

Thermo Scientific Pierce Protein Assay Technical Handbook



Total Protein Assays

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Quick Technical Summaries – Thermo Scientific Protein Assays

| Working Range (sample volume)* | Characteristics/Advantages | Applications | Disadvantages | Interfering Substances |
|--|---|--|---|---|
| Pierce [®] 660 nm Protein | Assay | | | |
| Standard Protocol: 25-2,000 μg/ml (65 μl) | Compatible with reducing agents, chelating agents and detergents | Ideal for measuring total protein concentration in | Use reagent with IDCR (Ionic Detergent | High levels of ionic detergents require the |
| Microplate Protocol: 50-2,000 μg/ml (10 μl) | Faster and easier to perform than BCA or Coomassie (Bradford) Assays | both reducing agents | Compatibility Reagent) with samples containing ionic detergents like SDS | addition of the Ionic Detergent Compatibility Reagent (IDCR). |
| | Excellent linearity of color development within the detection range | Used for quick, yet accurate estimation of protein | Greater protein-to-protein variability than the | - |
| | Less protein-to-protein variability than the Coomassie (Bradford) Assay | | BCA Assay | |
| | Reaches a stable end point | | | |
| | Compatible with Laemmli sample buffer containing bromophenol blue when using Compatibility Buffer | | | |
| The BCA Protein Assay | - Reducing Agent Compatible | | | |
| Standard Protocol: 125-2,000 μg/ml (25 μl) | Compatible with up to 5 mM DTT, 35 mM 2-mercaptoethanol or 10 mM TCEP | Allows the use of the superior BCA Assay in | Requires heating for color development | Compatible with all reducing agents and detergents found |
| Microplate Protocol: 125-2,000 μg/ml (9 μl) | No protein precipitation involved | situations in which it is normally unable to be read | | at concentrations routinely used in protein sample buffers |
| 125-2,000 µg/111 (9 µI) | Sample volume only 9 µl (microplate protocol) | No precipitation step | | |
| | Compatible with most detergents | means no worries about difficult-to-solubilize proteins | | |
| | Significantly less (14-23%) protein:protein variation than Bradford-based methods | | | |
| The BCA Protein Assay | | | | |
| Standard Protocol: 20-2,000 μg/ml (50 μl) | Two stable reagents used to make one working reagent | Adaptable for use with microplates | Not compatible with thiols/reducing agents | Reducing sugars and reducing agents |
| Enhanced Standard Protocol: 5-250 μg/ml | Working reagent stable for one week at room temperature | | Requires heating for color development Not a true end-point assay | Thiols Copper chelating agents |
| (50 µl) Microplate Protocol: | Compatible with detergents | Measure the amount of | | Ascorbic acid and uric acid |
| 20-2,000 μg/ml (25 μl) | Simple, easy to perform | protein covalently bound to affinity supports | | Tyrosine, cysteine and |
| | Less protein:protein variation than Coomassie dye methods | Determine copper levels using a reagent formulated | | tryptophan 50 mM Imidazole, 0.1 M Tris, |
| | Works with peptides (three amino acids or larger) | with BCA Reagent A⁴ | | 1.0 M glycine |
| | Flexible incubation protocols allow customization of reagent sensitivity and working range | | | |
| The Micro BCA Protein | Assay | | | |
| Standard Protocol: 60°C for 60 minutes | Three stable reagents used to make one working reagent | Suitable for determining protein concentration | More substances interfere at lower concentrations than with BCA Assay because the sample | Reducing sugars and reducing agents |
| 0.5-20 μg/ml (0.5 ml) Microplate Protocol: 37°C for 120 minutes 2-40 μg/ml (150 μl) | Working reagent stable for 24 hours at | in very dilute aqueous solutions | | Thiols |
| | room temperature | Adaptable for use with | volume-to-reagent volume | Copper chelating agents |
| | Compatible with most detergents | microplates ¹ | ration is 1:1 | Ascorbic acid and uric acid |
| | Simple, easy to perform | | 60°C water bath is needed | Tyrosine, cysteine and |
| | Less protein:protein variation than BCA, Coomassie dye or Lowry Methods | | | tryptophan 50 mM Imidazole, 0.1 M Tris, |
| | Works with peptides (three amino acids or larger) | | | 1.0 M glycine |
| | Linear color response to increasing protein concentration | | | |

* Sample volume per 1 ml total assay volume for measurement in 1 cm cuvette (Standard Protocol). Sample volume per 200-300 µl total volume for measurement in 96-well microplate.

Quick Technical Summaries – Thermo Scientific Protein Assays

| Working Range (sample volume)* | Characteristics/Advantages | Applications | Disadvantages | Interfering Substances |
|---|---|---|---|---|
| The Modified Lowry Pro | tein Assay | | | |
| Standard Protocol: 1-1,500 μg/ml | Two-reagent system – shelf life of at least one year | Lowry method is the most cited protein assay in the | Timed addition of Folin reagent adds complexity | Detergents (cause precipitation) |
| Microplate Protocol: | Two-step incubation requires precise | literature | Longer total assay time | Thiols, disulfides |
| 10-1,500 μg/ml (40 μl) | sequential timing of samples | Adaptable for use with microplates | Practical limit of about | Copper chelating reagents |
| | Color response read at 750 nm | morophatoo | 20 samples per run | Carbohydrates including |
| | Works with peptides (three amino acids or larger) | | | hexoseamines and their <i>N</i> -actyl derivatives |
| | Protein:protein variation similar to that seen with BCA Method | | | Glycerol, Tris, Tricine, K ¹⁺ ion |
| | Many authors have reported ways to deal with substances that interfere | | | |
| Coomassie Plus (Bradfor | rd) Assay | | | |
| Linear Range: | Simple/fast protocols | Standard assay ⁸ | Less linear color response | Detergents ¹⁸ |
| lgG: <i>125-1,500 μg/ml</i> BSA; <i>125-1,000 μg/ml</i> | Total preparation and assay time < 30 minutes | Micro assay ^{9,10,11} | in the micro assay | |
| , | One reagent system; stable for 12 months | Microplate format assay ¹² | Effect of interfering | |
| Standard Protocol: Sample-to-Reagent Ratio: 1:30 | Read-to-use formulation — no dilution or filtration needed | Assay of protein solutions | substances more pronounced in the micro assay | |
| Typical Working Range: 100-1,500 μg/ml (35 μl) | Nearly immediate color development at room temperature | Quantitation of immobilized protein ¹⁴ | Protein dye complex has tendency to adhere to | |
| Microplate Protocol: Sample-to-Reagent | Linear color response in standard assay (more accurate results) | Protein in permeabilized cells ¹⁵ | glass (easily removed with MeOH) ¹⁷ | |
| Ratio: 1:1 Typical Working Range: | Color response sensitive to changes in pH | | Protein must be > 3,000 Da | |
| <i>1-25 μg/ml (150 μl)</i> | Temperature dependence of color response | $NaCNBH_3$ determination ¹⁶ | | |
| | Compatible with buffer salts, metal ions, reducing agents, chelating agents | | | |
| | Low-odor formulation | | | |
| Coomassie (Bradford) Pr | rotein Assav | | | |
| Standard Protocol: | Simple-to-perform protocols | Standard assay ⁸ | Nonlinear color response | Detergents ¹⁸ |
| Sample-to-Reagent | One-reagent system, stable for 12 months | Micro assay ^{9,10,11} | More protein standard | 5 |
| Ratio: 1:50 100-1,500 μg/ml (20 μl) | Ready-to-use formulation | , Microplate format assay ¹⁹ | concentrations required to | |
| Microplate Protocol: | No dilution or filtration needed | Assay of protein solutions | cover working range | |
| Sample-to-Reagent Ratio: <i>1:1</i> 1-25 µg/ml (150 µl) | Fast, nearly immediate color development at room temperature | containing reducing agents Cell line lysates ²⁰ | Micro assay has potential for interference | |
| | Total preparation and assay time < 30 minutes | Protein recovery studies | Protein must be > 3,000 Da | |
| | Typical protein:protein variation expected for a Coomassie dye-based reagent | , | | |
| | Color response sensitive to pH | | | |
| | Temperature-dependent color response | | | |
| | Compatible with buffer salts, metal ions, reducing agents, chelating agents | | | |

* Sample volume per 1 ml total assay volume for measurement in 1 cm cuvette (Standard Protocol). Sample volume per 200-300 µl total volume for measurement in 96-well microplate.

Quick Technical Summaries – References

| Working Range | Characteristics/Advantages | Benefits |
|---|--|--|
| Pre-Diluted Protein | Assay Standard Sets | |
| Working Range : 125-2,000 μg/ml | Ready to use | No dilution series preparation |
| | 3.5 ml each of seven standard curve data | Dramatically improved speed to result |
| | points within the working range | General utility standards for BCA, Bradford and Lowry Assay methods |
| | Stable and sterile filtered | More reliable quantitation |
| | 15-35 standard test tube assays or 175-350 microplate assays | Standard set is treated as you would treat the sample |
| | | Unparalleled convenience |
| | | Economical for microplate format assays |
| protein assay for u <i>Biochem.</i> 153 , 267 | and Turley, R.B. (1986). Adaptation of the bicinchoninic acid use with microtiter plates and sucrose gradient fractions. <i>Anal</i> -271. Rrodback II (1986). A sensitive protein assay using microtiter | Simpson, I.A. and Sonne, O. (1982). A simple, rapid and sensitive method for measuring protein concentration in subcellular membrane fractions prepared by sucrose density ultracentrifugation. <i>Anal. Biochem.</i> 119, 424-427. Redinbaugh, M.G. and Campbell, W.H. (1985). Adaptation of the dye-binding protein assay to microtitor plates. <i>Anal. Biochem.</i> 117, 141-147. |

- Sorensen, K. and Brodbeck, U. (1986). A sensitive protein assay using microtiter plates. J. Immunol. Meth. 95, 291-293.
- 3. Stich, T.M. (1990). Determination of protein covalently bound to agarose supports using bicinchoninic acid. Anal. Biochem. 191, 343-346.
- 4. Brenner, A.J. and Harris, E.D. (1995). A quantitative test for copper using bicinchoninic acid. Anal. Biochem. 226, 80-84.
- 5. Akins, R.E. and Tuan, R.S. (1992). Measurement of protein in 20 seconds using a microwave BCA assay. Biotechniques. 12(4), 496-499.
- 6. Brown, R.E., et al. (1989). Protein measurement using bicinchoninic acid: elimination of interfering substances. Anal. Biochem. 180, 136-139.
- 7. Peterson, G.L. (1983). Meth. in Enzymol. Hirs, C.H.W. and Timasheff, S.N., eds. San Diego: Academic Press, 91, pp. 95-119.
- 8. Bradford, M.M. (1976). A rapid and sensitive method for the quantitation of microgram quantities of protein utilizing the principle of protein-dye binding. Anal. Biochem. 72, 248-254
- 9. Pande, S.V. and Murthy, S.R. (1994). A modified micro-Bradford procedure for elimination of interference from sodium dodecyl sulfate, other detergents and lipids. Anal. Biochem, 220, 424-426.
- 10. Brogdon, W.G. and Dickinson, C.M. (1983). A microassay system for measuring esterase activity and protein concentration in small samples and in high pressure liquid chromatography eluate fractions. Anal. Biochem. 131, 499-503.

- assay to microtiter plates. Anal. Biochem. 147, 144-147.
- 13. Ribin, R.W. and Warren, R.W. (1977). Quantitation of microgram amounts of protein in SDS-mercaptoethanol-Tris electrophoresis sample buffer. Anal. Biochem. 83, 773-777.
- 14. Bonde, M., Pontoppidan, H. and Pepper, D.S. (1992). Direct dye binding a quantitative assay for solid-phase immobilized protein. Anal. Biochem. 200, 195-198. 15. Alves Cordeiro, C.A. and Freire, A.P. (1994). Protein determination in permeabilized
- yeast cells using the Coomassie Brilliant Blue Dye Binding Assay. Anal. Biochem. 223, 321-323.
- 16. Sorensen, K. (1994). Coomassie Protein Assay Reagent used for quantitative determination of sodium cyanoborohydride (NaCNBH₃). Anal. Biochem. 218, 231-233.
- 17. Gadd, K.G. (1981). Protein estimation in spinal fluid using Coomassie blue reagent. Med. Lab. Sci. 38, 61-63.
- 18. Friedenauer, D. and Berlet, H.H. (1989). Sensitivity and variability of the Bradford protein assay in the presence of detergents. Anal. Biochem. 178, 263-268.
- 19. Splittgerber, A.G. and Sohl, J. (1989). Nonlinearity in protein assays by the Coomassie blue dye-binding method. Anal. Biochem. 179(1), 198-201.
- 20. Tsukada, T., et al. (1987). Identification of a region in the human vasoactive intestinal polypeptide gene responsible for regulation by cyclic AMP. J. Biol. Chem. 262(18), 8743-8747.

Introduction



Introduction

Protein quantitation is often necessary prior to handling protein samples for isolation and characterization. It is a required step before submitting protein samples for chromatographic, electrophoretic and immunochemical separation or analyses.

The most common methods for the colorimetric detection and quantitation of total protein can be divided into two groups based upon the chemistry involved. Protein assay reagents involve either protein-dye binding chemistry (Coomassie/Bradford) or protein-copper chelation chemistry. We offer numerous colorimetric assays for detection and quantitation of total protein. They are all well-characterized, robust assays that provide consistent, reliable results. Collectively, they represent the state-of-the-art for colorimetric detection and quantitation of total protein.

Selection of the Protein Assay

When it is necessary to determine the total protein concentration in a sample, one of the first issues to consider is the selection of a protein assay method. The choice among available protein assays usually is based upon the compatibility of the method with the samples to be assayed. The objective is to select a method that requires the least manipulation or pre-treatment of the samples containing substances that may interfere with the assay.

Table 1. Thermo Scientific Pierce Protein Assay Reagents and their working ranges.

| Reagent | Protocol Used | Estimated Working Range |
|---|--|-------------------------------|
| Pierce 600 nm | Standard tube | 25-2,000 μg/ml |
| Protein Assay | Standard microplate | 50-2,000 μg/ml |
| Coomassie (Bradford) | Standard tube or microplate | 100-1,500 μg/ml |
| Protein Assay | Micro tube or microplate | 1-25 μg/ml |
| Coomassie Plus | Standard tube or microplate | 100-1,500 μg/ml |
| (Bradford) Assay | Micro tube or microplate | 1-25 μg/ml |
| BCA Protein Assay – Reducing Agent Compatible | Standard tube or microplate | 125-2,000 µg/ml |
| BCA Protein Assay | Standard tube or microplate Enhanced tube | 20-2,000 μg/ml 5-250 μg/ml |
| Micro BCA | Standard tube | 0.5-20 μg/ml |
| Protein Assay | Standard microplate | 2-40 μg/ml |
| Modified Lowry | Standard protocol | 1-1,500 µg/ml |
| Protein Assay | Standard microplate | 10-1,500 µg/ml |

Each method has its advantages and disadvantages (see pages 1-3). Because no one reagent can be considered to be the ideal or best protein assay method, most researchers have more than one type of protein assay reagent available in their labs.

If the samples contain reducing agents or copper chelating reagents, either of the ready-to-use liquid Coomassie dye reagents (Coomassie [Bradford] Protein Assay or the Coomassie Plus Assay) would be excellent choices. The Modified Lowry Protein Assay offers all of the advantages of the original reagent introduced by Oliver Lowry in 1951 in a single, stable and ready-to-use reagent.

If the samples to be analyzed contain one or more detergents (at concentrations up to 5%), the BCA Protein Assay is the best choice. If the protein concentration in the detergent-containing samples is expected to be very low (< 20 μ g/ml), the Micro BCA Protein Assay may be the best choice. If the total protein concentration in the samples is high (> 2,000 μ g/ml), sample dilution can often be used to overcome any problems with known interfering substances.

Sometimes the sample contains substances that make it incompatible with any of the protein assay methods. The preferred method of dealing with interfering substances is to simply remove them. We offer several methods for performing this function, including dialysis, desalting, chemical blocking and protein precipitation followed by resolubilization. This handbook focuses on the last two methods. Chemical blocking involves treating the sample with something that prevents the interfering substance from causing a problem. Protein precipitation causes the protein to fall out of solution, at which time the interfering buffer can be removed and the protein resolubilized. The chemical treatment method, like that used in the BCA Protein Assay – Reducing Agent Compatible, or the Pierce 660 nm Protein Assay is generally preferred because, unlike protein precipitation, resolubilization of potentially hydrophobic proteins is not involved.

Selection of a Protein Standard

Selection of a protein standard is potentially the greatest source of error in any protein assay. Of course, the best choice for a standard is a highly purified version of the predominant protein found in the samples. This is not always possible or necessary. In some cases, all that is needed is a rough estimate of the total protein concentration in the sample. For example, in the early stages of purifying a protein, identifying which fractions contain the most protein may be all that is required. If a highly purified version of the protein of interest is not available or if it is too expensive to use as the standard, the alternative is to choose a protein that will produce a very similar color response curve with the selected protein assay method. For general protein assay work, bovine serum albumin (BSA) works well for a protein standard because it is widely available in high purity and is relatively inexpensive. Although it is a mixture containing several immunoglobulins, bovine gamma globulin (BGG) also is a good choice for a standard when determining the concentration of antibodies, because BGG produces a color response curve that is very similar to that of immunoglobulin G (IgG).

For greatest accuracy in estimating total protein concentration in unknown samples, it is essential to include a standard curve each time the assay is performed. This is particularly true for the protein assay methods that produce nonlinear standard curves. Determination of the number of standards and replicates used to define the standard curve depends upon the degree of nonlinearity in the standard curve and the degree of accuracy required. In general, fewer points are needed to construct a standard curve if the color response is linear. Typically, standard curves are constructed using at least two replicates for each point on the curve.

Preparation of Standards

Use this information as a guide to prepare a set of protein standards. Dilute the contents of one Albumin Standard (BSA) ampule into several clean vials, preferably using the same diluent as the sample(s). Each 1 ml ampule of 2.0 mg/ml Albumin Standard is

Preparation of Diluted Albumin (BSA) Standards for BCA Assay, BCA Reducing Agent-Compatible Assay and Pierce 660 nm Assay.

Dilution Scheme for Standard Test Tube Protocol and Microplate Procedure (Working Range = 20-2,000 $\mu g/ml)$

| Vial | Volume of Diluent | Volume and Source of BSA | Final BSA Concentration |
|------|----------------------|-----------------------------|----------------------------|
| А | 0 | 300 µl of stock | 2,000 µg/ml |
| В | 125 µl | 375 µl of stock | 1,500 µg/ml |
| С | 325 µl | 325 µl of stock | 1,000 µg/ml |
| D | 175 µl | 175 µl of vial B dilution | 750 µg/ml |
| E | 325 µl | 325 µl of vial C dilution | 500 µg/ml |
| F | 325 µl | 325 µl of vial E dilution | 250 µg/ml |
| G | 325 µl | 325 µl of vial F dilution | 125 µg/ml |
| Н | 400 µl | 100 µl of vial G dilution | 25 µg/ml |
| 1 | 400 µl | 0 | 0 µg/ml = Blank |

Dilution Scheme for Enhanced Test Tube Protocol (Working Range = $5-250 \mu g/ml$)

| Vial | Volume of Diluent | Volume and Source of BSA | Final BSA Concentration |
|------|----------------------|--------------------------------|----------------------------|
| А | 700 µl | 100 µl of stock | 250 µg/ml |
| В | 400 µl | 400 μl of vial A dilution | 125 µg/ml |
| С | 450 µl | 300 µl of vial B dilution | 50 µg/ml |
| D | 400 µl | 400 µl of vial C dilution | 25 µg/ml |
| Е | 400 µl | 100 µl of vial D dilution | 5 µg/ml |
| F | 400 µl | 0 | 0 µg/ml = Blank |

Preparation of Diluted Albumin (BSA) Standards for Micro BCA Assay.

| Vial | Volume of Diluent | Volume and Source of BSA | Final BSA Concentration |
|------|----------------------|-----------------------------|----------------------------|
| А | .45 ml | 0.5 µl of stock | 200 µg/ml |
| В | 8.0 ml | 2.0 µl of vial A dilution | 40 µg/ml |
| С | 4.0 ml | 4.0 µl of vial B dilution | 20 µg/ml |
| D | 4.0 ml | 4.0 µl of vial C dilution | 10 µg/ml |
| E | 4.0 ml | 4.0 µl of vial D dilution | 5 μg/ml |
| F | 4.0 ml | 4.0 μl of vial E dilution | 2.5 μg/ml |
| G | 4.8 ml | 3.2 µl of vial F dilution | 1 µg/ml |
| Н | 4.0 ml | 4.0 µl of vial G dilution | 0.5 µg/ml |
| 1 | 8.0 ml | 0 | 0 µg/ml = Blank |

sufficient to prepare a set of diluted standards for either working range suggested. There will be sufficient volume for three replications of each diluted standard.

Preparation of Protein Standards for Coomassie Plus (Bradford) Assay and Coomassie (Bradford) Assay.

Dilution Scheme for Standard Test Tube and Microplate Protocols (Working Range = $100-1,500 \mu g/ml$)

| Vial | Volume of Diluent | Volume and Source of BSA | Final BSA Concentration |
|------|----------------------|--------------------------------|----------------------------|
| А | 0 | 300 µl of stock | 2,000 µg/ml |
| В | 125 µl | 375 µl of stock | 1,500 µg/ml |
| С | 325 µl | 325 µl of stock | 1,000 µg/ml |
| D | 175 µl | 175 μl of vial B dilution | 750 µg/ml |
| E | 325 µl | 325 µl of vial C dilution | 500 µg/ml |
| F | 325 µl | 325 µl of vial E dilution | 250 µg/ml |
| G | 325 µl | 325 µl of vial F dilution | 125 µg/ml |
| Н | 400 µl | 100 μl of vial G dilution | 25 µg/ml |
| I | 400 µl | 0 | 0 µg/ml = Blank |

Dilution Scheme for Micro Test Tube or Microplate Protocols (Working Range = 1-25 $\mu g/ml)$

| Vial | Volume of Diluent | Volume and Source of BSA | Final BSA Concentration |
|------|----------------------|-----------------------------|----------------------------|
| А | 2,370 µl | 30 µl of stock | 25 µg/ml |
| В | 4,950 µl | 50 µl of stock | 20 µg/ml |
| С | 3,970 µl | 30 µl of stock | 15 µg/ml |
| D | 2,500 µl | 2,500 µl of vial B dilution | 10 µg/ml |
| E | 2,000 µl | 2,000 µl of vial D dilution | 5 μg/ml |
| F | 1,500 µl | 1,500 µl of vial E dilution | 2.5 µg/ml |
| G | 5,000 µl | 0 | 0 µg/ml = Blank |

Preparation of Diluted Albumin (BSA) for Modified Lowry Assay.

Dilution Scheme for Test Tube and Microplate Procedure (Working Range = 1-1,500 μg/ml)

| Vial | Volume of Diluent | Volume and Source of BSA | Final BSA Concentration |
|------|----------------------|-----------------------------|----------------------------|
| А | 250 µl | 750 µl of stock | 200 µg/ml |
| В | 625 µ | 625 µl of stock | 40 µg/ml |
| С | 310 µ | 310 µl of vial A dilution | 20 µg/ml |
| D | 625 µl | 625 µl of vial B dilution | 10 µg/ml |
| E | 625 µl | 625 µl of vial D dilution | 5 μg/ml |
| F | 625 µl | 625 µl of vial E dilution | 2.5 µg/ml |
| G | 800 µl | 200 µl of vial F dilution | 1 µg/ml |
| Н | 800 µl | 200 µl of vial G dilution | 0.5 µg/ml |
| 1 | 800 µl | 200 µl of vial H dilution | 0 µg/ml = Blank |
| J | 1,000 µl | 0 | 0 µg/ml = Blank |

Standards for Total Protein Assay

Bovine Serum Albumin Standard

The Thermo Scientific Pierce BSA Standard ... the most relied-upon albumin standard for total protein determination measurements.



Ordering Information

| Product | Description | Pkg. Size |
|---------|--|-----------|
| 23209 | Albumin Standard Ampules, 2 mg/ml Contains: Bovine Albumin in 0.9% NaCl solution containing sodium azide | 10 x 1 ml |
| 23210 | Albumin Standard, 2 mg/ml Contains: Bovine Albumin in 0.9% NaCl solution containing sodium azide | 50 ml |

Bovine Gamma Globulin Standard

Easy-to-use, 2 mg/ml BGG solution. Ampuled to preserve product integrity. An excellent choice for IgG total protein determination. Recommended for Coomassie (Bradford) Assays.

| Product | Description | Pkg. Size |
|---------|---|-----------|
| 23212 | Bovine Gamma Globulin Standard 2 mg/ml Contains: Bovine Gamma Globulin Fraction II in 0.9% NaCl solution containing sodium azide | 10 x 1 ml |

Mammalian Gamma Globulins for Standards:

| Description | Pkg. Size |
|-----------------------|--|
| Mouse Gamma Globulin | 10 mg |
| Rabbit Gamma Globulin | 10 mg |
| Rat Gamma Globulin | 10 mg |
| Goat Gamma Globulin | 10 mg |
| Human Gamma Globulin | 10 mg |
| | Mouse Gamma Globulin Rabbit Gamma Globulin Rat Gamma Globulin Goat Gamma Globulin |

Thermo Scientific Total Protein Assays

Pre-Diluted BSA and BGG Protein Assay Standard Sets

Construct a standard curve for most protein assay methods as fast as you can pipette.



Highlights:

- Stable and sterile filtered
- · Ideal for BCA and Bradford-based protein assays
- Standard curve range: 125-2,000 μg/ml
- · Seven data points within the range
- Sufficient materials to prepare 15-35 standard tube protocol curves or 175-350 standard microplate protocol curves running duplicate data points
- Convenient no need to prepare a diluted standard series for each determination
- Consistent no need to worry about variability in dilutions from day to day or person to person
- More reliable protein quantitation because of the assured accuracy of the concentrations of each standard
- Dramatically improved speed to result, especially with Bradford-based protein assays

Sample Preparation

Before a sample is analyzed for total protein content, it must be solubilized, usually in a buffered aqueous solution. The entire process is usually performed in the cold, with additional precautions taken to inhibit microbial growth or to avoid casual contamination of the sample by foreign debris such as hair, skin or body oils.

When working with tissues, cells or solids, the first step of the solubilization process is usually disruption of the sample's cellular structure by grinding and/or sonication or by the use of specially designed reagents (e.g., Thermo Scientific Pierce Cell Lysis Reagents) containing surfactants to lyse the cells. This is done in aqueous buffer containing one or more surfactants to aid the

solubilization of the membrane-bound proteins, biocides (antimicrobial agents) and protease inhibitors. After filtration or centrifugation to remove the cellular debris, additional steps such as sterile filtration, removal of lipids or further purification of the protein of interest from the other sample components may be necessary.

Nonprotein substances in the sample that are expected to interfere in the chosen protein assay method may be removed by dialysis with Thermo Scientific Slide-A-Lyzer Dialysis Cassettes or Thermo Scientific SnakeSkin Dialysis Tubing, gel filtration with Thermo Scientific Desalting Columns or D Detergent Removing Gel, or precipitation as in the Compat-Able[™] Protein Assays or SDS-Out Reagent.

Protein:protein Variation

Each protein in a sample is unique and can demonstrate that individuality in protein assays as variation in the color response. Such protein:protein variation refers to differences in the amount of color (absorbance) obtained when the same mass of various proteins is assayed concurrently by the same method. These differences in color response relate to differences in amino acid sequence, isoelectric point (pl), secondary structure and the presence of certain side chains or prosthetic groups.

Table 2 (page 9) shows the relative degree of protein:protein variation that can be expected with our different protein assay reagents. This differential may be a consideration in selecting a protein assay method, especially if the relative color response ratio of the protein in the samples is unknown. As expected, protein assay methods that share the same basic chemistry show similar protein:protein variation. These data make it obvious why the largest source of error for protein assays is the choice of protein for the standard curve.

Ordering Information

| Product | Description | Pkg. Size |
|---------|---|-----------|
| 23208 | Pre-Diluted Protein Assay Standards Bovine Serum Albumin (BSA) Set Diluted in 0.9% saline and preserved with 0.05% sodium azide Includes: 7 x 3.5 ml of standardized BSA solutions each at a specific concentration along a range from 125-2,000 μg/ml | Kit |
| 23213 | Pre-Diluted Protein Assay Standards: Bovine Gamma Globulin Fraction II (BGG) Set Diluted in 0.9% saline and preserved with 0.05% sodium azide Includes: 7 x 3.5 ml of standardized BGG solutions each at a specific concentration along a range from 125-2,000 µg/ml | Kit |

Total Protein Assays

For each of the six methods presented here, a group of 14 proteins was assayed using the standard protocol in a single run. The net (blank corrected) average absorbance for each protein was calculated. The net absorbance for each protein is expressed as a ratio to the net absorbance for BSA. If a protein has a ratio of 0.80, it means that the protein produces 80% of the color obtained for an equivalent mass of BSA. All of the proteins tested using the standard tube protocol with the BCA Protein Assay, the Modified Lowry Protein Assay, the Coomassie (Bradford) Assay and the Coomassie Plus (Bradford) Assay were at a concentration of 1,000 μ g/ml.

Compatible and Incompatible Substances

An extensive list of substances that have been tested for compatibility with each protein assay reagent can be found in the instruction booklet that accompanies each assay product. A copy can also be obtained from our web site. In summary, the Coomassie (Bradford) and the Coomassie Plus (Bradford) Assays will tolerate the presence of most buffer salts, reducing substances and chelating agents, but they will not tolerate the presence of detergents (except in very low concentrations) in the sample. Strong acids or bases, and even some strong buffers, may interfere if they alter the pH of the reagent.

The Modified Lowry Protein Assay is sensitive to the presence of reducing substances, chelating agents and strong acids or strong bases in the sample. In addition, the reagent will be precipitated by the presence of detergents and potassium ions in the sample.

The BCA Protein Assay is tolerant of most detergents but is sensitive to the presence of reducing substances, chelating agents and strong acids or strong bases in the sample. In general, the Micro BCA Protein Assay is more sensitive to the same substances that interfere with the BCA Protein Assay because less dilution of the sample is used.

Table 2. Protein:protein variation.

| | Pierce 660 nm Assay Ratio | BCA Ratio | Micro BCA Ratio | Modified Lowry Ratio | Coomassie (Bradford) Ratio | Coomassie Plus Ratio | Bio-Rad (Bradford) Ratio |
|-------------------------------|---------------------------------|--------------|-----------------------|----------------------------|----------------------------------|----------------------------|--------------------------------|
| 1. Albumin, bovine serum | 1.00 | 1.00 | 1.00 | 1.00 | 1.00 | 1.0 | 1.00 |
| 2. Aldolase, rabbit muscle | 0.83 | 0.85 | 0.80 | 0.76 | 0.76 | 0.74 | 0.97 |
| 3. α -Chymotrypsinogen | х | 1.14 | 0.99 | 0.48 | 0.48 | 0.52 | 0.41 |
| 4. Cytochrome C, horse heart | 1.22 | 0.83 | 1.11 | 1.07 | 1.07 | 1.03 | 0.48 |
| 5. Gamma Globulin, bovine | 0.51 | 1.11 | 0.95 | 0.56 | 0.56 | 0.58 | 0.58 |
| 6. IgG, bovine | х | 1.21 | 1.12 | 0.58 | 0.58 | 0.63 | 0.65 |
| 7. IgG, human | 0.57 | 1.09 | 1.03 | 0.63 | 0.63 | 0.66 | 0.70 |
| 8. lgG, mouse | 0.48 | 1.18 | 1.23 | 0.59 | 0.59 | 0.62 | 0.60 |
| 9. lgG, rabbit | 0.38 | 1.12 | 1.12 | 0.37 | 0.37 | 0.43 | 0.53 |
| 10. lgG, sheep | х | 1.17 | 1.14 | 0.53 | 0.53 | 0.57 | 0.53 |
| 11. Insulin, bov. pancreas | 0.81 | 1.08 | 1.22 | 0.60 | 0.60 | 0.67 | 0.14 |
| 12. Myoglobin, horse heart | 1.18 | 0.74 | 0.92 | 1.09 | 1.19 | 1.15 | 0.89 |
| 13. Ovalbumin | 0.54 | 0.93 | 1.08 | 0.32 | 0.32 | 0.68 | 0.27 |
| 14. Transferrin, human | 0.80 | 0.89 | 0.98 | 0.84 | 0.84 | 0.90 | 0.95 |
| Avg. ratio | 0.7364 | 1.02 | 1.05 | 0.68 | 0.68 | 0.73 | 0.60 |
| S.D. | 0.2725 | 0.15 | 0.12 | 0.26 | 0.26 | 0.21 | 0.28 |
| CV | 37% | 14.7% | 11.4% | 38.2% | 38.2% | 28.8% | 46% |
| | | | | | | | |

1. All of the proteins were tested using the standard tube protocol with the Micro BCA Protein Assay at a protein concentration of 10 µg/ml.

This table is a useful guideline to estimate the protein:protein variation in color response that can be expected with each method. It does not tell the whole story. However, because the comparisons were made using a single protein concentration, it is not apparent that the color response ratio also varies with changes in protein concentration.

Substances Compatible with Thermo Scientific Pierce Protein Assays

Concentrations listed refer to the actual concentration in the protein sample. A blank indicates that the material is incompatible with the assay; n/a indicates the substance has not been tested in that respective assay.

| | Pierce 660 nm Protein Assay | BCA – Reducing Agent Compatible Assay | BCA Assay | Micro BCA Assay | Modified Lowry Assay | Coomassie (Bradford) Assay | Coomassie Plus Assay |
|--|--------------------------------|--|---------------|--------------------|-------------------------|----------------------------------|-------------------------|
| Detergents | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) |
| Brij-35 | 5.0% | | 5.0% | 5.0% | 0.031% | .125% | 0.062% |
| Brij-56 | _ | _ | 1.0% | 1.0% | 0.062% | 0.031% | 0.031% |
| Brij-58 | _ | _ | 1.0% | 1.0% | 0.062% | 0.031% | 0.016% |
| CHAPS | 5.0% | 10.0% | 5.0% | 1.0% | 0.062% | 5.0% | 5.0% |
| CHAPSO | 4.0% | _ | 5.0% | 5.0% | 0.031% | 5.0% | 5.0% |
| Deoxycholic acid | | | 5.0% | 5.0% | n/a | 0.05% | 0.04% |
| Lubrol PX | _ | _ | 1.0% | 1.0% | 0.031% | 0.125% | 0.031% |
| Octyl glucoside | 5.0% | _ | 5.0% | 1.0% | 0.031% | 0.5% | 0.5% |
| Nonidet P-40 | 5.0% | _ | 5.0% | 5.0% | 0.016% | 0.5% | 0.5% |
| Octyl β-thioglucopyranoside | 10.0% | 10.0% | 5.0% | 5.0% | n/a | 3.0% | 3.0% |
| SDS (Lauryl) | 5.0% | 10.0% | 5.0% | 5.0% | 1.0% | 0.125% | 0.016% |
| Span 20 | | | 1.0% | 1.0% | 0.25% | 0.5% | 0.5% |
| Triton X-100 | 1.0% | 10.0% | 5.0% | 5.0% | 0.031% | 0.125% | 0.062% |
| Triton X-114 | | 2.0% | 1.0% | 0.5% | 0.031% | 0.125% | 0.062% |
| Triton X-305 | _ | | 1.0% | 1.0% | 0.031% | 0.5% | 0.125% |
| Triton X-405 | | · · · · · · · · · · · · · · · · · · · | 1.0% | 1.0% | 0.031% | 0.5% | 0.25% |
| Tween-20 | 10.0% | 10.0% | 5.0% | 5.0% | 0.062% | 0.062% | 0.031% |
| Tween-60 | _ | | 5.0% | 0.5% | n/a | 0.1% | 0.025% |
| Tween-80 | _ | | 5.0% | 5.0% | 0.031% | 0.062% | 0.016% |
| Zwittergent 3-14 | | ······ | 1.0% | | n/a | 0.025% | 0.025% |
| Salts/Buffers | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) |
| ACES, pH 7.8 | | | 25 mM | 10 mM | n/a | 100 mM | 100 mM |
| Ammonium sulfate | | · · · · · · · · · · · · · · · · · · · | 1.5 M | | | 1.0 M | 1.0 M |
| Asparagine | _ | | 1 mM | n/a | 5 mM | 10 mM | 10 mM |
| Bicine, pH 8.4 | | | 20 mM | 2 mM | n/a | 100 mM | 100 mM |
| Bis-Tris, pH 6.5 | _ | | 33 mM | 0.2 mM | n/a | 100 mM | 100 mM |
| Borate (50 mM), pH 8.5 | | | undiluted | 1:4 dilution* | n/a | undiluted | undiluted |
| B-PER Reagent | undiluted | 40.0% | undiluted | 1:10 dilution* | n/a | 1:2 dilution* | 1:2 dilution* |
| Calcium chloride in TBS, pH 7.2 | | | 10 mM | 10 mM | n/a | 10 mM | 10 mM |
| Na-Carbonate/Na-Bicarbonate (0.2 M), pH 9.4 | _ | | undiluted | undiluted | n/a | undiluted | undiluted |
| Cesium bicarbonate | | | 0.1 M | 0.1 M | 50 mM | 0.1 M | 0.1 M |
| CHES, pH 9.0 | | | 100 mM | 100 mM | n/a | 100 mM | 100 mM |
| Na-Citrate (0.6 M), Na-Carbonate (0.1 M), pH 9.0 | | | 1:8 dilution* | 1:600 dilution* | n/a | undiluted | undiluted |
| Na-Citrate (0.6 M), MOPS (0.1 M) pH 7.5 | _ | | 1:8 dilution* | 1:600 dilution* | n/a | undiluted | undiluted |
| Cobalt chloride in TBS, pH 7.2 | | | 0.8 mM | | n/a | 10 mM | 10 mM |
| EPPS, pH 8.0 | | | 100 mM | 100 mM | n/a | 100 mM | 100 mM |
| Ferric chloride in TBS, pH 7.2 | | | 10 mM | 0.5 mM | n/a | 10 mM | 10 mM |
| Glycine | 100 mM | 50 mM | 1 mM | n/a | 100 mM | 0.1 M | 0.1 M |
| Guanidine•HCl | 2.5 M | 2 M | 4 M | 4.0 M | n/a | 3.5 M | 3.5 M |
| HEPES, pH 7.5 | 100 mM | 500 mM | 100 mM | 100 mM | 1 mM | 0.1 M | 0.1 M |
| Imidazole, pH 7.0 | 200 mM | 50 mM | 50 mM | 12.5 mM | 25 mM | 200 mM | 200 mM |
| MES, pH 6.1 | 125 mM | 100 mM | 100 mM | 100 mM | 125 mM | 100 mM | 100 mM |
| MES (0.1 M), NaCl (0.9%), pH 4.7 | | _ | undiluted | 1:4 dilution* | n/a | undiluted | undiluted |
| MOPS, pH 7.2 | 125 mM | 200 mM | 100 mM | 100 mM | n/a | 100 mM | 100 mM |
| Molified Dulbecco's PBS, pH 7.4 | 125 mil | undiluted | undiluted | undiluted | n/a | undiluted | undiluted |
| Nickel chloride in TBS, pH 7.2 | | | 10 mM | 0.2 mM | n/a | 10 mM | 10 mM |
| PBS; Phosphate (0.1 M), NaCl (0.15 M), pH 7.2 | undiluted | undiluted | undiluted | undiluted | n/a | undiluted | undiluted |
| PIPES, pH 6.8 | | | 100 mM | 100 mM | n/a | 100 mM | 100 mM |
| RIPA lysis buffer; 50 mM Tris, 150 mM NaCl, | | | | | | | |
| 0.5% DOC, 1% NP-40, 0.1% SDS, pH 8.0 | undiluted | _ | undiluted | 1:10 dilution* | n/a | 1:10 dilution* | 1:40 dilution* |

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| | Pierce 660 nm Protein Assay | BCA – Reducing Agent Compatible Assay | BCA Assay | Micro BCA Assay | Modified Lowry Assay | Coomassie (Bradford) Assay | Coomassie Plus Assay |
|--|--------------------------------|--|---------------|--------------------|-------------------------|----------------------------------|-------------------------|
| Salts/Buffers (cont.) | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) |
| Sodium acetate | _ | _ | 0.2 M | 0.2 M | 0.2 M | 180 mM | 180 mM |
| Sodium azide | _ | _ | 0.2% | 0.2% | 0.2% | 0.5% | 0.5% |
| Sodium bicarbonate | _ | _ | 0.1 M | 0.1 M | 0.1 M | 0.1 M | 0.1 M |
| Sodium chloride | 1.25 M | | 1.0 M | 1.0 M | 1.0 M | 5.0 M | 5.0 M |
| Sodium citrate, pH 4.8 (or pH 6.4) | | _ | 200 mM | 5 mM (16.7 mM) | n/a | 200 mM | 200 mM |
| Sodium phosphate | _ | _ | 0.1 M | 0.1 M | 0.1 M | 0.1 M | 0.1 M |
| Tricine, pH 8.0 | | _ | 25 mM | 2.5 mM | n/a | 100 mM | 100 mM |
| Triethanolamine, pH 7.8 | _ | _ | 25 mM | 0.5 mM | n/a | 100 mM | 100 mM |
| Tris | 250 mM | 50 mM | 0.25 M | 0.05 M | 10 mM | 2.0 M | 2.0 M |
| TBS; Tris (25 mM), NaCl (0.15 M), pH 7.6 | _ | undiluted | undiluted | 1:10 dilution* | n/a | undiluted | undiluted |
| Tris (25 mM), Glycine (192 mM), pH 8.0 | _ | _ | 1:3 dilution* | 1:10 dilution* | n/a | undiluted | undiluted |
| Tris (25 mM), Glycine (192 mM), SDS (0.1%), pH 8.3 | _ | _ | undiluted | undiluted | n/a | 1:2 dilution* | 1:4 dilution* |
| Zinc chloride in TBS, pH 7.2 | | | 10 mM | 0.5 mM | n/a | 10 mM | 10 mM |
| Reducing agents | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) |
| N-acetylglucosamine in PBS, pH 7.2 | _ | _ | 10 mM | _ | n/a | 100 mM | 100 mM |
| Ascorbic acid | 500 mM | _ | _ | _ | 1 mM | 50 mM | 50 mM |
| Catecholamines | | | — | | n/a | n/a | n/a |
| Creatinine | _ | | | _ | n/a | n/a | n/a |
| Glucose | | _ | 10 mM | 1 mM | 0.1 mM | 1.0 M | 1.0 M |
| Melibiose | | | | n/a | 25 mM | 0.1 M | 0.1 M |
| Potassium thiocyanate | | _ | 3.0 M | n/a | 0.1 M | 3.0 M | 3.0 M |
| Thiol-containing agents | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) |
| Cysteine | 350 mM | _ | _ | | 1 mM | 10 mM | 10 mM |
| Dithioerythritol (DTE) | | | 1 mM | | | 1 mM | 1 mM |
| Dithiothreitol (DTT) | 500 mM | 5 mM | 1 mM | _ | _ | 5 mM | 5 mM |
| 2-Mercaptoethanol | 1 M | 35 mM | 0.01% | 1 mM | 1 mM | 1.0 M | 1.0 M |
| TCEP | 40 mM | 10 mM | | _ | _ | | _ |
| Thimerosal | | _ | 0.01% | _ | 0.01% | 0.01% | 0.01% |
| Chelating agents | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) |
| EDTA | 20 mM | 20 mM | 10 mM | 0.5 mM | 1 mM | 100 mM | 100 mM |
| EGTA | 20 mM | _ | | _ | 1 mM | 2 mM | 2 mM |
| Sodium citrate, pH 4.8 (6.4) | | 100 mM | 200 mM | 5 mM (16.7 mM) | 0.1 mM | 200 mM | 200 mM |
| Solvents/Misc. | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) | (≤) |
| Acetone | _ | _ | 10% | 1.0% | 10% | 10% | 10% |
| Acetonitrile | 50% | 30% | 10% | 1.0% | 10% | 10% | 10% |
| Aprotinin | _ | _ | 10 mg/L | 1 mg/L | 10 mg/L | 10 mg/L | 10 mg/L |
| DMF | | _ | 10% | 1.0% | 10% | 10% | 10% |
| DMSO | | _ | 10% | 1.0% | 10% | 10% | 10% |
| Ethanol | | | 10% | 1.0% | 10% | 10% | 10% |
| Glycerol (fresh) | 50% | 5% | 10% | 1.0% | 10% | 10% | 10% |
| Guanidine•HCl | 2.5 M | _ | 4.0 M | 4.0 M | 0.1 M | 3.5 M | 3.5 M |
| Hydrochloric acid | 125 mM | _ | 0.1 M | 0.01 M | 0.1 M | 0.1 M | 0.1 M |
| Leupeptin | _ | | 10 mg/L | 10 mg/L | 10 mg/L | 10 mg/L | 10 mg/L |
| Methanol | 50% | _ | 10% | 1.0% | 10% | 10% | 10% |
| Phenol Red | 0.5 mg/ml | 3.125 μg/ml | _ | n/a | 0.01 mg/ml | 0.5 mg/ml | 0.5 mg/ml |
| PMSF | | | 1 mM | 1 mM | 1 mM | 1 mM | 1 mM |
| Sodium hydroxide | 125 mM | | 0.1 M | 0.05 M | 0.1 M | 0.1 M | 0.1 M |
| Sucrose | 50% | 40% | 40% | 4% | 7.5% | 10% | 10% |
| TLCK | | | 0.1 mg/L | 0.1 mg/L | 0.01 mg/L | 0.1 mg/L | 0.1 mg/L |
| ТРСК | _ | — | 0.1 mg/L | 0.1 mg/L | 0.1 mg/L | 0.1 mg/L | 0.1 mg/L |
| Urea | 8 M | 4.0 M | 3.0 M | 3.0 M | 3.0 M | 3.0 M | 3.0 M |
| o-Vanadate (sodium salt) in PBS pH 7.2 | _ | _ | 1 mM | 1 mM | n/a | 1 mM | 1 mM |
| | | | | | | | |

n/a: not assayed

A blank indicates that the material is incompatible with the assay

* Diluted with distilled/deionized water

Time Considerations

The amount of time required to complete a total protein assay will vary for the different colorimetric, total protein assay methods presented. To compare the amount of time required to perform each assay, all seven assays were performed using 20 samples and eight standards (including the blank). Each sample or standard was assayed in duplicate using the standard tube protocol (triplicate using the plate). The estimates include times for both incubation(s) and handling:

- Preparing (diluting) the standard protein in the diluent buffer (10 minutes)
- Organizing the run and labeling the tubes (5 minutes)
- Pipetting the samples and reagents (10 minutes for 56 tubes, 1 minute per plate)
- · Mixing or incubating the tubes or plates (varies)
- Measuring the color produced (15 minutes for 56 tubes or 1 minute per plate)
- Graphing the standard curve, calculating, recording and reporting the results (30 minutes)

Table 3. Times required to assay 20 samples and 8 standards using the test tube procedure; handling times are considerably less using the microplate procedure.

| Method | Product # | Incubation Time | Total Assay Time |
|---|-----------|------------------------------|------------------|
| Pierce 600 nm Protein Assay | 23250 | 5 minutes | 75 minutes |
| Coomassie Plus (Bradford) Assay | 23236 | 10 minutes | 80 minutes |
| Coomassie (Bradford) Assay | 23200 | 10 minutes | 80 minutes |
| BCA Assay | 23225 | 30 minutes | 100 minutes |
| Modified Lowry Assay | 23240 | 10 minutes and 30 minutes | 110 minutes |
| BCA Protein Assay – Reducing Agent Compatible | 23250 | 45 minutes | 115 minutes |
| Micro BCA Assay | 23235 | 60 minutes | 130 minutes |

Calculation of Results

When calculating protein concentrations manually, it is best to use point-to-point interpolation. This is especially important if the standard curve is nonlinear. Point-to-point interpolation refers to a method of calculating the results for each sample using the equation for a linear regression line obtained from just two points on the standard curve. The first point is the standard that has an absorbance just below that of the sample and the second point is the standard that has an absorbance just above that of the sample. In this way, the concentration of each sample is calculated from the most appropriate section of the whole standard curve. Determine the average total protein concentration for each sample from the average of its replicates. If multiple dilutions of each sample have been assayed, average the results for the dilutions that fall within the most linear portion of the working range.

When analyzing results with a computer, use a quadratic curve fit for the nonlinear standard curve to calculate the protein concentration of the samples. If the standard curve is linear, or if the absorbance readings for your samples fall within the linear portion of the standard curve, the total protein concentrations of the samples can be estimated using the linear regression equation.

Most software programs allow one to construct and print a graph of the standard curve, calculate the protein concentration for each sample, and display statistics for the replicates. Typically, the statistics displayed will include the mean absorbance readings (or the average of the calculated protein concentrations), the standard deviation (SD) and the coefficient of variation (CV) for each standard or sample. If multiple dilutions of each sample have been assayed, average the results for the dilutions that fall in the most linear portion of the working range.

References

Krohn, R.I. (2002). The colorimetric detection and quantitation of total protein, Current Protocols in Cell Biology, A3.H.1-A.3H.28, John Wiley & Sons, Inc.

Krohn, R.I. (2001). The colorimetric determination of total protein, *Current Protocols in Food Analytical Chemistry*, B1.1-B1.1.27, John Wiley & Sons, Inc.

Thermo Scientific Pierce 660 nm Protein Assays

Rapid, reproducible and colorimetric.

Accurate protein concentration measurements are required to study many biochemical processes. Although there are several methods for quantifying proteins, colorimetric or chromogenic methods remain popular because of their relative simplicity and speed. The most commonly used dye-binding protein assay is the Bradford assay,¹ which is based on coomassie dye binding to proteins. The Bradford assay, however, is prone to inaccuracy from its typical non-linear standard curves. Moreover, the assay is not compatible with samples containing detergents at commonly used concentrations. The new Pierce 660 nm Protein Assay is highly reproducible, rapid and more linear than the Bradford method. Furthermore, it is compatible with commonly used detergents and reducing agents.

Highlights:

- Accurate results standard curves are more linear than the Bradford method
- Versatile compatible with commonly used detergents and reducing agents and with samples lysed in Laemmli sample buffer
- Fast single reagent with a simple mix-and-read assay
- Flexible available in test tube and microplate formats
- **Economical** use small volumes of valuable samples: 10 μ l in microplate and 100 μ l in standard procedures
- **Convenient** room temperature storage means no waiting for the reagent to warm-up before use

Every protein assay has limitations depending on the application and the specific protein sample analyzed. The most useful features to consider when choosing a protein assay are sensitivity (lower detection limit), compatibility with common substances in samples (e.g., detergents, reducing agents, chaotropic agents, inhibitors, salts and buffers), standard curve linearity and protein-to-protein variation. Current methods for the colorimetric determination of protein concentration in solution include the Coomassie Blue G-250 dye-binding assay,¹ Biuret method,² the Lowry method,³ the bicinchoninic acid (BCA) assay⁴ and colloidal gold protein assay.⁵

The new Pierce 660 nm Protein Assay is based on the binding of a proprietary dye-metal complex to protein in acidic conditions that causes a shift in the dye's absorption maximum, which is measured at 660 nm. To demonstrate the effect of protein binding to the dye-metal complex, we performed spectral analysis of the dye with and without metal and in the presence and absence of BSA. The absorption maximum of the dye-metal complex shifts proportionally upon binding to BSA (Figure 1).

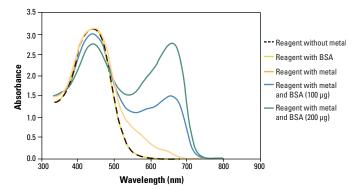


Figure 1. The absorption maximum of the reagent-metal complex shifts proportionally upon binding to BSA. The absorption spectra were recorded for the Thermo Scientific Pierce 660 nm Protein Assay Reagent from 340 to 800 nm using a Varian Cary[®] Spectrophotometer. The assay reagent is a proprietary dye-metal complex that binds to protein in acidic conditions, which shifts the dye's absorption maximum.

The dye-metal complex is reddish-brown that changes to green upon protein binding. The color produced in the assay is stable and increases in proportion to a broad range of increasing protein concentrations. The color change is produced by deprotonation of the dye at low pH facilitated by protein-binding interactions through positively charged amino acid groups and the negatively charged deprotonated dye-metal complex.

The linear detection ranges for BSA are 25-2,000 μ g/ml for the test tube assay and 50-2,000 μ g/ml for the microplate assay. The linear range for BGG is 50-2,000 μ g/ml for both the test tube and microplate assays (Figures 2 and 3). The assay has a moderate protein-to-protein variation of 37% and is more linear compared with the Bradford assay and, thus, produces more accurate results (Figure 4). The Pierce 660 nm Protein Assay color development is significantly greater with BSA than with most other proteins, including BGG. Therefore, BSA is a suitable standard if the sample contains primarily albumin, or if the protein being assayed has a similar response to the dye as BSA. For a color response that is typical of globulins, BGG is an appropriate standard protein.

The new Pierce 660 nm Protein Assay is compatible with high concentrations of most detergents, reducing agents and other commonly used reagents. Additionally, by simply adding Ionic Detergent Compatibility Reagent (IDCR) to the assay reagent, the assay is compatible with samples containing Laemmli SDS sample buffer with bromophenol blue and many common ionic detergents. IDCR completely dissolves by thorough mixing and does not have any affect on the assay. In conclusion, the Pierce 660 nm Protein Assay is a detergent- and reducing agent-compatible protein assay that is linear over wide range of concentrations. The simple mix-and-read format is easy to use, providing researchers a fast method for accurate protein quantitation.

Methods:

Spectral Analysis: The absorption spectra from 340 to 800 nm were recorded using a Varian Cary Spectrophotometer of the following component combinations: the Pierce 660 nm Protein Assay Reagent alone and in the presence of the transition metal; 100 µg of bovine serum albumin (BSA) and reagent with and without metal; and 200 µg of BSA with the reagent and metal.

Typical Response Curves

Test Tube Procedure: To each test tube containing 0.1 ml of BSA or BGG standard replicate (25, 50, 125, 250, 500, 750, 1000, 1500 and 2000 µg/ml) in saline, 1.5 ml of the Pierce 660 nm Protein Assay Reagent was added, mixed well and incubated at room temperature for 5 minutes. The absorbance of all samples and controls was measured at 660 nm. The average absorbance for the blank replicates (control) was subtracted from the absorbance for individual standard replicates. A standard curve was generated by plotting the average blank-corrected 660 nm measurement for each standard versus its concentration. For a comparison study, the standard Bio-Rad Bradford Assay was performed as per manufacturer's directions.

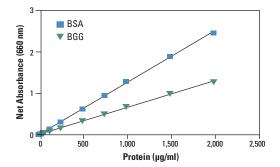


Figure 2. Typical color response curves using the test tube procedure. The linear detection ranges are 25-2,000 μ g/ml for bovine serum albumin (BSA) and 50-2,000 μ g/ml for bovine gamma globulin (BGG). The average absorbance for the blank replicates (control) was subtracted from the absorbance for individual standard replicates.

Microplate Procedure: To each well containing 0.01 ml of BSA or BGG standard replicate (25, 50, 125, 250, 500, 750, 1000, 1500 and 2000 μ g/ml) in saline, 0.15 ml of the Pierce 660 nm Protein Assay Reagent was added. The plate was covered with sealing tape, mixed for one minute on a plate shaker and incubated at room temperature for five minutes. The plate reader was set to 660 nm and using the control as a blank, the absorbance of all samples was measured. A standard curve was generated by plotting the average blank-corrected 660 nm measurement for each standard versus its concentration.

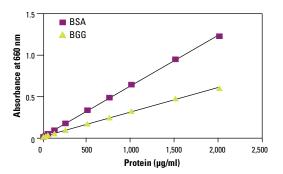


Figure 3. Typical color response curves using the microplate procedure. The linear detection range is 50-2,000 μ g/ml for bovine serum albumin (BSA) and bovine gamma globulin (BGG). The average absorbance for the blank replicates (control) was subtracted from the absorbance for individual standard replicates.

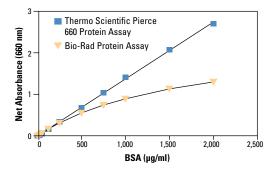


Figure 4. Performance comparison of Bradford Protein Assay versus the Thermo Scientific Pierce 660 nm Protein Assay. Assays were performed according to the standard test-tube procedure using 100 μ l of BSA. The Pierce 660 nm Protein Assay has a greater linear range of 25-2,000 μ g/ml, compared with the Bradford Assay, which has a linear range of only 125-1,000 μ g/ml.

References

- Bradford, M.M. (1976). A rapid and sensitive method for the quantitation of microgram quantities of protein utilizing the principle of protein-dye binding. *Anal. Biochem.* 72 248-254
- Gornall, A.G. (1949). Determination of serum proteins by means of the biuret reaction. J. Biol. Chem. 177, 751-766.
- Lowry, O.H. (1951). Protein measurement with Folin-Phenol reagent. J. Biol. Chem. 193, 265-275.
- Smith, P.K., et al. (1985). Measurement of protein using bicinchoninic acid. Anal. Biochem. 150, 76-85.
- 5. Stoscheck, C.M. (1987). Protein assay sensitive at nanogram levels. *Anal. Biochem.* **160**, 301-305.

Ordering Information

| Product | Description | Pkg. Size |
|---------|--|-----------|
| 22660 | Pierce 660 nm Protein Assay Reagent Sufficient reagent for 500 standard assays and 5,000 microplate assays. | 750 ml |
| 22662 | Pierce 660 nm Protein Assay Kit Sufficient reagent to perform 300 standard assays and 3,000 microplate assays. | Kit |
| | Contains: Pierce 660 nm Protein Assay Reagent Pre-Diluted Protein Assay Standards, Bovine Serum Albumin (BSA) Set 3.5 ml each of 125- 2,000 mg/ml BSA | 450 ml |
| 22663 | Ionic Detergent Compatibility Reagent Sufficient for treating 100 ml Pierce 660 nm Protein Assay Reagent. | 5 x 1 g |
| | 5 pouches, 1 gram each | |

Bicinchoninic Acid (BCA)-based Protein Assays

In 1985, Paul K. Smith, *et al.* introduced the BCA Protein Assay. Since then it has become the most popular method for colorimetric detection and quantitation of total protein. The BCA Protein Assay has a unique advantage over the Modified Lowry Protein Assay and any of the Coomassie dyebased assays – it is compatible with samples that contain up to 5% surfactants (detergents).

Briefly, the sample is added to the tube or plate containing the prepared BCA Working Reagent and after a 30-minute incubation at 37°C and cooling to room temperature, the resultant purple color is measured at 562 nm. The protocol is similar for the Micro BCA Protein Assay, except the ratio of sample volume to working reagent is different and the tubes are incubated for 60 minutes at 60°C.

Chemistry of BCA-based Protein Assays

The BCA Protein Assay combines the well-known reduction of Cu²⁺ to Cu¹⁺ by protein in an alkaline medium with the highly sensitive and selective colorimetric detection of the cuprous cation (Cu¹⁺) by bicinchoninic acid (Figure 1). The first step is the chelation of copper with protein in an alkaline environment to form a blue colored complex. In this reaction, known as the biuret reaction, peptides containing three or more amino acid residues form a colored chelate complex with cupric ions in an alkaline environment containing sodium potassium tartrate. This became known as the biuret reaction because a similar complex forms with the organic compound biuret (NH₂-CO-NH-CO-NH₂) and the cupric ion. Biuret, a product of excess urea and heat, reacts with copper to form a light blue tetradentate complex (Figure 2). Single amino acids and dipeptides do not give the biuret reaction, but tripeptides and larger polypeptides or proteins will react to produce the light blue to violet complex that absorbs light at 540 nm. One cupric ion forms a colored coordination complex with four to six nearby peptides bonds.

The intensity of the color produced is proportional to the number of peptide bonds participating in the reaction. Thus, the biuret reaction is the basis for a simple and rapid colorimetric reagent of the same name for quantitatively determining total protein concentration. Since the working range for the biuret assay is from 5 to 160 mg/ml, the biuret assay is used in clinical laboratories for the quantitation of total protein in serum.

Step 1.

Protein +
$$Cu^{2+}$$
 \longrightarrow Cu^{2+}

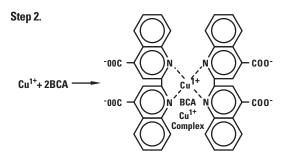


Figure 1. Reaction schematic for the bicinchoninic acid (BCA)-containing protein assay.

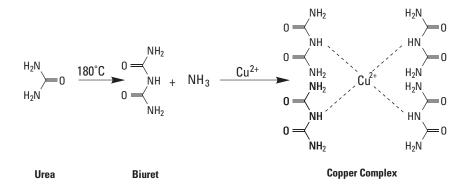


Figure 2. Biuret reaction schematic.

In the second step of the color development reaction, BCA Reagent, a highly sensitive and selective colorimetric detection reagent reacts with the cuprous cation (Cu¹⁺) that was formed in step 1. The purple colored reaction product is formed by the chelation of two molecules of BCA Reagent with one cuprous ion (Figure 1). The BCA/Copper Complex is water-soluble and exhibits a strong linear absorbance at 562 nm with increasing protein concentrations. The purple color may be measured at any wavelength between 550-570 nm with minimal (less than 10%) loss of signal. The BCA Reagent is approximately 100 times more sensitive (lower limit of detection) than the biuret reagent. The reaction that leads to BCA Color Formation as a result of the reduction of Cu²⁺ is also strongly influenced by the presence of any of four amino acid residues (cysteine or cystine, tyrosine, and tryptophan) in the amino acid sequence of the protein. Unlike the Coomassie dye-binding methods that require a minimum mass of protein to be present for the dye to bind, the presence of only a single amino acid residue in the sample may result in the formation of a colored BCA-Cu¹⁺ Chelate. This is true for any of the four amino acids cited above. Studies performed with di and tripeptides indicate that the total amount of color produced is greater than can be accounted for by the color produced with each BCA Reagent-reactive amino acid. Therefore, the peptide backbone must contribute to the reduction of copper as well.

The rate of BCA Color Formation is dependent on the incubation temperature, the types of protein present in the sample and the relative amounts of reactive amino acids contained in the proteins. The recommended protocols do not result in end-point determinations, the incubation periods were chosen to yield maximal color response in a reasonable time frame.

Advantages of the BCA Protein Assay

The primary advantage of the BCA Protein Assay is that most surfactants (even if present in the sample at concentrations up to 5%) are compatible. The protein:protein variation in the amount of color produced with the BCA Protein Assay is relatively low, similar to that observed for the Modified Lowry Protein Assay (Table 2, page 9).

The BCA Protein Assay produces a linear response curve $(r^2 > 0.95)$ and is available in two formulations based upon the dynamic range needed to detect the protein concentration of an unknown sample. The BCA Assay is less complicated to perform than the Lowry Protein Assay for both formulations. The standard BCA Protein Assay (Figure 3) detects protein concentrations from 20 to 2,000 µg/ml and is provided with Reagent A (carbonate buffer containing BCA Reagent) and Reagent B (cupric sulfate solution). A working solution (WS) is prepared by mixing 50 parts of BCA Reagent A with 1 part of BCA Reagent B (50:1, Reagent A:B). The working solution is an apple green color that turns purple after 30 minutes at 37°C in the presence of protein. The ratio of sample to WS used is 1:20. The Micro BCA Protein Assay (Figure 4) is more sensitive and has a narrower dynamic range of 0.1-25 µg/ml. To prepare the Micro BCA WS, three reagents (A, B and C) are mixed together at a ratio of 25 parts Micro Reagent A to 24 parts Micro Reagent B and 1 part Micro Reagent C. The Micro BCA WS is mixed with the sample or standard at a 1:1 volume ratio. The purple color response is read at 562 nm after 1 hour at 60°C.

Since the color reaction is not a true end-point reaction, considerable protocol flexibility is allowed with the BCA Protein Assay. By increasing the incubation temperature, the sensitivity of the assay can be increased. When using the enhanced tube protocol (incubating at 60°C for 30 minutes), the working range for the assay shifts to 5-250 μ g/ml and the minimum detection level becomes 5 μ g/ml.

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Both BCA Protein Assay formulations have less protein:protein variability than the Coomassie-based assays. The color response obtained for a seven point standard curve with the standard BCA Protein Assay using BSA or BGG standards shows less than a 20% variation between these two proteins (Figure 3). The Coomassie assay demonstrates > 30% variation in the signal generated between BSA and BGG (Table 2, page 9). There is even less variation (< 12%) when comparing these protein standards with the Micro BCA Protein Assay (Figure 4). In general, the BCA Protein Assay provides one of the most accurate measurements of protein concentration in biological samples, is detergentcompatible and simple to perform.

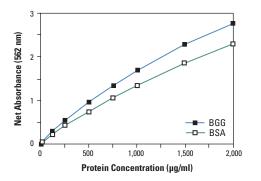


Figure 3. Color response curves obtained with the Thermo Scientific BCA Protein Assay using bovine serum albumin (BSA) and bovine gamma globulin (BGG). The standard tube protocol was performed and the color was measured at 562 nm.

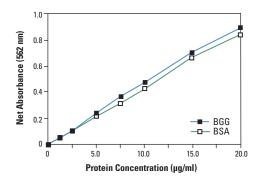


Figure 4. Color response curves obtained with the Thermo Scientific Micro BCA Protein Assay using bovine serum albumin (BSA) and bovine gamma globulin (BGG). The standard tube protocol was performed and the color was measured at 562 nm.

Disadvantages of the BCA Protein Assay

Substances that reduce copper will also produce color in the BCA Assay, thus interfering with the accuracy of the protein quantitation. Reagents that chelate the copper also interfere by reducing the amount of BCA Color produced with protein. Certain single amino acids (cysteine or cystine, tyrosine and tryptophan) will also produce color and interfere in BCA Assays.

BCA Protein Assay – Reducing Agent Compatible

The Thermo Scientific Pierce BCA Assay is always compatible with more detergents, buffers/salts and solvents than any other colorimetric protein assay. Now it's compatible with reducing agents at concentrations routinely used in protein sample buffers!



The BCA Assay provides one of the most accurate measurements of protein concentration in biological samples available. Although the BCA Assay is compatible with more detergents, buffers/salts and solvents than any colorimetric protein assay, the presence of disulfide reducing agents, including dithiothreitol (DTT) and 2-mercaptoethanol interferes with the assay. The BCA Protein Assay Kit – Reducing Agent Compatible (Product # 23250) provides all the advantages of the original BCA Assay as well as compatibility with reducing agents at concentrations routinely used in protein sample buffers (Figures 1 and 2).

Highlights:

- Compatible with up to 5 mM DTT, 35 mM 2-mercaptoethanol or 10 mM TCEP
- No protein precipitation required
- Linear working range: 125-2,000 μg/ml
- Sample volume: 25 µl
- Compatible with most ionic and nonionic detergents
- Significantly less protein:protein variation than coomassie (Bradford)-based methods
- Colorimetric method; measure at 562 nm
- Easy-to-use protocol (Figure 2)

References

Smith, P.K., et al. (1985). Measurement of protein using bicinchoninic acid. Anal. Biochem. 150, 76-85.

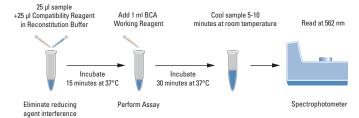


Figure 2. Thermo Scientific Pierce BCA Protein Assay – Reducing Agent Compatible protocol.

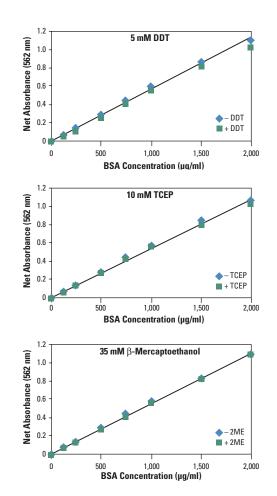


Figure 1. Thermo Scientific Pierce BCA Protein Assay – Reducing Agent Compatible produces a linear standard curve in the presence of reducing agents. Color response curves for BSA after treatment with Reducing Agent Compatible Reagent in the presence and absence of 5 mM DTT, 35 mM β -mercaptoethanol and 10 mM TCEP.

Ordering Information

| Product | Description | Pkg. Size |
|---------|--|--|
| 23250 | BCA Protein Assay Kit – Reducing Agent Compatible Sufficient reagents to perform 250 standard tube assays. | 750 ml |
| | Includes: BCA Reagent A BCA Reagent B Compatibility Reagent Reconstitution Buffer Albumin Standard (2 mg/ml) | 250 ml 25 ml 10 x 20 mg 15 ml 10 x 1 ml ampules |
| 23252 | Microplate BCA Protein Assay Kit – Reducing Agent Compatible Sufficient reagents for 1,000 microplate assays. | Kit |
| | Includes: BCA Reagent A BCA Reagent B Compatibility Reagent Reconstitution Buffer Albumin Standard (2 mg/ml) | 250 ml 25 ml 48 x 10 mg 15 ml 10 x 1 ml ampules |
| | 96-Well Microplates | 20/pkg. |

The Original BCA Protein Assay

Used in more labs than any other detergentcompatible formulation.



Highlights:

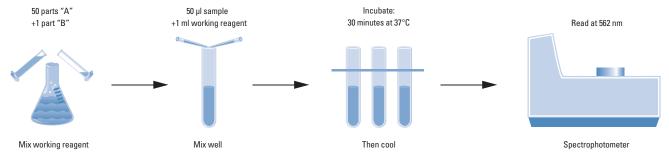
- · Colorimetric method; read at 562 nm
- Compatible with most ionic and nonionic detergents
- Four times faster and easier than the classical Lowry method
- · All reagents stable at room temperature for two years
- Working reagent stable for 24 hours
- Linear working range for BSA from 20 to 2,000 µg/ml
- Minimum detection level of 5 µg/ml with the enhanced protocol
- Convenient microplate or cuvette format
- Less protein:protein variation than dye-binding methods

| Irdoring | Information |
|----------|-------------|
| Ulucillu | mumanun |
| | |

| Product | Description | Pkg. Size |
|---------|---|-------------------|
| 23225 | BCA Protein Assay Kit Sufficient reagents to perform 500 standard tube assays or 5,000 microplate assays. | Kit |
| | Includes: Reagent A | 2 x 50 mg |
| | Reagent B | 25 ml |
| | Albumin Standard (2 mg/ml) | 10 x 1 ml ampules |
| 23227 | BCA Protein Assay Kit Sufficient reagents to perform 250 standard tube assays or 2,500 microplate assays. | Kit |
| | Includes: Reagent A | 1 x 500 ml |
| | Reagent B | 25 ml |
| | Albumin Standard (2 mg/ml) | 10 x 1 ml ampules |
| 23221 | BCA Protein Assay Reagent A Contains: BCA and tartrate in an alkaline carbonate buffer | 250 ml |
| 23223 | BCA Protein Assay Reagent A Contains: BCA and tartrate in an alkaline carbonate buffer | 1,000 ml |
| 23222 | BCA Protein Assay Reagent A Contains: BCA and tartrate in an alkaline carbonate buffer | 3.75 liter |
| 23224 | BCA Protein Assay Reagent B Contains: 4% CuSO ₄ •5H ₂ O | 25 ml |
| 23230 | BCA Protein Assay Reagent A Recrystallized purified powder | 25 g |
| 23228 | BCA Protein Assay Reagent A Contains: BCA and tartrate in an alkaline carbonate buffer | 500 ml |

References

Smith, P.K., et al. (1985). Anal. Biochem. 150, 76-85.
Sorensen, K. (1992). BioTechniques 12(2), 235-236.
Ju, T., et al. (2002). J. Biol. Chem. 277, 178-186.
Shibuya, T., et al. (1989). J. Tokyo Mid. College 47(4), 677-682.
Hinson, D.L. and Webber, R.J. (1988). BioTechniques 6(1), 14, 16, 19.
Akins, R.E. and Tuan, R.S. (1992). BioTechniques 12(4), 469-499.
Tyllianakis, P.E., et al. (1994). Anal. Biochem. 219(2), 335-340.
Gates, R.E. (1991). Anal. Biochem. 196(2), 290-295.
Stich, T.M. (1990). Anal. Biochem. 191, 343-346.
Tuszynski, G.P. and Murphy, A. (1990). Anal. Biochem. 184(1), 189-191.



Thermo Scientific Pierce BCA Protein Assay protocol.

Micro BCA Protein Assay

Most sensitive BCA formulation measuring dilute protein solutions from 0.5 to 20 µg/ml.

Highlights:

- Colorimetric method; read at 562 nm
- Compatible with most ionic and nonionic detergents
- A very sensitive reagent for dilute protein samples
- Linear working range for BSA: 0.5-20 µg/ml
- Less protein:protein variation than dye-binding methods
- All kit reagents stable at room temperature for two years
- Working reagent is stable for 24 hours
- Convenient microplate or cuvette format

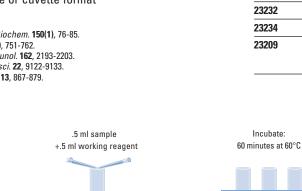
References

Smith, P.K., et al. (1985). Anal. Biochem. 150(1), 76-85.
 Kang, D.E., et al. (2002). Cell 110, 751-762.
 Rawadi, G., et al. (1999). J. Immunol. 162, 2193-2203.
 Blum, D., et al. (2002). J. Neurosci. 22, 9122-9133.
 Paratcha, G., et al. (2003). Cell 113, 867-879.

50 parts "MA" 48 parts "MB"

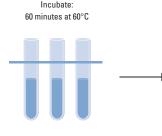
2 parts "MC"

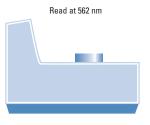
Mix working reagent



Ordering Information

| Description | Pkg. Size |
|---|--|
| Micro BCA Protein Assay Kit Sufficient reagents to perform 480 standard tube assays or 3,200 microplate assays. | Kit |
| Includes: Micro Reagent A (MA) (Sodium carbonate, sodium bicarbonate, and sodium tartrate in 0.2 N NaOH) | 240 ml |
| Micro Reagent B (MB) (4% BCA in water) | 240 ml |
| Micro Reagent C (MC) (4% cupric sulfate pentahydrate in water) Albumin Standard Ampules (2 mg/ml) | 12 ml |
| Micro BCA Reagent A (MA) | 240 ml |
| Micro BCA Reagent B (MB) | 240 ml |
| Micro BCA Reagent C (MC) | 12 ml |
| Albumin Standard Ampules, 2 mg/ml Contains: Bovine Albumin Fraction V in 0.9% NaCl solution containing sodium azide | Kit |
| | Micro BCA Protein Assay Kit Sufficient reagents to perform 480 standard tube assays or 3,200 microplate assays. Includes: Micro Reagent A (MA) (Sodium carbonate, sodium bicarbonate, and sodium tartrate in 0.2 N NaOH) Micro Reagent B (MB) (4% BCA in water) Micro Reagent C (MC) (4% cupric sulfate pentahydrate in water) Albumin Standard Ampules (2 mg/ml) Micro BCA Reagent B (MB) Micro BCA Reagent C (MC) Albumin Standard Ampules, 2 mg/ml Contains: Bovine Albumin Fraction V in 0.9% |





Then cool

Spectrophotometer

Thermo Scientific Pierce Micro BCA Protein Assay protocol.

Mix well

Coomassie Dye-based Protein Assays (Bradford Assays)

Use of Coomassie G-250 Dye in a colorimetric reagent for the detection and quantitation of total protein was first described by Dr. Marion Bradford in 1976. Both the Coomassie (Bradford) Protein Assay Kit (Product # 23200) and the Coomassie Plus (Bradford) Assay Kit (Product # 23236) are modifications of the reagent first reported by Dr. Bradford.

Chemistry of Coomassie-based Protein Assays

In the acidic environment of the reagent, protein binds to the Coomassie dye. This results in a spectral shift from the reddish/ brown form of the dye (absorbance maximum at 465 nm) to the blue form of the dye (absorbance maximum at 610 nm) (Figure 1). The difference between the two forms of the dye is greatest at 595 nm, so that is the optimal wavelength to measure the blue color from the Coomassie dye-protein complex. If desired, the blue color can be measured at any wavelength between 575 nm and 615 nm. At the two extremes (575 nm and 615 nm) there is a loss of about 10% in the measured amount of color (absorbance) compared to that obtained at 595 nm.

Development of color in Coomassie dye-based protein assays has been associated with the presence of certain basic amino acids (primarily arginine, lysine and histidine) in the protein. Van der Waals forces and hydrophobic interactions also participate in the binding of the dye by protein. The number of Coomassie dye ligands bound to each protein molecule is approximately proportional to the number of positive charges found on the protein. Free amino acids, peptides and low molecular weight proteins do not produce color with Coomassie dye reagents. In general, the mass of a peptide or protein must be at least 3,000 daltons to be assayed with this reagent. In some applications this can be an advantage. The Coomassie (Bradford) Protein Assay has been used to measure "high molecular weight proteins" during fermentation in the beer brewing industry.

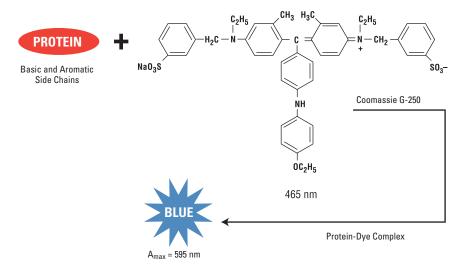


Figure 1. Reaction schematic for the Coomassie dye-based protein assays (the Coomassie [Bradford] Protein Assay and the Coomassie Plus (Bradford) Assay.

Advantages of Coomassie-based Protein Assays

Coomassie dye-binding assays are the fastest and easiest to perform of all protein assays. The assay is performed at room temperature and no special equipment is required. Briefly, for either the Coomassie (Bradford) Protein Assay or the Coomassie Plus Assay, the sample is added to the tube containing reagent and the resultant blue color is measured at 595 nm following a short room-temperature incubation. The Coomassie dye-containing protein assays are compatible with most salts, solvents, buffers, thiols, reducing substances and metal chelating agents encountered in protein samples.

Disadvantages of Coomassie-based Protein Assays

The main disadvantage of Coomassie-based protein assays is their incompatibility with surfactants at concentrations routinely used to solubilize membrane proteins. In general, the presence of a surfactant in the sample, even at low concentrations, causes precipitation of the reagent. Since the Coomassie dye reagent is highly acidic, a small number of proteins cannot be assayed with this reagent due to their poor solubility in the acidic reagent. Also, Coomassie reagents result in about twice as much protein:protein variation as copper chelation based assay reagents (Table 2, page 9). In addition, Coomassie dye stains the glass or quartz cuvettes used to hold the solution in the spectrophotometer while the color intensity is being measured. (Cuvettes can be cleaned with strong detergent solutions and/or methanol washes, but use of disposable polystyrene cuvettes eliminates the need to clean cuvettes.)

General Characteristics of Coomassie-based Protein Assays (Bradford Assays)

Coomassie-based protein assays share a number of characteristics. The Coomassie (Bradford) Protein Assay produces a nonlinear standard curve. The Coomassie Plus (Bradford) Assay has the unique advantage of producing a linear standard curve over part of its total working range. When using bovine serum albumin (BSA) as the standard, the Coomassie Plus Assay is linear from 125 to 1,000 μ g/ml. When using bovine gamma globulin (BGG) as the standard, the Coomassie Plus Assay is linear from 125 to 1,500 μ g/ml. The complete working range of the Coomassie Plus Assay covers the concentration range from 125 to 1,000 μ g/ml for the tube protocol and from 1 to 25 μ g/ml for the micro protocol (Figures 2-3).

Coomassie dye-based protein assays must be refrigerated for long-term storage. If ready-to-use liquid Coomassie dye reagents will be used within one month, they may be stored at ambient temperature (18-26°C). Coomassie protein assay reagent that has been left at room temperature for several months will

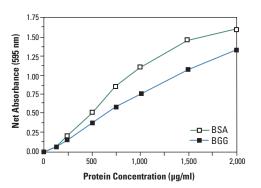


Figure 2. Color response curves obtained with Thermo Scientific Pierce Coomassie Plus (Bradford) Assay using bovine serum albumin (BSA) and bovine gamma globulin (BGG). The standard tube protocol was performed and the color was measured at 595 nm.

have a lower color response, especially at the high end of the working range. Coomassie protein assay reagents that have been stored refrigerated must be warmed to room temperature before use. Using either cold plates or cold liquid Coomassie dye reagent will result in low absorbance values.

The ready-to-use liquid Coomassie dye reagents must be mixed gently by inversion just before use. The dye in these liquid reagents spontaneously forms loose aggregates upon standing. These aggregates may become visible after the reagent has been standing for as little as 60 minutes. Gentle mixing of the reagent by inversion of the bottle will uniformly disperse the dye. After binding to protein, the dye also forms protein-dye aggregates. Fortunately, these protein-dye aggregates can be dispersed easily by mixing the reaction tube. This is common to all liquid Coomassie dye reagents. Since these aggregates form relatively quickly, it is also best to routinely mix (vortex for 2-3 seconds) each tube or plate just before measuring the color.

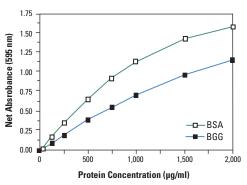


Figure 3. Color response curves obtained with Thermo Scientific Pierce Coomassie (Bradford) Protein Assay using bovine serum albumin (BSA) and bovine gamma globulin (BGG). The standard tube protocol was performed and the color was measured at 595 nm.

Coomassie Plus (Bradford) Protein Assay

As fast as the original Coomassie Assay, with increased accuracy ... the high-performance Bradford reagent.

• Easier, quicker preparation

Working reagent is ready to use. No tedious dilution, no filtration of a dye concentrate and no mess to clean up.

Lower cost per assay

Just 23¢ per sample with the standard protocol, and less than 5¢ per sample with the microplate protocol.

• Faster assay

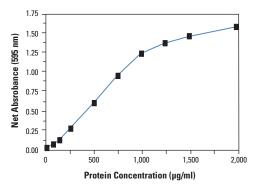
Total assay time is less than 10 minutes!

• More accurate results

Substantially increased linearity of response, and only half the expected protein:protein variation of other commercial formulations.

Highlights:

- Detects protein concentrations from 1 to 1,500 µg/ml
- Ready-to-use dye-binding reagent formulation
- Fast (almost immediate) color development read at 595 nm
- Compatible with reducing sugars, reducing substances and thiols
- · Refrigerated reagent is stable for up to two years
- Superior linear response over the range of 125-1,500 µg/ml
- · Convenient microplate or cuvette format
- \bullet Micro protocol useful for protein concentrations from 1 to 25 $\mu g/ml$



Typical color response curve for BSA using the Thermo Scientific Pierce Coomassie Plus (Bradford) Protein Assay Reagent.

References

Bradford, M. (1976). *Anal. Biochem.* **72**, 248-254. Glover, B.P. and McHenry, C.S. (2001). *Cell* **105**, 925-934. Kagan, A., *et al.* (2000). *J. Biol. Chem.* **275**, 11241-11248. Goel, R., *et al.* (2002). *J. Biol. Chem.* **277**, 18640-18648.

Ordering Information

| Product | Description | Pkg. Size |
|-----------|---|---------------------|
| 23236 | Coomassie Plus (Bradford) Assay Kit Sufficient reagents to perform 630 standard assays or 3,160 microplate assays. | Kit |
| | Includes: Coomassie Plus Protein Assay Reagent Albumin Standard (2 mg/ml) | 950 ml 10 x 1 ml |
| | | ampules |
| 23238 | Coomassie Plus (Bradford) Reagent Sufficient reagents to perform 200 standard assays or 1,000 microplate assays. Albumin Standard not included. | 300 ml |
| Related P | roducts | |
| Product | Description | Pkg. Size |
| 23239 | Coomassie Plus Compat-Able Protein Assay Kit | Kit |



Thermo Scientific Pierce Coomassie Plus (Bradford) Assay protocol. The protocol is simple, fast and very easy to perform.

Compatible Substances

Reagents compatible with Coomassie Plus Assay using the standard protocol. Interferences may be observed at the stated concentration when using the Micro Assay Procedure.

| Ammonium Sulfate | 1.0 M | 2-Mercaptoethanol | 1.0 M |
|------------------|--------|-------------------|--------|
| Azide | 0.5% | MES | 100 mM |
| Brij-56 | 0.03% | NaCl | 5.0 M |
| Brij-35 | 0.06% | NaOH | 0.1 M |
| Brij-58 | 0.016% | NP-40 | 0.5% |
| CHAPS | 5.0% | SDS | 0.016% |
| CHAPSO | 5.0% | Sucrose | 10.0% |
| Citrate | 200 mM | Tris | 2.0 M |
| EDTA | 100 mM | Triton X-100 | 0.06% |
| Glucose | 1.0 M | Triton X-114 | 0.06% |
| Glycine | 0.1 M | Triton X-405 | 0.25% |
| Guanidine•HCl | 3.5 M | Tween-20 | 0.03% |
| HCI | 0.1 M | Tween-80 | 0.016% |
| KSCN | 3.0 M | Urea | 3.0 M |
| | | | |

Coomassie Dye-based Protein Assays

Coomassie (Bradford) Protein Assay

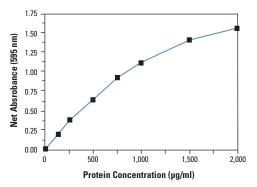
The Bradford method workhorse ... ready-to-use, allowing total protein determination in seconds!

This ready-to-use formulation more closely resembles in performance, the reagent published by Bradford.¹ It demonstrates the typical assay characteristics known for Coomassie dye-based formulations.²

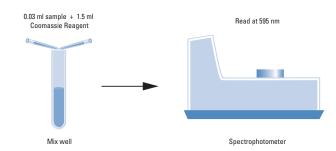
Highlights:

24

- · Ready-to-use dye-binding reagent formulation
- Fast (almost immediate) color development; read at 595 nm
- Compatible with reducing substances and chelating agents
- Refrigerated reagent is stable for 12 months
- Determine protein concentration from 100 to 1,500 µg/ml
- Micro method for the range of 1 to 25 µg/ml
- Convenient microplate or cuvette format



Thermo Scientific Pierce Coomassie (Bradford) Protein Assay Reagent: typical color response curve for BSA.



Thermo Scientific Pierce Coomassie (Bradford) Protein Assay protocol.

Ordering Information

| Product | Description | Pkg. Size |
|---------|--|-----------|
| 23200 | Coomassie (Bradford) Protein Assay Kit (Ready-to-use Coomassie Blue G-250 based reagent) Sufficient reagents to perform 630 standard tube assays or 3,800 microplate assays. | Kit |
| | Includes: Coomassie Protein Assay Reagent | 950 ml |
| | Albumin Standard Ampules (2 mg/ml) | 10 x 1 ml |

References

- 1. Bradford, M. (1976). Anal. Biochem. 72, 248-254.
- 2. VanKley, H. and Hale, S.M. (1977). Anal. Biochem. 81, 485-487.
- Messenger, M.M., et al. (2002). J. Biol Chem. 277, 23054-23064.

Removing Interfering Substances

Virtually every protein detection method known exhibits sensitivity to the presence of particular reagents in the protein sample. Proteins are typically found in solutions that contain detergents, buffer salts, denaturants, reducing agents, chaotropic agents and/or anti-microbial preservatives. These additives may affect the results of an assay. When a component of a protein solution artificially increases or decreases the signal of any assay, the component is considered to be an interfering substance.

Interfering substances can affect the protein assay in the following ways:

- They can suppress the response of an assay
- They can enhance the response of an assay
- . They can result in an elevated background reading

A small amount of interference from many common substances can be compensated for in the blank designed for a specific assay. To compensate for the interference, the protein samples for the standard curve must be diluted in the same buffer as the protein being assayed.

Often, interfering substances can overwhelm the assay, making it difficult or impossible to perform. The two most popular assay methods, Lowry- or Bradford-based assays, are both strongly affected by various components found in standard sample buffers. Lowry-based methods are incompatible with reducing and chelating agents; DTT, β -mercaptoethanol, cysteine, EDTA and some sugars while Bradford-based methods are incompatible with most detergents. Unfortunately, many common sample buffers contain both reducing agents and detergents, Laemmli buffer for example.

In these situations, the interfering substance can be removed by a variety of means, of which gel filtration and dialysis are the most common. However both of these methods are time-consuming and can result in diluted protein samples. The Compat-Able Protein Assay Preparation Set (page 26) was developed to solve this problem. The Compat-Able Reagents render potentially interfering substances virtually invisible to either a Lowry- or Bradford-based assay. These unique reagents dispose of any possible interfering substances in your sample by selectively precipitating out the protein, allowing the non-protein sample components to be removed easily. Precipitated protein is recovered in water or an assay-compatible buffer and then assayed by any method.

In one round of treatment, Compat-Able Reagents can remove most any interfering substance, including but not limited to:

- Laemmli buffer
- 3.0 M Tris
- 20% glycerol
- 4% SDS
- 3.6 M magnesium chloride
- 1.25 M sodium chloride
- 350 mM dithiothreitol (DTT)
- 5% Triton X-100
- 5% Tween-20
- 125 mM sodium citrate
- 200 mM glucose
- 200 mM sodium acetate
- 5% β -mercaptoethanol
- 200 mM EDTA
- 1.0 M imidazole

If concentrations of these or other interfering components exceed this level, more than one round of pre-treatment can be performed.

Removal of Interfering Substances

Compat-Able Protein Assays

Excellent choice for use with samples prepared for 1-D or 2-D electrophoresis.

These Thermo Scientific Kits pair BCA and Coomassie Plus (Bradford) Assays, recognized around the world as the best detergent- and reducing agent-compatible assays (respectively) for total protein analysis, with a great sample preparation reagent. These unique reagents dispose of any interfering substances in your sample by selectively precipitating the protein, allowing the nonprotein components to be removed easily. Precipitated protein is recovered in water and assayed with the BCA Protein Assay or Coomassie Plus Assay.

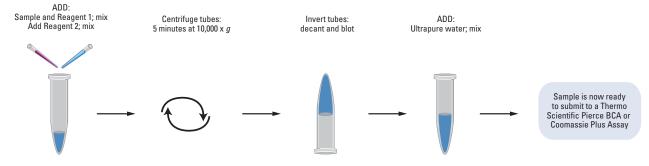
Highlights:

- Removes interfering substances prior to any downstream application
- Ready-to-use sample preparation reagents save time and effort
- Four-step protocol takes less than 10 minutes to complete
- Room temperature-stable sample preparation reagents can be stored on your bench top so they won't get lost in the cold room or hidden in the lab refrigerator
- Precipitates protein out of solution, leaving potentially interfering substances to be decanted away without dialysis or gel filtration, saving time and avoiding sample loss or dilution
- Easily adaptable to pre-treatment of many samples at one time
- Adaptable to both a test tube and microcentrifuge tube sample preparation protocol, to allow for 50 μl or 100 μl sample volumes
- Sample prep reagents are available with the BCA or Coomassie Assays or sold separately



Ordering Information

| Product | Description | Pkg. Size |
|---------|--|------------|
| 23229 | BCA Compat-Able Protein Assay Kit Contains one each of the following: Product # 23227, BCA Protein Assay Kit Sufficient reagents to perform 250 standard tube assays or 2,500 microplate assays. | Kit |
| | BCA Reagent A | 2 x 250 ml |
| | BCA Reagent B | 25 ml |
| | BSA Standards (2 mg/ml) Product # 23215, Compat-Able | 10 x 1 ml |
| | Protein Assay Preparation Reagent Set | |
| | (see description below) | |
| 23239 | Coomassie Plus Compat-Able Protein Assay Reagent Kit | Kit |
| | Contains one each of the following: Product # 23236, Coomassie Plus Protein Assay | |
| | Reagent Kit Sufficient materials for 630 standard assays, | |
| | <i>950 microassays or 3,160 microplate assays.</i> Coomassie Plus Reagent Formulation | 950 ml |
| | BSA Standards (2 mg/ml) | 10 x 1 ml |
| | Product # 23215, Compat-Able | |
| | Protein Assay Preparation Reagent Set (see description below) | |
| 23215 | Compat-Able Protein Assay Preparation Reagent Set | Kit |
| | Two-reagent set with sufficient material to pre-treat up to 500 samples prior to total | |
| | protein assay. | |
| | Compat-Able Protein Assay Preparation Reagent 1 | 250 ml |
| | Compat-Able Protein Assay Preparation Reagent 2 | 250 ml |



Thermo Scientific Compat-Able Protein Assay protocol. Make almost any protein sample compatible with the Thermo Scientific Pierce BCA or Coomassie Plus (Bradford) Assays in four simple steps.

Modified Lowry Protein Assay

Although the mechanism of color formation for the Modified Lowry Protein Assay is similar to that of the BCA Protein Assay, there are several significant differences between the two.

In 1951 Oliver H. Lowry introduced this colorimetric total protein assay method. It offered a significant improvement over previous protein assays and his paper became one of the most cited references in the life science literature. The Modified Lowry Protein Assay uses a stable reagent that replaces two unstable reagents described by Dr. Lowry. The Modified Lowry assay is easy to perform because the incubations are done at room temperature and the assay is sensitive enough to allow the detection of total protein in the low microgram per milliliter range. Essentially, the Modified Lowry protein assay is an enhanced biuret assay involving copper chelation chemistry.

Chemistry of the Modified Lowry Protein Assay

Although the mechanism of color formation for the Modified Lowry Protein Assay is similar to that of the BCA Protein Assay, there are several significant differences between the two. The exact mechanism of color formation in the Modified Lowry Protein Assay remains poorly understood. It is known that the color-producing reaction with protein occurs in two distinct steps. As seen in Figure 1, protein is first reacted with alkaline cupric sulfate in the presence of tartrate during a 10-minute incubation at room temperature. During this incubation, a tetradentate copper complex forms from four peptide bonds and one atom of copper. The tetradentate copper complex is light blue in color (this is the "biuret reaction"). Following the incubation, Folin phenol reagent is added. It is believed that the color enhancement occurs when the tetradentate copper complex transfers electrons to the phosphomolybdic/phosphotungstic acid complex (the Folin phenol reagent).

The reduced phosphomolybdic/phosphotungstic acid complex produced by this reaction is intensely blue in color. The Folin phenol reagent loses its reactivity almost immediately upon addition to the alkaline working reagent/sample solution. The blue color continues to intensify during a 30-minute room temperature incubation. It has been suggested by Lowry, *et al.* and by Legler, *et al.* that during the 30-minute incubation, a rearrangement of the initial unstable blue complex leads to the stable final blue colored complex that has higher absorbance.

For small peptides, the amount of color increases with the size of the peptide. The presence of any of five amino acid residues (tyrosine, tryptophan, cysteine, histidine and asparagine) in the peptide or protein backbone further enhances the amount of color produced because they contribute additional reducing equivalents to further reduce the phosphomolybdic/phosphotungstic acid complex. With the exception of tyrosine and tryptophan, free amino acids will not produce a colored product with the Modified Lowry Reagent; however, most dipeptides can be detected. In the absence of any of the five amino acids listed above in the peptide backbone, proteins containing proline residues have a lower color response with the Modified Lowry Reagent due to the amino acid interfering with complex formation.

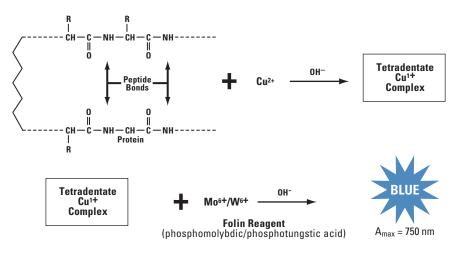


Figure 1. Reaction schematic for the Modified Lowry Protein Assay.

Advantages of the Modified Lowry Protein Assay

The final blue color is optimally measured at 750 nm, but it can be measured at any wavelength between 650 nm and 750 nm with little loss of color intensity. It is best to measure the color at 750 nm because few other substances absorb light at that wavelength. The amount of light absorbed at 750 nm is directly proportional to the amount of protein in the sample, but the color response curve produced is nonlinear. The sensitivity of the Modified Lowry Protein Assay is greatly enhanced over that of the biuret reagent. The working range of the method extends from 5 to 2,000 mg/ml.

The Modified Lowry Protein Assay demonstrates less protein:protein variability than Coomassie-based assays. When comparing the standard curve responses between BSA and BGG, there is less than a 15% variation in the signal generated with these two standard proteins (Figure 2). The Coomassie Protein Assay demonstrates > 30% variation in the signal generated between BSA and BGG (Table 2, page 9).

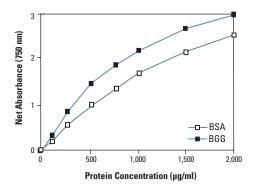


Figure 2. Color response curves obtained with the Thermo Scientific Pierce Modified Lowry Protein Assay Reagent using bovine serum albumin (BSA) and bovine gamma globulin (BGG). The standard tube protocol was performed and the color was measured at 750 nm.

Disadvantages of the Modified Lowry Protein Assay

The Modified Lowry Protein Assay will form precipitates in the presence of detergents or potassium ions. The problem of precipitation caused by the presence of potassium ions in the sample can sometimes be overcome by centrifuging the tube and measuring the color in the supernatant. Most surfactants will cause precipitation of the reagent even at very low concentrations. One exception is sodium dodecyl sulfate (SDS), which is compatible with the reagent at concentrations up to 1% in the sample. Chelating agents interfere by binding copper and preventing formation of the copper peptide bond complex. Reducing agents and free thiols also interfere by reducing the phosphotungstate-phosphomolybdate complex, immediately forming an intensely blue colored product upon their addition to the Modified Lowry Protein Assay Reagent.

General Characteristics of the Modified Lowry Protein Assay

The Modified Lowry Protein Assay Reagent must be refrigerated for long-term storage. If the entire bottle of reagent will be used within one month, it may be stored at room temperature (18-26°C). Reagent that has been left at room temperature for more than one month may produce lower color response, especially at the higher end of the working range. If the reagent has been stored refrigerated, it must be warmed to room temperature before use. Using cold Modified Lowry Protein Assay Reagent will result in low absorbance values.

The protocol requires that the Folin phenol reagent be added to each tube precisely at the end of the 10-minute incubation. At the alkaline pH of the Lowry reagent, the Folin phenol reagent is almost immediately inactivated. Therefore, it is best to add the Folin phenol reagent at the precise time while simultaneously mixing each tube. Because this is somewhat cumbersome, some practice is required to obtain consistent results. This also limits the total number of samples that can be assayed in a single run. If a 10-second interval between tubes is used, the maximum number of tubes that can be assayed within 10 minutes is 60 (10 seconds/tube x 60 tubes = 600 seconds or 10 minutes).

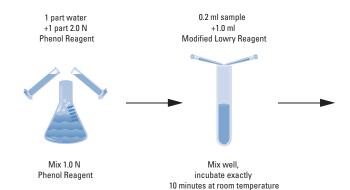
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Modified Lowry Protein Assay Reagent

All the accuracy of the Lowry, but modified so it's ready-to-use and stable for at least one year!

Highlights:

- The most widely cited colorimetric method; read at 750 nm
- Ready-to-use reagent for the loyal Lowry method user
- Preformulated cupric sulfate-tartrate reagent stable for one year at room temperature
- Linear results from 1 to 1,500 µg/ml for BSA
- Adaptable to microplates
- Less protein:protein variation than dye-binding methods



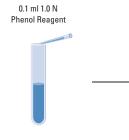
Thermo Scientific Pierce Modified Lowry Protein Assay Reagent protocol.

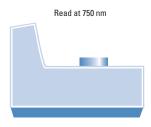
Ordering Information

| Product | Description | Pkg. Size |
|---------|--|------------------------------|
| 23240 | Modified Lowry Protein Assay Kit Sufficient reagents to perform 480 standard tube assays or 2,400 microplate assays. | Kit |
| | Includes: Modified Lowry Protein Assay Reagent 2 N Folin-Ciocalteu Phenol Reagent Albumin Standard Ampules (2 mg/ml) | 480 ml 50 ml 10 x 1 ml |

References

Lowry, O.H., et al. (1951). J. Biol. Chem. 193, 76-85. Temel, R.E., et al. (2003). J. Biol. Chem. 278, 4792-4799.





Mix well, incubate 30 minutes at room temperature

Spectrophotometer

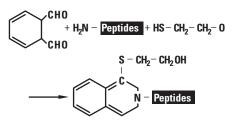
o-Phthalaldehyde [OPA] Fluorescent Protein Assay

The Thermo Scientific Pierce Fluoraldehyde Protein/Peptide Assay is an *o*-phthalaldehyde-based reagent developed to detect minute amounts of protein and peptides. Fluoraldehyde reactions are complete in less than one minute with sensitivity down to 50 ng/ml. While some solutions interfere with protein/peptide measurement at 280 nm, our Fluoraldehyde Assay is compatible with many substances that interfere with other protein assays, such as detergents and reducing agents. Amine-containing buffers must be avoided, however, when performing assays using this chemistry.

In the standard assay mode, the fluoraldehyde ready-to-use formulation can measure protein concentration in the range of 10 to 500 μ g/ml, while the micro-assay working range is 50 ng/ml to 25 μ g/ml.

Our Fluoraldehyde Protein/Peptide Assay Reagent requires only 200 µl of sample for use in a microplate assay, saving valuable sample and time. Fluoraldehyde assays require an excitation wavelength of 360 nm and emission wavelength of 455 nm.

OPA will react only with primary amines. When reacted with primary amines in the presence of mercaptoethanol, OPA yields an intense blue colored fluorescent product that has a maximum wavelength of excitation of 340 nm and emission at 455 nm.^{1,2} Wavelengths from 330-375 nm have been used for excitation and 436-490 nm for measuring emission. Protein concentrations



The reaction of *o*-Phthalaldehyde with a primary amine on a peptide in the presence of 2-mercaptoethanol to form a fluorescent-labeled peptide.

Fluoraldehyde o-Phthalaldehyde Crystals

An easy, economical way to detect amino acids in pre- and postcolumn chromatographic effluents.

Stable in aqueous solution

Rapid analysis, no heating required

· Highly sensitive, low

background

Highlights:

- 0 H O
- Fluoraldehyde o-Phthalaldehyde M.W. 134.13

as low as 50 ng/ml can be measured with an OPA assay. The inherent sensitivity and speed of OPA, along with its broad linear range, makes it a useful protein and peptide assay reagent.

OPA is ideal for assaying peptides that do not contain tyrosine residues, or for other applications in which absorbance at 280 nm cannot be used. Proteins and peptides tested yield linear results over a wide range of concentrations using both standard and microassay protocols.

There is considerable protein:protein and peptide:peptide variation with the OPA assay; therefore, it is best to use a purified sample of the particular protein or peptide as the standard. When this is not possible, the next best option is to use a protein or peptide that gives a response similar to the sample. Alternatively, a commonly accepted standard protein such as bovine serum albumin can be used.

Reducing agents and metal chelators do not interfere with an OPA-based assay, provided they are included in the blanks and standards. In addition, most detergents do not interfere. Any common sample buffers and constituents are also compatible, but primary amines such as Tris or glycine buffers will interfere with OPA and must be avoided. Acetylated and other primary amineblocked peptides will not give a response with OPA.

References

1. Ogden, G. and Foldi, P. (1987). *LC•GC* 5(1), 28-38. 2. Roth, M. (1971). *Anal. Chem.* 43, 880-882.

| Ordering Information | | |
|----------------------|--|-----------|
| Product | Description | Pkg. Size |
| 26015 | Fluoraldehyde <i>o</i> -Phthalaldehyde Crystals | 5 g |

References

Lindroth, P. and Mopper, K. (1979). Anal. Chem. 51, 1667-1674. Lee, K.S. and Drescher, D.G. (1979). J. Biol. Chem. 254, 6248-6251. van Eijk, H.M., et al. (1988). Clin. Chem. 34, 2510-2513. Graser, T.A., et al. (1988). Anal. Biochem. 151, 142-152. Cooper, J.D., et al. (1984). Anal. Biochem. 142, 98-102. Krishnamurti, C.R., et al. (1984). J. Chromatogr. 315, 321-331. Jones, B.N., et al. (1983). J. Chromatogr. 266, 471-482. Lee, H., et al. (1979). Anal. Biochem. 96, 298-307. Chen, R.F., et al. (1981). J. Lig. Chrom. 4, 565-586.

Fluoraldehyde *o*-Phthalaldehyde Reagent Solution

Excellent sensitivity – an ideal choice when working with limited amounts of purified protein or peptides.



Fluoraldehyde o-Phthalaldehyde Reagent Solution M.W. 134.13 $\lambda ex = 340 \text{ nm}$ $\lambda em = 455 \text{ nm}$

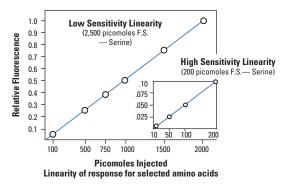
Thermo Scientific Pierce Fluoraldehyde Reagent Solution contains a stabilized, highly purified preparation of *o*-phthalaldehyde, Brij-35 Detergent and mercaptoethanol in a specially formulated borate buffer. It is a highly sensitive, ready-to-use reagent solution that exhibits excellent linear response (Figure 1) and offers outstanding shelf life (Figure 2). In addition, when compared to other *o*-phthalaldehyde detection reagents, our solution exhibits decreased background over time and a high signal:noise ratio.

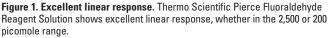
Highlights:

- A ready-to-use, highly sensitive fluorescent pre- or post-column reagent for amino acid detection and quantitation
- Provides an accurate measure of both composition and absolute protein/peptide content
- · Ready-to-use with no processing needed
- · Reacts with all primary amine-containing analytes
- · High sensitivity; low background

Application Note:

For even greater sensitivity, use a combination of OPA with Fmoc-Chloride with automated pre-column derivatization, detecting both primary and secondary amines. with this application, primary amino acids are first derivatized with OPA, while non-reacted secondary amino acids are then reacted with Fmoc-Chloride, resulting in extraordinary amino acid detection sensitivity and accuracy.¹²





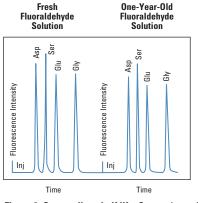


Figure 2. Outstanding shelf life. Comparison of fluorescence response of selected amino acids after reaction with recently prepared and one-year-old Thermo Scientific Pierce Fluoreldehyde Reagent Solutions.

Ordering Information

| Product | Description | Pkg. Size | |
|---------|--|-----------|--|
| 26025 | Fluoraldehyde <i>o</i> -Phthalaldehyde Reagent Solution | 945 ml | |

References

1. Godel, H., et al. (1992). LC-GC International 5, 44-49.

2. Schuster, R. (1988). J. Chromatogr. 431, 271-284.

Jones, B.N. and Gilligan, J.P. (1983). American Biotechnology Laboratory, Dec. Issue, 46-51.

Benson, J.R. and Woo, D.J. (1984). J. Chromatogr. Sci. 22, 386-399.

Specialty Assays – Histidine-tagged Proteins

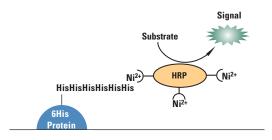
Histidine-tagged Protein Detection

Thermo Scientific HisProbe-HRP Western blotting probe takes advantage of the affinity of histidine for the Ni²⁺ cation.

HisProbe-HRP is a nickel (Ni²⁺)-activated derivative of horseradish peroxidase (HRP). This product has been optimized for direct detection of recombinant histidine-tagged proteins and other histidine-rich proteins. The active ligand is a tridentate chelator that allows Ni²⁺ to be bound in active form for subsequent interaction and detection of target molecules. The active chelator has similar binding capabilities to that reported for iminodiacetic acid, which has long been used for immobilized metal affinity chromatography (IMAC).

Highlights:

- Yields lower background than anti-histidine antibodies
- · Pierce HRP is a high-activity enzyme
- Stripping and reprobing is possible
- HisProbe-HRP (Ni²⁺) can be used for detection of histidine-tagged proteins

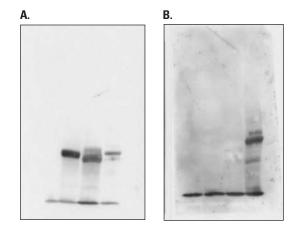


Detection of histidine-tagged fusion proteins with Thermo Scientific HisProbe-HRP.

Ordering Information

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| Product | Description | Pkg. Size |
|---------|---|----------------|
| 15165 | HisProbe HRP | 1 mg |
| 15168 | SuperSignal [®] West Pico HisProbe Kit | Kit |
| | Includes: HisProbe -HRP | 2 mg |
| | SuperSignal West Pico Chemiluminescent Substrate | 500 ml |
| | BSA in TBS (10X) | 1 x 125 ml |
| | BupH™ Tris Buffered Saline Packs | 10 x 500 ml |
| | Surfact-Amps [®] 20 (10%) | 6 x 10 ampules |



Panel A using Thermo Scientific HisProbe-HRP shows high specific binding and low background.

Panel B using anti-polyHis failed to recognize two of the three fusion proteins.

References

- Adler, J. and Bibi, E. (2004). Determinants of substrate recognition by the *Escherichia coli* multidrug transporter MdfA identified on both sides of the membrane. *J. Biol. Chem.* **279**, 8957-8965.
- Adler, J. and Bibi, E. (2005). Promiscuity in the geometry of electrostatic interactions between the *Escherichia coli* multidrug resistance transporter MdfA and cationic substrates. J. Biol. Chem. 280, 2721-2729.
- Boulant, S., et al. (2003). Unusual multiple recoding events leading to alternative forms of hepatitis C virus core protein from genotype 1b. J. Biol. Chem. 278, 45785-45792.
- Kanaya, E., et al. (2001). Zinc release from the CH₂C₆ zinc finger domain of filamentous flower protein from Arabidopsis thaliana induces self-assembly. J. Biol. Chem. 276, 7383-7390.
- Robalino, J., et al. (2004). Two zebrafish eIF4E family members are differentially expressed and functionally divergent. J. Biol. Chem. 279, 10532-10541.
- Robichon, C., et al. (2005). Depletion of apolipoprotein N-acyltransferase causes mislocalization of outer membrane lipoproteins in *Escherichia coli. J. Biol. Chem.* 280, 974-983.
- Segawa, H., et al. (2005). Reconstitution of GDP-mannose transport activity with purified Leishmania LPG2 protein in liposomes. J. Biol. Chem. 280, 2028-2035.
- Sundberg-Smith, L., et al. (2005). Adhesion stimulates direct PAK1/ERK2 association and leads to ERK-dependent PAK1 Thr212 phosphorylation. J. Biol. Chem. 280, 2055-2064.
- Wagner, C., et al. (2005). Dimerization of NO-sensitive guanylyl cyclase requires the $\alpha 1$ N terminus. J. Biol. Chem. 280, 17687-17693.
- Wann, E., et al. (2000). The fibronectin-binding MSCRAMM FnbpA of Staphylococcus aureus is a bifunctional protein that also binds to fibrinogen. J. Biol. Chem. 275, 13863-13871.

Easy-Titer IgG and IgM Assay Kits

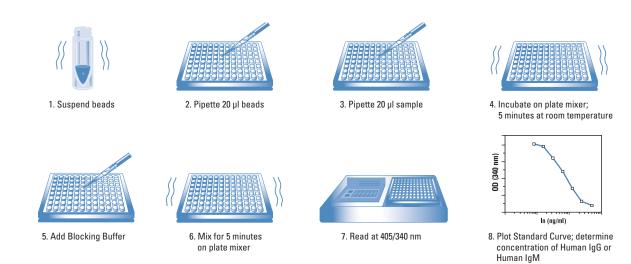
Simply the fastest, easiest way to quantitate antibodies ... ever!

It is no longer necessary to wait or to rely on inaccurate and insensitive UV or colorimetric IgG determination methods. It is not necessary to struggle with the inadequacies of methods that titrate antibody activity. It is even possible to avoid the tedious, time-consuming ELISA approach to determine antibody titer. Thermo Scientific Easy-Titer IgG Assay Kits make it possible to detect IgG in less time and with greater specificity and sensitivity than ever before.

Our Easy-Titer[®] Assay Kits do not cross-react with antibodies from other species such as bovine antibodies present in the media used to culture antibody-producing hybridoma cells. This remarkable specificity allows the measurement of human IgG concentrations from a variety of sample types such as culture supernatants, ascites or body fluids without first purifying the antibody from other contaminants.

Highlights:

- Easy-to-use particle-based antibody titer determination kit
- · Start of assay to recovery of result in less than one hour
- Four times faster than classical ELISA-based protocols
- Convenient design perform the assay in a 96-well plate and measure the result in a microplate reader
- Measures antibodies from culture supernatants ascites or body fluids
- Measures humanized antibodies and chimeras with intact Fc regions
- · No cross-reactivity with Ig from other species



Thermo Scientific Easy-Titer IgG and IgM Assay Kit protocol. A simple assay makes for an easy-to-perform assay protocol. Easy-Titer IgG Assay Kits feature a simple procedure that reduces hands-on time and requires fewer steps that lead to more reproducible results. The entire process can be completed easily in about 30 minutes.

Specialty Assays – Antibodies

Performance Specifications:

Specificity

· Against all IgG subclasses (human, mouse or rabbit)

Sensitivity

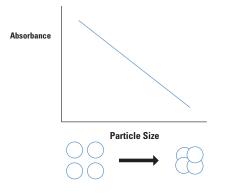
- Detection limit: 15 ng/ml
- Detection range (standard curve): 15 to 300 ng/ml

Coefficient of Variation (intra- and interassay): < 5%

Reaction time: 10 minutes

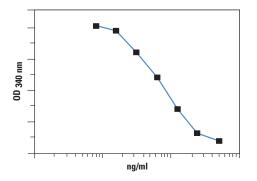
• Read results at 340 nm or 405 nm

Standard curve calculations are compatible with software supplied for use with microplate readers.



How the assay works:

- Monodisperse beads sensitized with a specific antibody absorb at 340 and 405 nm
- The beads agglutinate in the presence of human IgG or IgM
- Larger diameter clusters form that absorb less efficiently at 340 and 405 nm
- This decrease in absorbance is proportional to antibody concentration



Typical standard curve for Thermo Scientific Easy-Titer Kit. The unknown concentration of IgG is easily determined on a standard curve constructed with serial dilutions of a standard sample.

Ordering Information

| Product | Description | Pkg. Size |
|---------|--|----------------|
| 23310 | Easy-Titer Human IgG Assay Kit* Sufficient reagents for 96 tests (87 determinations and one standard curve). | Kit |
| | Includes: Goat Anti-Human IgG Sensitized Polystyrene Beads [Monodispersed, polystyrene IgG (Fc) sensitized beads are supplied suspended in a phosphate buffer, pH 7.4 and stabilized with BSA and 0.1% sodium azide] | 2 ml |
| | Easy-Titer Dilution Buffer Easy-Titer Blocking Buffer | 30 ml 15 ml |
| 23315 | Easy-Titer Human IgG Assay Kit* Includes: Goat Anti-Human IgG Sensitized Beads | Kit 2 ml |
| | Easy-Titer Dilution Buffer Easy-Titer Blocking Buffer | 30 ml 15 ml |
| 23300 | Easy-Titer Mouse IgG Assay Kit* Includes: Goat Anti-Mouse IgG Sensitized Beads | Kit 2 ml |
| | Easy-Titer Dilution Buffer Easy-Titer Blocking Buffer | 30 ml 15 ml |
| 23305 | Easy-Titer Rabbit IgG Assay Kit* Includes: Goat Anti-Rabbit IgG Sensitized Beads | Kit 2 ml |
| | Easy-Titer Dilution Buffer Easy-Titer Blocking Buffer | 30 ml 15 ml |
| 23325 | Easy-Titer Human IgG Assay Kit* Includes: Goat Anti-Human IgG | Kit 2 ml |

* Note: An IgG or IgM Standard is not included in these kits. Select the appropriate standard from the Related Products listed below.

Related Products

IgG Standards for Easy-Titer Kits

| Product | Description | Pkg. Size |
|---------|----------------------------|-----------|
| 31154 | Human IgG, Whole Molecule | 10 mg |
| 31146 | Human IgM, Whole Molecule | 2 mg |
| 31204 | Mouse IgG, Whole Molecule | 5 mg |
| 31235 | Rabbit IgG, Whole Molecule | 10 mg |

| Microplate Accessories | | | |
|------------------------|---|------------|--|
| Product | Description | Pkg. Size | |
| 15041 | 96-Well Plates Corner Notch | 100 plates | |
| 15031 | 8-Well Strip Plates Corner Notch Includes one strip well ejector per package. | 100 plates | |
| 23325 | Easy-Titer Human IgG (Gamma Chain) Assay Kit | Kit | |

Reference

Brown, M.A., et al. (2000). J. Biol. Chem. 275, 19795-19802.

Pierce Protease Assay Kits – Colorimetric and Fluorometric

Detects protease as low as 2 ng/ml in less than one hour!

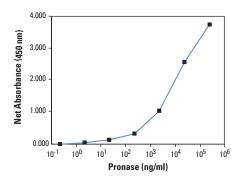
Thermo Scientific Pierce Protease Assay Kits are an ideal choice for performing routine assays necessary during the isolation of proteases, or for identifying the presence of contaminating proteases in protein samples. These protease assays are also ideal for studying pH or temperature vs. activity profiles of purified proteases.

Highlights:

- No corrosive precipitants used
- Entire assay can be run in microplates
- 1,000 times more sensitive, three times faster and uses half the sample of unmodified casein-based protease assays
- Total elapsed time to result less than one hour
- Measure multiple samples simultaneously in ELISA plate readers
- Time/temperature/pH easily manipulated to optimize sensitivity

The colorimetric Protease Assay Kit uses fully succinylated casein as substrate for this assay. Hydrolysis of this readily soluble casein substrate in the presence of protease results in the release of peptide fragments with free aminoterminal groups. Evidence of protease activity is obtained by reaction of these peptides with trinitrobenzene sulfonic acid (TNBSA), followed by measurement of the absorbance increase that is due to the formation of yellow colored TNB-peptide adducts. A standard protease is provided, allowing you to determine the concentration of protease in samples undergoing analysis.

Our Fluorescent Protease Assay Kit is based on a FITC-labeled casein. This sensitive assay can be used in either FRET or FP modes.



Sensitivity of the colorimetric Thermo Scientific Pierce Protease Assay.

Ordering Information

| Product | Description | Pkg. Size |
|---------|---|--------------------------|
| 23263 | Protease Assay Kit Sufficient material for 250 assays | Kit |
| | Includes: Succinylated Casein (supplied as a lyophilized salt-free powder) | 5 x 10 mg |
| | 2,4,6-Trinitrobenzene sulfonic acid (TNBSA) | 2 ml |
| | TPCK Trypsin standard (40 BAEE units/mg) | 50 mg |
| | BupH Borate Buffer Pack (makes 500 ml) | 1 pack |
| 23266 | Fluorescent Protease Assay Kit Sufficient material for at least 1,000 assays in a 96-well format. | Kit |
| | Includes: FITC-Casein, Lyophilized | 2.5 mg |
| | TPCK Trypsin BupH Tris Buffered Saline | 50 mg 1 pack |
| 23267 | FITC-Casein | 2.5 mg (1,000 assays) |

Reference

Rao, S.K., et al. (1997). Anal. Biochem. 250(2), 222-227.

Specialty Assays – Glycoproteins

Glycoprotein Carbohydrate Estimation Kit

Direct approach to the estimation of carbohydrate content in proteins with Thermo Scientific Glycoprotein Carbohydrate Estimation Kit.



Highlights:

- · Enables quick and easy identification of an unknown protein sample as a glycoprotein
- Estimates the percent carbohydrate content of a glycoprotein when run against a set of glycoprotein standards with known carbohydrate content
- · Complementary to electrophoresis, Western blotting and ELISA-based procedures often used to detect glycoprotein
- Determines carbohydrate content in three easy steps: (1) oxidize, (2) react and (3) read
- Entire assay performed in less than 75 minutes
- · All you need is this kit, a microplate and a plate reader to determine carbohydrate content

Ordering Information

| Product | Description | Pkg. Size |
|---------|--|-------------|
| 23260 | Glycoprotein Carbohydrate Estimation Kit Sufficient reagents for 250 microplate assays or 60 standard test tube assays. | Kit |
| | Includes: Sodium meta-Periodate | 500 mg |
| | Glycoprotein Detection Reagent | 500 mg |
| | Glycoprotein Assay Buffer Negative Controls: | 250 ml |
| | Lysozyme and BSA Positive Controls: | 2.5 mg each |
| | Ovalhumin | 2.5 mg |
| | Apo-Transferrin | 2.5 mg |
| | Fetuin | 0.25 mg |
| | α ₁ -Acid Glycoprotein | 0.25 mg |
| 23259 | Fluorescent Protease Assay Kit Sufficient material for at least 1,000 assays in a 96-well format. | Set |
| | Includes: Negative Controls: | |
| | Lysozyme and BSA Positive Controls: | 2.5 mg each |
| | Ovalbumin | 2.5 mg |
| | Apo-Transferrin | 2.5 mg |
| | Fetuin | 0.25 mg |
| | α ₁ -Acid Glycoprotein | 0.25 mg |
| 23262 | Glycoprotein Detection Agent | 1 g |

Assay Principle

The protein sample under analysis is oxidized and reacted with the exclusive Glycoprotein Detection Reagent. The resulting colored complex is read at 550 nm. From the absorbance of the resulting complex at 550 nm the approximate percentage of carbohydrate in the glycoprotein under analysis can be estimated.

1. Add 50 µl of protein standard or sample to each well.



3. Mix and incubate for 10 minutes at room temperature (RT).

2. Add 25 µl of 10 mM Sodium meta-Periodate

- 4. Add 150 μl of a 0.5% solution of Thermo Scientific Glycoprotein Detection Reagent in 1.0 M NaOH.
- 5. Mix and incubate at BT for 60 minutes.
- 6. Read the plate in a micro-plate reader at 550 nm. Interpolate the results of the unknown with the results of the standard proteins.





The Thermo Scientific Phosphoprotein Phosphate Estimation Assay microplate protocol.

in assay buffer.

Phosphoprotein Phosphate Estimation Kit

Get some basic questions about your target protein answered without having to perform a Western blot.

The novel protein characterization tool, that gives today's protein analyst the ability to quickly and reliably determine whether a purified target protein is phosphorylated and, if so, the extent of phosphorylation compared to a phosphoprotein of known phosphorus content. This easy-to-perform assay is specific for estimating phosphoserine or phosphothreonine post-translational modifications and has been adapted to both a tube and convenient microplate format. The Thermo Scientific Phosphoprotein Phosphate Estimation Assay provides answers that a traditional Western blot simply cannot, and you can get answers about five times faster, too.

Unique advantage of the assay chemistry

The specificity of this assay toward seryl and threonyl phosphate ester modifications can indirectly "detect" a phosphotyrosine modification should the result of the assay be negative. A negative result on a pure protein preparation can suggest that the protein is not phosphorylated or that the protein is, in fact, phosphorylated, but modified by way of the tyrosyl side chains. Further Western blot analysis can verify which conclusion is correct.

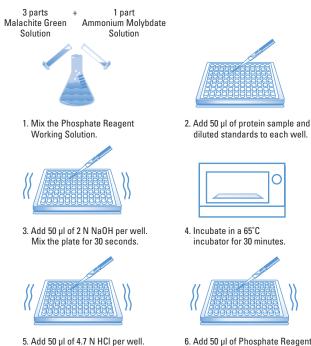
In addition, the Phosphoprotein Phosphate Estimation Assay Kit can also be used to determine the amount of a purified known phosphoprotein in a sample. A standard curve can be constructed using a purified preparation of the known protein.

Highlights:

- Easy-to-prepare working reagent
- Colorimetric detection
- · Use as qualitative or semi-quantitative assay
- Test tube or microplate assay option
- Estimate extent of phosphoserine/phosphothreonine modification
- Calculate the moles of phosphate (phosphorus) per mole of purified protein
- · Use as quantitative assay for known pure phosphoproteins
- · Results in about one hour
- Room temperature stability of kit components saves refrigerator and freezer space

Assay Principle

The Phosphoprotein Phosphate Estimation Assay is based on the alkaline hydrolysis of phosphate from seryl and threonyl residues in phosphoprotein and the quantification of the released phosphate by the use of malachite green and ammonium molybdate.



6. Add 50 μl of Phosphate Reagent per well. Mix for 30 seconds.



 Read the plate in a microplate reader at 650 nm. Plot the results of the unknown against the results of the standard protein provided. Calculate the approximate number of phosphorylation sites.

The Thermo Scientific Phosphoprotein Phosphate Estimation Assay microplate protocol.

Mix for 30 seconds.

7. Incubate for 30 minutes

at room temperature.

| Ordering Information | | |
|----------------------|---|-----------|
| Product | Description | Pkg. Size |
| 23270 | Phosphoprotein Phosphate Estimation Kit Sufficient reagents for 20 x 96-well microplate assays or 500 test tube assays. | Kit |
| | Includes: Ammonium Molybdate Solution | 25 ml |
| | Malachite Green Solution | 75 ml |
| | Phosvitin Positive Control | 1 mg |
| | BupH Tris Buffered Saline | 1 pack |
| Related P | roducts | |
| Product | Description | Pkg. Size |
| 24550 | GelCode [®] Phosphoprotein Staining Kit | Kit |

Pierce Quantitative Peroxide Assay Kits

Quickly measure peroxide contamination in various biological samples.

Highlights:

- · Fast and easy to use
- Peroxidase independent
- No lipid extraction necessary
- Spectrophotometric analysis
- · No heating required

Thermo Scientific Pierce Quantitative Peroxide Assays are the simplest assays for detecting the presence of peroxides in both aqueous and lipid-containing laboratory reagents. The basis of these assays is the complexing of ferric ion (Fe²⁺) by H_2O_2 in the presence of xylenol orange. Peroxides in the sample oxidize Fe²⁺ to Fe³⁺, and the Fe³⁺ will form a colored complex with xylenol orange that can be read at 560 nm.

The presence of hydrogen peroxide (H_2O_2) can now be detected to monitor any peroxide contamination that may be harmful to biological samples. When performed on a routine basis, our Quantitative Peroxide Assay can prevent inadvertent introduction of peroxides into your valuable samples. If the effects of peroxide cannot be avoided in a particular system, these assays will help you assess the risk to your sample.

References

Coutant, F., et. al. (2002). J. Immunol. **169**, 688-1695. Goyer, A., et. al. (2002). Eur. J. Biochem. **269**, 272-282. Requena, J. (2001). Proc. Nat. Acad. Sci., U.S.A. **98**, 69-74.

Comparison of Assay Protocols for Lipid Peroxide Content

Thermo Scientific Pierce Quantitative Peroxidase Assay

- 1. Mix one volume of Reagent A with 100 volumes of Reagent C to prepare Working Reagent.
- 2. Add 950 µl of Working Reagent to 50 µl of sample.
- 3. Incubate at room temperature for 30 minutes.
- 4. Read at 560 nm (or 595 nm for ELISA plate readers).

Total Time: 35 Minutes

Ordering Information

| Product | Description | Pkg. Size |
|---------|--|-----------|
| A | Quantitative Peroxide Assay Kit Aqueous compatible formulation. | Kit |
| | Includes: Reagent A (25 mM Ammonium Ferrous Sulfate) Reagent B (125 µM Xylenol Orange in water with Sorbitol) | 2 x 50 ml |
| 23285 | Quantitative Peroxide Assay Kit Lipid-compatible formulation. | Kit |
| | Includes: Reagent A (25 mM Ammonium Ferrous Sulfate) Reagent B (125 µM Xylenol | 4 x 25 ml |
| | Orange in methanol with BHT) | |

Thiobarbituric Acid Assay

- 1. Mix 0.1 ml sample, 0.4 ml H_2 O and 0.2 ml 7% SDS.
- 2. Stir gently and add 2 ml 0.1 N HCI.
- 3. Add 0.3 ml 10% phosphotungstic acid.
- 4. Incubate 5 minutes at room temperature.
- 5. Add 1 ml 0.67% thiobarbituric acid (TBA) and acetic acid.
- 6. Heat 45 minutes at 95°C.
- 7. Cool in ice bath.
- 8. Add 5 ml butanol.
- 9. Vortex and centrifuge for 15 minutes.
- 10. Determine lipid peroxide concentration in butanol layer by fluorescence at 515 nm excitation and 553 nm emission.

Total Time: 80-90 Minute

Pierce SILAC Protein Quantitation Kits

New options for absolute protein quantitation for challenging research situations.

Stable isotope labeling using amino acids in cell culture (SILAC) is a powerful method to identify and quantify relative differential changes in complex protein samples. The SILAC method uses *in vivo* metabolic incorporation of "heavy" ¹³C- or ¹⁵N-labeled amino acids into proteins followed by mass spectrometry (MS) analysis for accelerated comprehensive identification, characterization and quantitation of proteins.

Thermo Scientific Pierce SILAC Applications:

- Quantitative analysis of relative changes in protein abundance from different cell treatments
- Quantitative analysis of proteins for which there are no antibodies available
- · Protein expression profiling of normal vs. disease cells
- Identification and quantification of hundreds to thousands of proteins in a single experiment

Highlights:

- Efficient 100% label incorporation into proteins of living cells
- **Reproducible** eliminates intra-experimental variability caused by differential sample preparation
- Flexible media deficient in both L-lysine and L-arginine, allowing for more complete proteome coverage through dual amino acid isotope labeling
- Compatible label proteins expressed in a wide variety of mammalian cell lines adapted to grow in DMEM or RPMI 1640 medium, including HeLa, 293T, COS7, U2OS, A549, A431, HepG2, NIH 3T3, Jurkat and others

SILAC requires growing mammalian cells in specialized media supplemented with light or heavy forms of essential amino acids; i.e., ¹²C₆ and ¹³C₆ L-lysine, respectively. A typical experiment involves arowing one cell population in medium containing light amino acids (control), while the other population is grown in the presence of heavy amino acids (experimental). The heavy and light amino acids are incorporated into proteins through natural cellular protein synthesis. After alteration of the proteome in one sample through chemical treatment or genetic manipulation, equal amounts of protein from both cell populations are then combined, separated by SDS-polyacrylamide gel electrophoresis and digested with trypsin before MS analysis. Because peptides labeled with heavy and light amino acids are chemically identical, they co-elute during reverse-phase column prefractionation and, therefore, are detected simultaneously during MS analysis. The relative peak intensities of multiple isotopically distinct peptides from each protein are then used to determine the average change in protein abundance in the treated sample (Figure 1).

Thermo Scientific Pierce SILAC Kits are compatible with mammalian cell lines adapted to grow in either DMEM or RPMI 1640 media. Each kit includes all necessary reagents to isotopically label cells, including media, heavy and light amino acids, and dialyzed serum. The heavy $^{13}C_6$ $^{15}N_4$ L-Arginine is available separately and can be added to SILAC media to enhance peptide isotope label coverage. When combined with Thermo Scientific Protein/Peptide Sample Enrichment Products, Pierce SILAC Protein Quantitation Kits also enable MS analysis of low-abundance proteins such as cell-surface proteins, organelle-specific proteins and protein post-translational modifications such as phosphorylation or glycosylation.

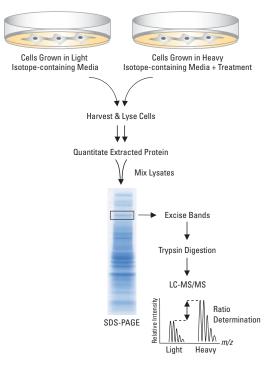


Figure 1. Schematic of SILAC workflow. A549 cells adapted to DMEM were grown for six passages (10 days) using SILAC DMEM (Product # 89983) containing 0.1 mg/ml heavy ¹⁵C₆ L-lysine-2HCl or light L-lysine-HCl supplemented with 10% dialyzed FBS. After 100% label incorporation, ¹³C₆ L-lysine-labeled cells were treated with 5 μ M camptothecin for 24 hours. Cells from each sample (light and heavy) were lysed using Thermo Scientific M-PER Mammalian Protein Extraction Reagent (Product # 78501). Samples were normalized for protein concentration using the Thermo Scientific Pierce BCA Protein Assay (Product # 23225), and 50 μ g of each sample were equally mixed before 4-20% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) analysis. Gels were stained with Thermo Scientific GelCode Blue Stain Reagent (Product # 24592) and proteins were digested and alkylated using the Thermo Scientific Pierce In-Gel Tryptic Digestion Kit (Product # 8871) before analysis using a Thermo Scientific LTQ Orbitrap Hybrid Mass Spectrometer.

Mass Spec Protein Detection

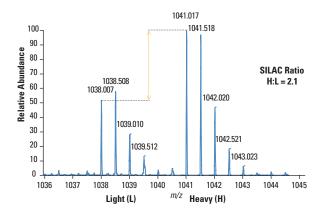


Figure 2. Representative MS spectra generated using SILAC. Light and heavy ($^{13}C_6$) L-lysine-containing peptides (AEDNADTLALVFEAPNQEK) from PCNA were analyzed by MS. Mass spectra of heavy peptides containing $^{13}C_6$ L-lysine have an increased mass of 6 Da and are shifted to the right of light peptide spectra by a mass to charge ratio (m/z) of 3 caused by a +2 ionization of peptides.

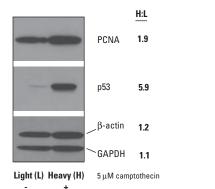


Figure 3. Comparison of A549 protein levels detected by Western blotting after camptothecin treatment. Ten micrograms of each light (L) and heavy (H) sample (Figure 1) were analyzed by 4-20% SDS-PAGE and Western blotting using specific antibodies.

References

- Everly, P.A., et al. (2004). Quantitative cancer proteomics: Stable isotope labeling with amino acids (SILAC) as a tool for prostate cancer research. Mol & Cell Proteomics 3.7, 729-735.
- 2. Mann, M. (2006). Functional and quantitative proteomics using SILAC. *Nature Reviews* 7, 952-959.
- 3. Levine, A.J. (1997). p53, the cellular gatekeeper for growth and division. Cell 88, 323-331.

Ordering Information

| Product | Description | Pkg. Size |
|---------|--|------------------|
| 89982 | Pierce SILAC Protein Quantitation Kit – RPMI 1640 | Kit |
| | Includes: SILAC RPMI Media | 2 x 500 ml |
| | Dialyzed FBS | 2 x 50 ml |
| | ¹³ C ₆ L-Lysine-2HCl L-Lysine-2HCl | 50 mg |
| | L-Arginine-HCl | 50 mg |
| 89983 | Pierce SILAC Protein | 2 x 50 mg Kit |
| 03303 | Ouantitation Kit – DMEM | KIL |
| | Includes: SILAC DMEM Media | 2 x 500 ml |
| | Dialyzed FBS | 2 x 50 ml |
| | ¹³ C ₆ L-Lysine-2HCI | 50 mg |
| | L-Lysine-2HCI | 50 mg |
| | L-Arginine-HCI | 2 x 50 mg |
| 89984 | SILAC RPMI Media | 500 ml |
| | (RPMI-1640 Medium minus L-Lysine | |
| | and L-Arginine) | |
| 89985 | SILAC DMEM Media | 500 ml |
| | (DMEM Medium minus L-Lysine | |
| | and L-Arginine) | |
| 89986 | Dialyzed FBS | 50 ml |
| 89987 | L-Lysine-2HCl | 50 mg |
| 89988 | ¹³ C ₆ L-Lysine-2HCl | 50 mg |
| 89989 | L-Arginine-HCl | 50 mg |
| 89990 | ¹³ C ₆ ¹⁵ N ₄ L-Arginine-HCI | 50 mg |

The purchase of this product conveys a non-transferable license to the Purchaser to use this product in methods protected under U.S. Patent 6,653,076 (owned by University of Washington) for research purposes only.

TMT Isobaric Mass Tagging Kits and Reagents

Complete kits for quantitative protein expression analysis.

Changes in protein expression and post-translational modifications are essential mechanisms of biological regulation and disease. Advancements in mass spectrometry (MS) instrumentation, bioinformatics and quantification methods, such as label-free quantification, metabolic labeling and chemical tagging, now enable researchers to identify and quantitatively analyze thousands of proteins in a given sample with a high degree of accuracy.

Highlights:

- Enables protein identification and quantitation from multiple samples of cells, tissues and biological fluids
- Consistent chemistry allows efficient transition from method development to multiplex quantitation, enabling biomarker discovery research
- Efficient labeling of membrane and post-translationally modified proteins
- Expandable system allows concurrent multiplexing of up to six different samples in a single experiment (Figure 2)
- Optimized fragmentation and fully supported quantitation with Thermo Scientific Proteome Discoverer 1.0 for all Thermo Scientific LC MS/MS platforms, such as Thermo Scientific LTQ XL and LTQ Orbitrap XL Systems (Figures 3-4)

Applications:

- Protein identification and quantitation from multiple samples of cells, tissue or biological fluids
- Protein expression profiling of normal vs. disease states or control vs. treated
- Multiplex up to six different samples concurrently in a single experiment
- Quantitative analysis of proteins for which no antibodies are available
- Identification and quantitation of membrane and post-translationally modified proteins
- Identification and quantification of hundreds to thousands of proteins in a single experiment

The Thermo Scientific TMT Isobaric Mass Tagging Kits and Reagents enable concurrent identification and quantitation of proteins in different samples using tandem mass spectrometry. The TMT Isobaric Tags are small chemical molecules with identical structure that covalently attach to the amino groups of lysine residues and the N-terminal of peptides, thereby labeling various peptides in a given sample (Figure 1). During the MS/MS analysis, each isobaric tag produces a unique reporter ion signature that makes quantitation possible. In the first MS analysis, the labeled peptides are indistinguishable from each other; however, in the tandem MS mode during which peptides are isolated and fragmented, each tag generates a unique reporter ion. Protein quantitation is then accomplished by comparing the intensities of the six reporter ions in the MS/MS spectra.

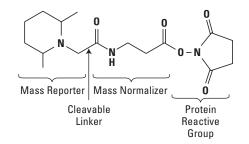


Figure 1. Structural design of a tandem mass tag. Mass reporter: Each member has a unique mass and reports sample-specific abundance of a labeled peptide during MS/MS analysis. Cleavable linker: Preferentially fragments under typical MS/MS conditions to release the mass reporter. Mass normalizer: Each member has a unique mass that balances the mass reporter, ensuring the same overall mass for all tags in a set. Reactive group: Reactive NHS ester provides high-efficiency amine-specific labeling of proteins/peptides.

Our complete family of isobaric tandem mass tags (TMT) are uniquely designed to enable a rapid and cost-effective transition from method development to high-throughput protein quantitation. The tags consist of TMTzero[™], the TMTduplex[™] set and the TMTsixplex[™] set (Figure 3).

Mass Spec Protein Detection

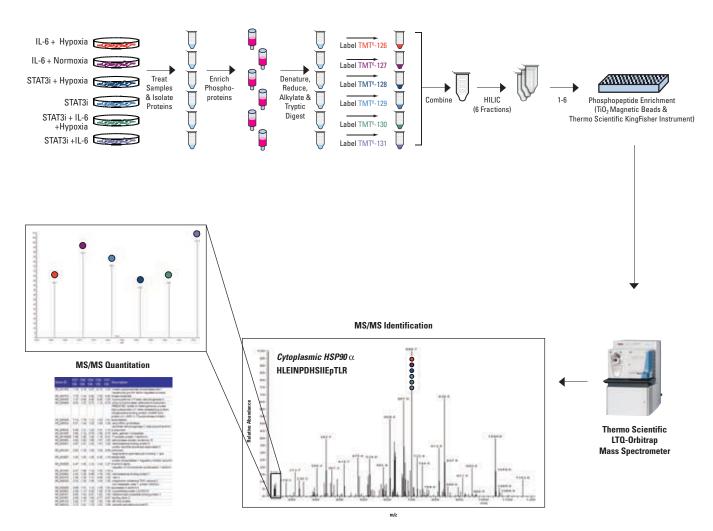
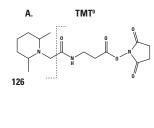
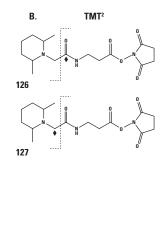


Figure 2. Quantitative phosphoproteomic workflow. The procedure combines phosphoprotein enrichment, isobaric mass tag (TMT) labeling, and phosphopeptide enrichment before mass spectrometry.





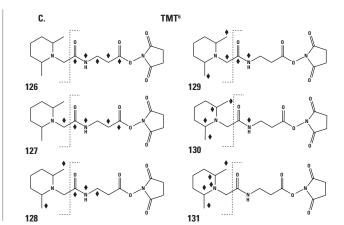


Figure 3. The TMT family of isobaric tag reagents. A. TMTzero has no isotopic substitutions and is used for method development. B. A pair of isobaric mass labels with a single isotopic substitution per tag is used for simple pairwise comparisons of relative protein expression. C. A sixplex of isobaric mass labels each with five isotopic substitutions per tag is used. Used for complex analyses including multiplex patient screening, time-course analysis or dose escalation studies.

The TMTzero Label Reagent allows testing and optimization of sample preparation, labeling, fractionation and MS fragmentation for peptide identification and reporter detection without using the more costly isotope-labeled compounds. The TMTduplex Reagent Set allows twoplex protein profiling for small studies. The TMTsixplex Reagent Set allows sixplex protein profiling for multiple conditions, including time courses, dose responses, replicates or multiple sample comparisons. Each tag is based on the same chemical structure, eliminating the need to modify labeling conditions or HPLC separation conditions between experiments.

This allows the TMT tags to be used as mass tags because the TMTzero and TMTsixplex tags have the same structure, but differ in mass by 5 Da. These tags can also be integrated into complex workflows, such as phosphoprotein and phosphopeptide enrichment (Figure 2).

The tags are provided as standalone sets or in optimized kit formats containing all necessary reagents and controls for maximum flexibility, convenience and reliability. The TMT Reagents combined with Thermo Scientific instruments and software provide a complete and integrated solution to perform absolute quantitation of target proteins (Figure 2, 4).

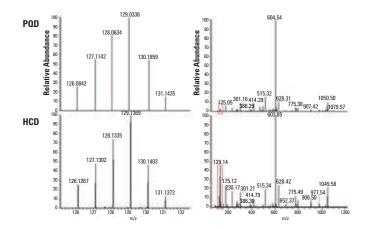


Figure 4. Analysis of a TMTsixplex-labeled peptide by Pulsed Q Dissociation (PQD) and High Energy Collision Dissociation (HCD). TMTsixplex-labeled A*ITIFQER (2+) from rabbit glyceraldehyde-3-phosphate dehydrogenase in a 10-protein sample were mixed at a 1:2:3:4:2:0.5 ratio. Shown are the MS/MS spectra for the peptide fragment and reporter ion regions. PQD fragmentation was performed on a LTQ XL[™] Ion Trap and by HCD on a high-resolution Orbitrap XL Mass Spectro.

References

- Thompson, A., et al. (2003). Tandem mass tags: a novel quantification strategy for comparative analysis of complex protein mixtures by MS/MS. Anal. Chem. 75(8), 1895-1904.
- Dayon, L., et al. (2008). Relative quantification of proteins in human cerebrospinal fluids by MS/MS using 6-plex isobaric tags. Anal. Chem. 80(8), 2921-2931.

Ordering Information

| Product | Description | Pkg. Size |
|---------|--|-----------|
| 90063 | TMTduplex Isobaric Mass Tagging Kit Labeling Reagents for Multiplexed and Absolute Protein Quantification | Kit |
| 90064 | TMTsixplex Isobaric Mass Tagging Kit Labeling Reagents for Multiplexed and Absolute Protein Quantification | Kit |
| 90065 | TMTduplex Label Reagent Set Labeling Reagents for Multiplexed and Absolute Protein Quantification | Kit |
| 90066 | TMTsixplex Label Reagent Set Labeling Reagents for Multiplexed and Absolute Protein Quantification | Kit |
| 90067 | TMTzero Label Reagent Labeling Reagent for Multiplexed and Absolute Protein Quantification | 5 vials |

Protein Assay Selection Guide Poster

The updated Thermo Scientific Pierce Protein Assay Selection Guide Poster (Product # 1601652) measures 32 5/8" x 24 3/8" (~83 cm x 62 cm) and provides an overview of all of our protein assays, including the recently introduced BCA, Reducing Agent Compatible and Pierce 660 nm Protein Assay. The poster outlines each protein assay's advantages, applications, standard assay protocol, precautions, reaction schemes, typical standard curves, detec-



tion range and interfering substance information. If your lab performs protein assays, you'll want to have this poster on your wall.

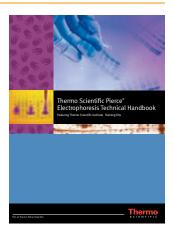
Protein Purification Technical Handbook

This 81-page handbook provides protocols and technical and product information to help maximize results for protein purification. It also includes background and trouble shooting advice for covalent coupling of affinity ligands to chromatography supports, avidin:biotinbinding, affinity purification of antibodies, IP and co-IP, affinity procedures for contaminant removal, and related procedures.



Electrophoresis Technical Handbook

This 44-page reference guide provides information to improve the speed, convenience and sensitivity of your protein gel electrophoresis and staining applications. The handbook covers all aspects of electrophoresis – from sample and gel preparation to choice of molecular weight markers. In addition, it contains an extensive section on protein gel-staining techniques and products.



Cell Lysis Technical Handbook

This handbook provides protocols and technical and product information to help maximize results for Protein/Gene Expression studies. The handbook provides background, helpful hints and troubleshooting advice for cell lysis, protein purification, cell fractionation, protease inhibitors and protein refolding. The handbook is an essential resource for any laboratory studying Protein/ Gene Expression.



To request a copy of any of these pieces, visit *www.thermo.com/ pierce* or contact your local office or distributor.

Tandem Mass Tags and methods of their use are protected by pending patents applications and granted patents worldwide including European Patent EP # 1,275,004, U.S. Patent # 7,294,456 and U.S. Patent Application 10/489,341.

SuperSignal® Technology is protected by U.S. Patent # 6,432,662.

Micro BCA[™] and BCA[™] Assay Technologies are protected by U.S. Patent # 4,839,295.

B-PER® Technology is protected by U.S. Patent # 6,174,704.

Easy-Titer® IgG Assay Technology is protected by U.S. Patent # 5,043,289 and European Patent # 0266278B1. Slide-A-Lyzer Dialysis Cassette Technology is protected by U.S. Patent 5,503,741 and 7,056,440; CA 2,170,738;

and EP 0 720 508 B1.

U.S. Patents pending on HisProbe[™]-HRP, GelCode Blue Stain Technology and Pierce 660 nm Protein Assay.

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