nm measurement for each BSA standard vs. its concentration in µg/mL. Use the standard curve to determine the protein concentration of each unknown sample.

Microplate Procedure

A. Standard Microplate Protocol (Working Range = 100-1500µg/mL)

1. Pipette 5µL of each standard or unknown sample into the appropriate micro -plate wells
2. Add 250µL of the Coomassie Reagent to each well and mix with plate shaker for 30 seconds.
3. Remove plate from shaker. For the most consistent results, incubate plate for 10 minutes at room temperature (RT).
4. Measure the absorbance at or near 595nm with a plate reader.
5. Subtract the average 595nm measurement for the Blank replicates from the 595nm measurements of all other individual standard and unknown sample replicates.
6. Prepare a standard curve by plotting the average Blank-corrected 595nm measurement for each BSA standard vs. its concentration in µg/mL. Use the standard curve to determine the protein concentration of each unknown sample.

Note: When compared to the Standard Test Tube Protocol, 595nm measurements obtained with the Microplate Protocols are lower because the light path used is shorter. Consequently, this may increase the minimum detection level of the assay. If higher 595nm measurements are required, use 7-10µL of standard or sample and 250µL of Coomassie Reagentper well.

ote: If using curve-fitting algorithms associated with a microplate reader, a four-parameter (quadratic) or best-fit curve will provide more accurate results than a purely linear fit. If plotting results by hand, a point-to-point curve is preferable to a linear fit to the standard points.

B. Micro Microplate Protocol (Working Range = 1-25µg/mL)

1. Pipette 150µL of each standard or unknown sample into the appropriate mic Droplate wells.
2. Add 150µL of the Coomassie Reagent to each well and mix with plate shaker for 30 seconds.
3. Remove plate from shaker. For the most consistent results, incubate plate for 10 minutes at room temperature (RT).
4. Measure the absorbance at or near 595nm on a plate reader.
5. Subtract the average 595nm measurement for the Blank replicates from the 595nm measurements of all other individual standard and unknown sample replicates.
6. Prepare a standard curve by plotting the average blank corrected 595nm measurement for each BSA standard vs. its concentration in µg/mL. Using the standard curve, determine the protein concentration estimate for each unknown sample.

Note: If using curve-fitting algorithms associated with a microplate reader, a four-parameter (quadratic) or best-fit curve will provide more accurate results than a purely linear fit. If plotting results by hand, a point-to-point curve is preferable to a linear fit to the standard points.

Troubleshooting

Problem	Possible Cause	Solution
Absorbance of Blank is OK, but remaining standards and samples yield lower values than expected	Improper reagent storage	Store reagent refrigerated
	Reagent still cold	Allow Reagent to warm to RT
	Absorbance measured at incorrect wavelength	Measure absorbance near 595nm
Absorbances of Blank and standards are OK, but samples yield lower values than expected	Sample protein (peptide) has a low molecular weight (e.g., less than 3000)	Use the BCA or Lowry Protein Assay
A precipitate forms in all tubes	Sample contains a surfactant (detergent)	Dialyze or dilute sample Remove interfering substances from sample using Protein Assay Preparation Reagent Set
	Samples not mixed well or left to stand for extended time, allowing aggregates to form with the dye	Mix samples immediately prior to measuring absorbances
All tubes (including Blanks) are dark blue	Strong alkaline buffer raises pH of formulation, or sample volume too large, thereby raising reagent pH	Dialyze or dilute sample Remove interfering substances from sample .
Need to read absorbances at a different wavelength	Spectrophotometer or plate reader does not have 595nm filter	Color may be read at any wavelength between 575nm and 615nm, although the slope of standard curve and overall assay sensitivity will be reduced

A. Interfering substances

Certain substances are known to interfere with coomassie-based protein assays including most ionic and nonionic detergents, which reduce color development and can cause precipitation of the assay reagent. Other substances interfere to a lesser extent. These have only minor (tolerable) effects below a certain con -centration in the original sample. Maximum compatible concentrations for many substances in the Standard Test Tube Protocol are listed in Table 2 (see last page). Substances were compatible in the Standard Test Tube Protocol if the error in protein concentration estimation (of BSA at 1000µg/mL) caused by the presence of the substance in the sample was less than or equal to 10%. The Blank-corrected 595nm absorbance measurements (for the 1000µg/mL BSA standard + substance) were compared to the net 595nm absorbances of the 1000µg/mL BSA standard prepared in 0.9% saline

B. Strategies for eliminating or minimizing the effects of interfering substances

The effects of interfering substances in the Coomassie Assay may be overcome by several methods.

- Remove the interfering substance by dialysis or desalting.
- Dilute the sample until the substance no longer interferes.
- Precipitate proteins with acetone or trichloroacetic acid (TCA).

Upon precipitation the liquid containing the substance that interfered is discarded and the protein pellet is solubilized in a small amount of ultrapure water or directly in the Coomassie Reagent.

Note: For greatest accuracy, the protein standards must be treated identically to the sample(s).

Additional Information

A. Response characteristics for different proteins

Each of the commonly used total protein assay methods exhibits some degree of varying response toward different proteins. These differences relate to amino acid sequence, isoelectric point, structure and the presence of certain side chains or prosthetic groups that can dramatically alter the protein's color response. Most protein assay methods utilize BSA or immunoglobulin (IgG) as the standard against which the concentration of protein in the sample is determined. Albumin dard curve should be prepared from a pure sample
of the target protein to be measured.

Table 3 shows typical protein-to-protein variation in color response. All proteins were tested at a concentration of 1000µg/mL using the Standard Test Tube Protocol. The average net color response for BSA was normalized to 1.00 and the average net color response of the other proteins is expressed as a ratio to the response of BSA. The protein-to-protein variation observed with the Coomassie Reagent is significantly less than that seen with other Bradford-type coomassie dye formulations.

B. Measuring Absorbances at Wavelengths other than 595nm

If a photometer or plate reader is not available with a 595nm filter, the blue color may be measured at any wavelength between 570nm and 610nm. The maximum sensitivity of the assay occurs when the absorbance of the dye-protein complex is measured at 595nm. Measuring the absorbance at any wavelength other than 595nm will result in a lower slope for the standard curve and may increase the minimum detection level for the protocol.

C. Effect of Temperature on 595nm Absorbance

Absorbance measurements at 595nm obtained with the Coomassie Reagent are dependent on the temperature of the reagent to some extent. As the reagent temperature increases to room temperature, the 595nm measurements will increase. Therefore, it is important that the Coomassie Reagent remain at a constant temperature (i.e., RT) during the assay.

D Cleaning and Re-using Glassware

Care must be exercised when cleaning glassware that will be used again for protein assays. Thorough cleaning often requires the use of a detergent which must be completely removed in the final rinse. The coomassie dye will stain glass or quartz cuvettes. Disposable polystyrene cuvettes are a convenient a lternative.

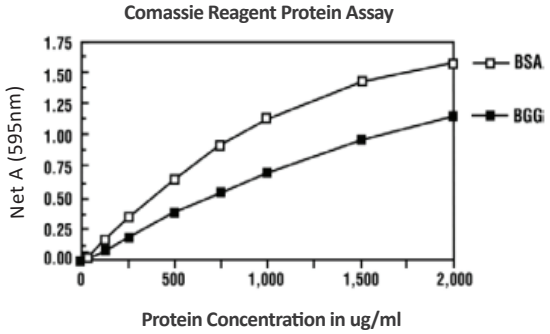

Figure 1. Typical color response curves for BSA and BGG using the Standard Test Tube Protocol of the Coomassie Assay.

Table 3. Protein-to-Protein Variation. Absorbance ratios (595nm) for proteins relative to BSA using the Standard Test Tube Protocol in the Coomassie Assay.

*** Ratio = (Avg "test" net Abs.) / (avg. BSA net Abs.)**

Table 4. Compatible substance concentrations in the Coomassie Assay (see text for details).

Substance (Salts/Buffer)	Compatible Concentration
ACES, pH 7.8	100mM
Ammonium sulfate	1M
Asparagine	10mM
Bicine, pH 8.4	100mM
Bis-Tris, pH 6.5	100mM
Borate (50mM), pH 8.5	undiluted
B-PER™ Reagent	1/2 dilution*
Calcium chloride in TBS, pH 7.2	10mM
Na-Carbonate/Na-Bicarbonate (0.2M), pH 9.4	undiluted
Cesium bicarbonate	100mM
CHES, pH 9.0	100mM
Na-Citrate (0.6M), Na-Carbonate (0.1M),pH 9.0	undiluted
Na-Citrate (0.6M), MOPS (0.1M), pH 7.5	undiluted
Cobalt chloride in TBS, pH 7.2	10mM
EPPS, pH 8.0	100mM
Ferric chloride in TBS, pH 7.2	10mM
Glycine	100mM
Guanidine•HCl	3.5M
HEPES, pH 7.5	100mM
Imidazole, pH 7.0	200mM
MES, pH 6.1	100mM
MES (0.1M), NaCl (0.9%), pH 4.7	undiluted
MOPS, pH 7.2	100mM
Modified Dulbecco's PBS, pH 7.4	undiluted
Nickel chloride in TBS, pH 7.2	10mM
PBS; Phosphate (0.1M), NaCl (0.15M), pH 7.2	undiluted
PIPES, pH 6.8	100mM
RIPA lysis buffer; 50mM Tris, 150mM NaCl, 0.5% DOC, 1% NP-40, 0.1% SDS, pH 8.0	1/10 diution*
Sodium acetate, pH 4.8	180mM
Sodium azide	0.5%
Sodium bicarbonate	100mM
Sodium chloride	5.0M
Sodium citrate, pH 4.8 or pH 6.4	200mM
Sodium phosphate	100mM
Tricine, pH 8.0	100mM
Triethanolamine, pH 7.8	100mM
Tris	2M
TBS; Tris (25mM), NaCl (0.15M), pH 7.6	undiluted
Tris (25mM), Glycine (192mM), pH 8.0	undiluted
Tris (25mM), Glycine (192mM), SDS (0.1%), pH 8.3	1/2 dilution*
Zinc chloride in TBS, pH 7.2	10mM

Substance Detergents	Compatible Concentration
Brij™-35	0.125%
Brij-56, Brij-58	0.031%
CHAPS, CHAPSO	5.0%
Deoxycholic acid	0.05%
Lubrol™ PX	0.125%
Octyl β-glucoside	0.5%
Nonidet P-40 (NP-40)	0.5%
Octyl β-thioglucopyranoside	3.0%
SDS	0.125%
Span™ 20	0.5%
Triton™ X-100, X-114	0.125%
Triton X-305, X-405	0.5%
Tween™-20	0.062%
Tween-60	0.1%
Tween-80	0.062%
Zwittergent™ 3-14	0.025%
Chelating agents	
EDTA	100mM
EGTA	2mM
Sodium citrate	200mM
Reducing & Thiol-Containing Agents	
N-acetylglucosamine in PBS, pH 7.2	100mM
Ascorbic acid	50mM
Cysteine	10mM
Dithioerythritol (DTE)	1mM
Dithiothreitol (DTT)	5mM
Glucose	1M
Melibiose	100mM
2-Mercaptoethanol	1M
Potassium thiocyanate	3M
Thimerosal 0.01%	0.01%
Misc. Reagents & Solvents	
Acetone	10%
Acetonitrile	10%
Aprotinin	10mg/L
DMF, DMSO	10%
Ethanol	10%
Glycerol (Fresh)	10%
Hydrochloric Acid	100mM
Leupeptin	10mg/mL
Methanol	10%
Phenol Red	0.5mg/mL
PMSF	1mM
Sodium Hydroxide	100mM
Sucrose	10%
TLCK	0.1mg/L
TPCK	0.1mg/mL
Urea	3M
o-Vanadate (sodium salt), in PBS, pH 7.2	1mM

*Diluted with ultrapure water.

Related Products

Product Name	Cat No
Albumin, Ultra pure Bovine Serum Albumin	A0100
Xpert Protease Inhibitor Cocktail Solution (100X)	P3100
Xpert Prestained protein marker	P8502
Tween 20, Molecular Biology Grade	P9100
West-Ez Blocking Buffer, 3% BSA	W3710
West-Q Chemiluminescent Substrate Kit	W3650
West-Q Chemiluminescent Substrate kit, Plus	W3651
West-Ez Stripping Buffer	S2100

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