



Featuring the **Gold Standards** in Band Identification



Now includes
BenchMark™ and
MagicMark™

With Invitrogen's protein standards you can expect:

- Clear band resolution
- Accurate, easy-to-read results
- A time-saving, load-and-go format

Setting the Standard

Invitrogen's complete line of protein markers sets the standard in electrophoresis band identification. Eight different protein standards are available, each with unique advantages, all offering maximum convenience. With every standard, you'll get:

- Unambiguous identification—proteins resolve into clear, sharp bands for precise results
- Wide size range of protein markers—enables you to identify diverse protein molecular weights easily
- Load-and-go format—standards are supplied ready to use, with no need to mix, reduce, or heat before using
- Consistent high quality—standards are strictly quality controlled to ensure consistent band intensity

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A standard for every application

A broad selection of protein standards is available to meet your electrophoresis needs. Whether you're approximating molecular weight, verifying transfer efficiency, or determining protein isoelectric point, you're sure to find a standard that meets your application needs (Table 1). Each protein standard is supplied in a ready-to-use format, eliminating the need to dilute, mix, or heat before loading.

Molecular weight standards for SDS-PAGE

Protein molecular weight standards provide the means to estimate molecular weight as well as to confirm electrophoresis and transfer runs in SDS-PAGE (sodium dodecyl sulfate polyacrylamide gel electrophoresis). By constructing a standard curve with a series of standards, you can estimate the apparent molecular weight of a sample protein based on its relative mobility. Unstained protein molecular weight standards (*i.e.*, Mark12™ Unstained Standard, BenchMark™ Protein Ladder) provide more

accurate size estimation than pre-stained standards. Pre-stained standards, however, are well suited for confirming electrophoresis runs and estimating transfer efficiency. They also offer good molecular weight approximation if they are calibrated according to the gel/buffer system in use.

Marker for western blots

MagicMark™ Western Protein Standard provides an easy and convenient means to accurately estimate protein molecular weight directly on western blots. You can visualize MagicMark™ standard bands simultaneously with your target protein using the same antibody conjugate and protocol.

IEF marker for isoelectric focusing gel

The SERVA™ Liquid Mix IEF Marker enables you to determine the isoelectric point (pI) of unknown protein samples on horizontal and vertical isoelectric focusing (IEF) gels.

Table 1 - Advantages of Invitrogen's protein standards

Standard	MW/pI range	Advantage
MultiMark™ Multi-Colored Standard	4-250 kDa†	Multi-colored bands for at-a-glance identification
SeeBlue™ Plus2 Pre-Stained Standard	4-250 kDa†	Sharply resolved bands, including two colored bands for easy analysis
SeeBlue™ Pre-Stained Standard	4-250 kDa†	Sharpest, most consistent pre-stained bands
BenchMark™ Pre-Stained Protein Ladder	~ 10-190 kDa†	Sharp blue bands with one pink band for easy band identification
Mark12™ Unstained Standard	2.5-200 kDa	Accurate estimation of molecular weight with the broadest molecular weight range
BenchMark™ Protein Ladder	10-220 kDa	Accurate estimation of molecular weight including two triple-intensity bands for easy reference
MagicMark™ Western Protein Standard	20-120 kDa	Accurate molecular weight estimation directly on western blots
SERVA™ Liquid Mix IEF Marker	pI 3.5-10.7	Accurate determination of protein isoelectric points

† Actual range is dependent upon gel type and buffer system

For the **easiest** molecular weight band identification

The **MultiMark® Multi-Colored Standard** gives you a colorful alternative to blue, pre-stained molecular weight markers. It consists of 9 multi-colored bands (Figure 1).

Because each band is a different color, you'll be able to easily and immediately identify the molecular weight of each protein.

Figure 1 - Apparent molecular weights¹ of the MultiMark® Multi-Colored Standard

Protein	SDS-PAGE System				
	Tris-Glycine	Tricine	NuPAGE® Bis-Tris (MES)	NuPAGE® Bis-Tris (MOPS)	NuPAGE® Tris Acetate
Myosin	250	208	185	188	209
Phosphorylase B	148	105	98	97	111
Glutamic Dehydrogenase	60	53	52	52	52
Carbonic Anhydrase	42	34	31	33	34
Myoglobin-Blue	30	23	19*	21*	n/a
Myoglobin-Red	22	17	17*	19*	n/a
Lysozyme	17	13	11	12	n/a
Aprotinin	6	7	6	n/a	n/a
Insulin, B chain	4	4	3	n/a	n/a

▲
NuPAGE® 4-12% Bis-Tris Gel w/MES SDS Buffer

Approximate Molecular Weights (kDa)

* Note: The 2 migration patterns of Myoglobin Red and Blue are reversed in the NuPAGE® Bis-Tris MES and MOPS Buffers compared to the Tris-Glycine and Tricine Systems.

¹ Migration patterns in several buffer systems are shown because protein bands will have different mobilities in different SDS-PAGE buffer systems. For more information on this phenomenon, contact a Technical Service Representative at 800.955.6288, ext. 2 or review the technical note entitled "Accurate calibration of molecular weight standards for different buffer systems" on our web site at www.invitrogen.com.

Sharp bands and easy analysis

The SeeBlue® Plus2 Pre-Stained Standard consists of 10 pre-stained protein markers—8 blue and 2 colored—that resolve into sharp distinct bands

(Figure 2). The two colored bands make it easy to immediately identify the protein markers.

Figure 2 - Apparent molecular weights* of the SeeBlue® Plus2 Pre-Stained Standard

Protein	SDS-PAGE System				
	Tris-Glycine	Tricine	NuPAGE® Bis-Tris (MES)	NuPAGE® Bis-Tris (MOPS)	NuPAGE® Tris-Acetate
Myosin	250	210	188	191	210
Phosphorylase B	148	105	98	97	111
BSA	98	78	62	64	71
Glutamic Dehydrogenase	64	55	49	51	55
Alcohol Dehydrogenase	50	45	38	39	41
Carbonic Anhydrase	36	34	28	28	n/a
Myoglobin-Red	22	17	17	19	n/a
Lysozyme	16	16	14	14	n/a
Aprotinin	6	7	6	n/a	n/a
Insulin, B chain	4	4	3	n/a	n/a

▲
NuPAGE® 4-12% Bis-Tris Gel w/MES SDS Buffer

Approximate Molecular Weights (kDa)

* Migration patterns in several buffer systems are shown because protein bands will have different mobilities in different SDS-PAGE buffer systems. For more information on this phenomenon, contact a Technical Service Representative at 800 955 6288, ext. 2 or review the technical note entitled "Accurate calibration of molecular weight standards for different buffer systems" on our web site at www.invitrogen.com.

Sharpest band resolution

For sharp, consistent pre-stained bands, the **SeeBlue[®] Pre-Stained Standard** is the molecular weight standard of choice. It consists of 9 individual protein bands—all

blue—that provide high resolution in any SDS-PAGE system (Figure 3). Since SeeBlue[®] is supplied pre-stained, there is no need to stain the gel in order to visualize the standard.

Figure 3 - Apparent molecular weights* of the SeeBlue[®] Pre-Stained Standard

Protein	SDS-PAGE System				
	Tris-Glycine	Tricine	NuPAGE [®] Bis-Tris (MES)	NuPAGE [®] Bis-Tris (MOPS)	NuPAGE [®] Tris Acetate
Myosin	250	210	188	191	210
BSA	98	78	62	64	71
Glutamic Dehydrogenase	64	55	49	51	55
Alcohol Dehydrogenase	50	45	38	39	41
Carbonic Anhydrase	36	34	28	28	n/a
Myoglobin	30	23	18	19	n/a
Lysozyme	16	16	14	14	n/a
Aprotinin	6	7	6	n/a	n/a
Insulin, B chain	4	4	3	n/a	n/a

Approximate Molecular Weights (kDa)

▲ NuPAGE[®] 4-12% Bis-Tris Gel w/MES SDS Buffer

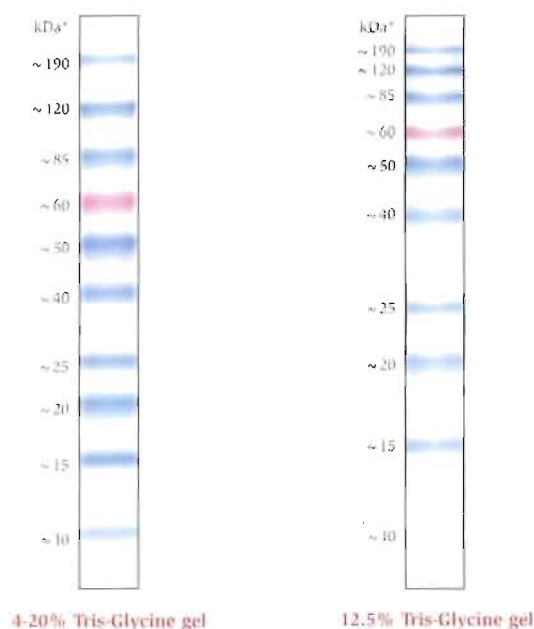
Migration patterns in several buffer systems are shown because protein bands will have different mobilities in different SDS-PAGE buffer systems. For more information on this phenomenon, contact a Technical Service Representative at 800 955 6288, ext. 2 or review the technical note entitled "Accurate calibration of molecular weight standards for different buffer systems" on our web site at www.invitrogen.com.

Monitor electrophoretic separation in real time

The **BenchMark™ Pre-Stained Protein Ladder** allows you to monitor the progress and quality of an electrophoretic separation. Like other pre-stained protein standards, you can also use the BenchMark™ Pre-Stained Ladder to estimate the efficiency of protein transfer when performing western blotting.

The standard proteins are affinity-purified and covalently coupled with dye. You'll see superb band sharpness (Figure 4), get easy orientation with a pink reference band, and approximate molecular weight without difficulty.

Figure 4 - BenchMark™ Pre-Stained Protein Ladder



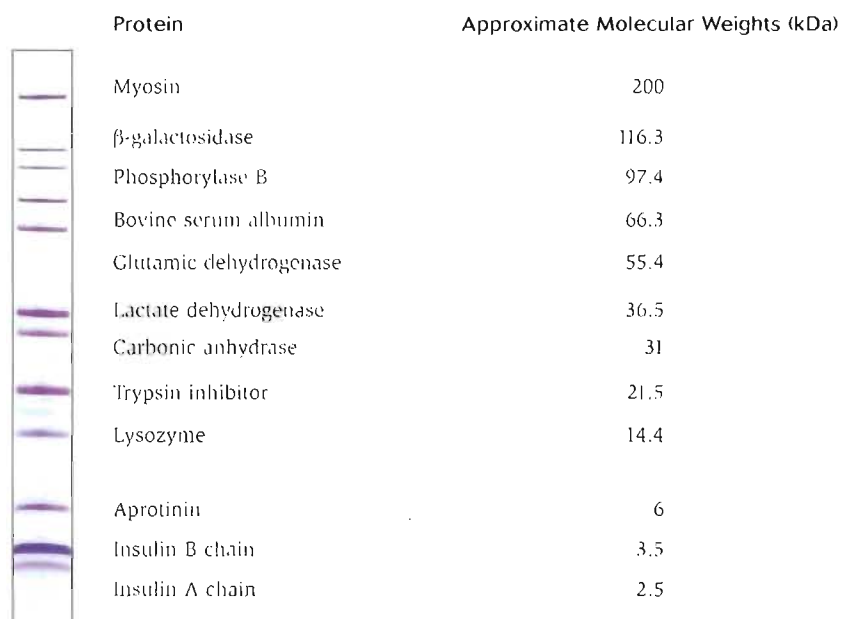
* Coupling of the chromophores to the proteins affects their apparent molecular weight in SDS-PAGE relative to unstained standards. Each band in the pre-stained ladder is calibrated against unstained BenchMark™ Protein Ladder on a 4-20% Tris-Glycine gel and the apparent molecular weight is reported on the product profile. The pre-stained protein ladder should only be used to determine an approximate size molecular weight.

Most accurate estimation of molecular weight

The 12 protein bands on the **Mark12™ Unstained Standard** migrate the closest to their true molecular weight. That's because the dye used in pre-stained standards can affect band migration patterns, resulting in apparent molecular weights that are different from those of patterns in their unstained state. Since the proteins in the Mark12™ stan-

dard are unstained, their migration pattern is not modified by the dye, allowing you to achieve the most accurate estimation of molecular weight. The Mark12™ bands appear sharp and distinct when visualized with Coomassie® (Figure 5) or silver stain.

Figure 5 - The Mark12™ Unstained Standard



▲
NuPAGE® 4-12% Bis-Tris
Gel w/MES stained with
Coomassie® R-250

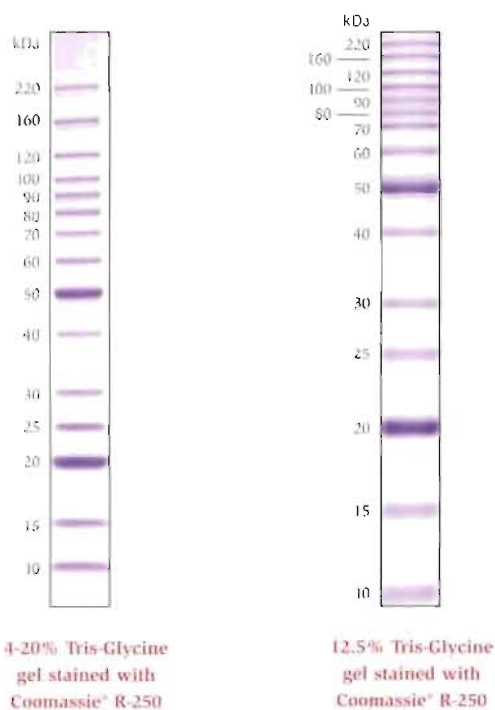
Note: The apparent molecular weights stated above apply to the Tris-glycine, Tricine, and NuPAGE® Systems.

The benchmark of protein ladders

BenchMark[®] Protein Ladders are ideal for estimation of molecular weight of unknown proteins by SDS-polyacrylamide gel electrophoresis. Affinity-purified proteins generate sharp, intense bands without background for accuracy. You can visualize bands using

either Coomassie Brilliant Blue R-250 stain (Figure 6) or silver stain. Standard bands, including two triple-intensity reference bands, are in easy-to-identify increments for proper band identification.

Figure 6 - BenchMark[®] Protein Ladder

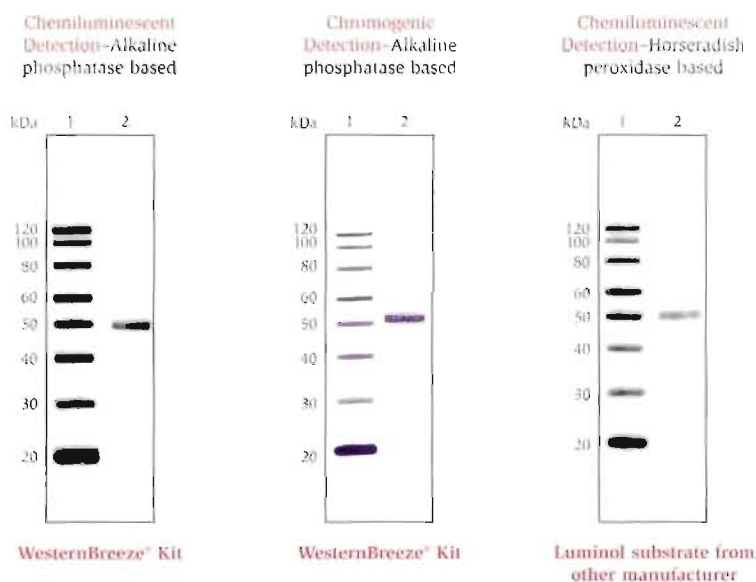


Easy, accurate western blot analysis

MagicMark™ Western Protein Standard lets you accurately estimate molecular weight directly on western blots. Each protein of the MagicMark™ Standard contains an IgG binding site. You can visualize MagicMark™ protein bands simultaneously with your target protein using the same antibody conjugate and protocol (Figure 7). Use

chemiluminescent, fluorescent, or colorimetric detection methods for your analysis. With MagicMark™, you'll bypass steps required in conventional methods, yet obtain sharp bands and precise molecular weight estimation on your western blots.

Figure 7 - Sharp bands of MagicMark™ detected simultaneously with target protein



MagicMark™ Standard and an expressed protein containing a V5 epitope tag were separated on a NuPAGE™ 4-12% Bis-Tris Gel and transferred to a nitrocellulose membrane. The blots were probed with a 1:5,000 dilution of mouse anti-V5 primary antibody and detected with the indicated western detection systems.

Lane 1: 5 µl of MagicMark™ standard
Lane 2: 2 ng of protein

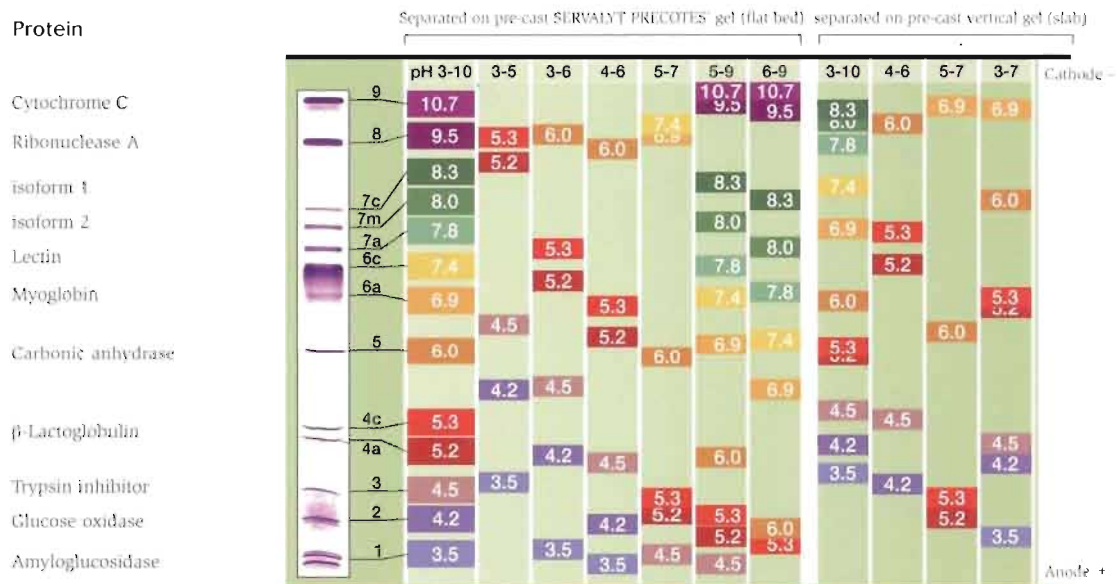
For more information on MagicMark™ Western Protein Standard, request the MagicMark™ brochure at www.invitrogen.com.

Accurate estimation of isoelectric points

The **SERVA® Liquid Mix IEF Marker** provides 9 different proteins with 13 isoforms (Figure 8) for determining the isoelectric point (pI) of a full range of unknown protein samples in vertical or horizontal IEF gels. Since the standards are salt-free, you'll get

straight bands for precise results. Unlike other IEF markers, the SERVA® Marker contains bromophenol blue and methyl red dyes, so you can visualize the progress of the markers during electrophoresis.

Figure 8 - Schematic representation of SERVA® IEF markers in various pH fractions



A standard to meet **your needs**

At Invitrogen you're sure to find a standard for every protein electrophoresis application (Table 2). Each standard provides clear band resolution so you can accurately estimate molecular weights or

isoelectric points. In addition, you'll save time with the suitable load-and-go format. Call and order today.

Table 2 - Choosing a protein standard

	MultiMark™	SeeBlue® Plus2/ SeeBlue®	BenchMark™ Pre-stained	Mark12™	BenchMark™	MagicMark™	SERVA® IEF Marker 3-10
Application							
SDS-PAGE	Good	Good	Good	Good	Good	Good	n/a
IEF Gel	n/a	n/a	n/a	n/a	n/a	n/a	Best!
Western Blot	Good	Good	Good	Good	Good	Best!	n/a
Immediate Band Identification	Best!	Good	Good	*	*	*	*
Sharp Bands	Good	Best!	Good	Best!	Best!	Best!	Good
MW Estimation	Good	Good	Good	Best!	Best!	Best!	pI Estimation
Monitor Migration during Electrophoresis	Good	Good	Good	n/a	n/a	n/a	n/a
Silver Staining	Good	Good	Good	Best!	Best!	Good	Good
MW/pI Range	4-250 kDa	4-250 kDa	~ 10-190 kDa	2.5-200 kDa	10-220 kDa	20-120 kDa	pI 3.5-10.7
Type	Natural Proteins	Natural Proteins	Recombinant Proteins	Natural Proteins	Recombinant Proteins	Recombinant Proteins	Natural Proteins

* Bands visible only after staining.

Description	Quantity	Cat. no.
MultiMark™ Multi-Colored Standard	500 µl	LC5725
SeeBlue™ Plus2 Pre-Stained Standard	500 µl	LC5925
SeeBlue™ Pre-Stained Standard	500 µl	LC5625
BenchMark™ Pre-Stained Protein Ladder	2 x 250 µl	10748-010
Mark12™ Unstained Standard	1 ml	LC5677
BenchMark™ Protein Ladder	2 x 250 µl	10747-012
MagicMark™ Western Protein Standard	250 µl	LC5600
SERVA™ IEF 3-10 Marker	500 µl	39212-01

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