TECHNICAL MANUAL

TNT® Quick Coupled Transcription/Translation Systems

Instructions for use of Products
L1170, L1171, L2080 and L2081

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TNT® Quick Coupled Transcription/Translation Systems

All technical literature is available at: www.promega.com/protocols/
Visit the web site to verify that you are using the most current version of this Technical Manual.
E-mail Promega Technical Services if you have questions on use of this system: techserv@promega.com

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1. Description

The TNT® Quick Coupled Transcription/Translation Systems are convenient single-tube, coupled transcription/translation reactions for eukaryotic in vitro translation. The original TNT® Coupled Reticulocyte Lysate Systems simplified the process and reduced the time required to obtain in vitro translation results compared with standard rabbit reticulocyte lysate systems (1). Standard rabbit reticulocyte systems commonly use RNA synthesized in vitro from SP6, T3 or T7 RNA polymerase (1). The TNT® Quick Coupled Transcription/Translation System further simplifies the process by combining the RNA polymerase, nucleotides, salts and Recombinant RNasin® Ribonuclease Inhibitor with the reticulocyte lysate solution to form a single TNT® Quick Master Mix (Figure 1). For most gene constructs, the TNT® Quick reaction produces significantly more protein (two- to sixfold) in a 60- to 90-minute reaction than a standard in vitro rabbit reticulocyte lysate reaction using RNA templates.

The TNT® Quick Coupled Transcription/Translation System is available in two configurations for transcription and translation of genes cloned downstream from either the T7 or SP6 RNA polymerase promoters. To use these systems, 0.2–2.0µg of circular plasmid DNA containing a T7 or SP6 promoter, or a PCR-generated fragment containing a T7 promoter, is added to an aliquot of the TNT® Quick Master Mix and incubated in a 50µl reaction volume for 60–90 minutes at 30°C. The synthesized proteins are then analyzed by SDS-polyacrylamide gel electrophoresis (SDS-PAGE) and detected. Included with the TNT® Quick System is a luciferase-encoding control plasmid and Luciferase Assay Reagent, which can be used in a non-radioactive assay for rapid (<30 seconds) detection of functionally active luciferase protein. Starting with either circular plasmid DNA or PCR-generated DNA, in vitro transcription/translation results may be obtained easily in 5–6 hours.

(!) PCR-generated fragments are not recommended for use with the SP6 promoter. Use the T7 promoter.

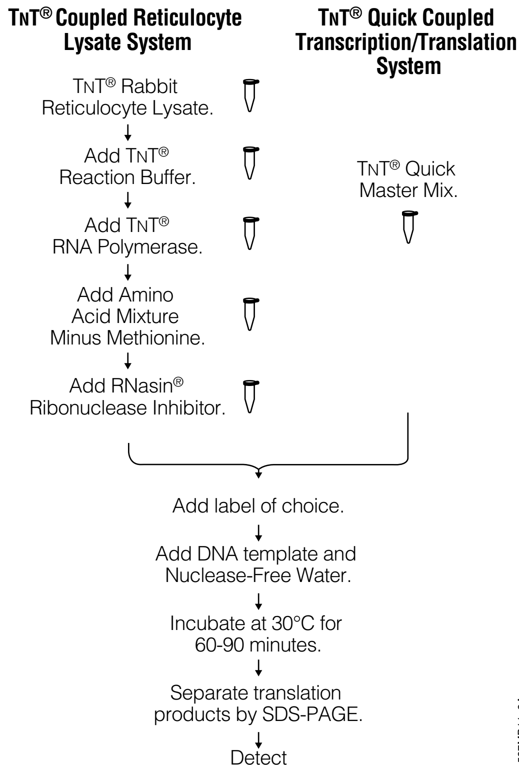

Figure 1. Comparison of the TNT® Coupled Reticulocyte Lysate System and the TNT® Quick Coupled Transcription/Translation System protocols.

1. Description (continued)

In addition to verifying the expected molecular weight of a gene construct, the TNT® Quick System is ideal for screening large numbers of constructs for either naturally occurring or deliberately engineered mutations. Applications of the system include:

- Truncation mutation analysis [e.g., the Protein Truncation Test (PTT)] (2)
- Drug screening (affecting translation rates)
- Mutation and detection analysis (i.e., enzyme kinetics)
- Protein:protein interactions (using GST pull-downs)
- Immunoprecipitation of protein complexes
- Protein dimerization assays
- Ligand-binding region determination/confirmation/competition assays
- In vitro expression cloning (IVEC) (functional genomics)
- Protein structure analysis
- Electrophoretic mobility shift assays (EMSAs) for DNA:protein interactions
- DNA footprinting and protein cross-linking studies
- Protein-RNA binding assays
- Post-translational modification tests
- Verification/characterization of cloned genes

The TNT® Quick Coupled Transcription/Translation Systems are also useful for detecting protein:protein interactions in vitro. [^{35}S]methionine-labeled proteins labeled using TNT® Quick Coupled Transcription/Translation System can be used as probes to detect interactions with suspected protein partners that have been expressed as GST-(glutathione-S-transferase) or epitope-tagged fusion proteins (3). [^{35}S]methionine-labeled proteins can be synthesized using coupled in vitro reactions from either full-length cDNAs or deletion mutants. The fusion proteins can be bound to an affinity matrix along with the radioactive proteins with which they interact (4–6). The bound radioactive proteins are then eluted and analyzed by SDS-PAGE or Western analysis (Figure 2; 6). The fusion tag approach has been used to study receptor-mediated control of apoptosis (7).

Alternatively, a non-radioactive approach may be used; the protein is labeled with biotinylated lysine (e.g., Transcend™ Biotinylated tRNA) or is fluorescently tagged (e.g., FluoroTect™ Green$_{Lys}$ System BODIPY®-FL-labeled tRNA [Cat.# L5001]) and combined with a GST-tagged protein. The biotinylated protein is detected by methods similar to those used in Western blotting (8,9). The fluorescently tagged protein can be detected from within the gel (10).

For a complete list of references for these and other applications, see reference 6 or visit the Promega Technical Resource Center citations at: **www.promega.com/citations/**

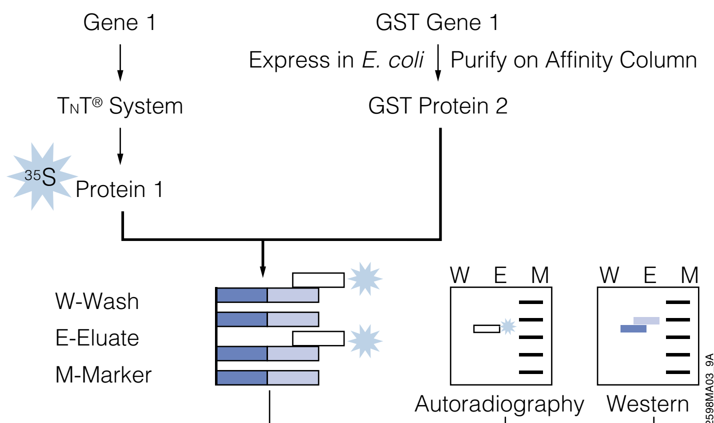

Figure 2. The study of protein:protein interactions using the TNT® Systems (6). This schematic shows translation of one protein with radioactive [^{35}S]methionine in a TNT® System reaction. Large amounts of the suspected partner protein are expressed and purified. A fusion tag (most commonly GST) is incorporated into this second protein to facilitate purification and subsequent capture steps. After the GST fusion protein is immobilized on sepharose (GST pulldowns), it is mixed with the protein produced in the TNT® reaction. The sepharose is washed to remove unbound protein, and the remaining bound proteins are eluted and analyzed on a gel. This technique allows measurement of the protein:protein interactions for both proteins and is often used to verify the in vivo results obtained from yeast two-hybrid experiments. Promega offers the MagneGST™ Pull-Down System (Cat.# V8870) for GST pull-down experiments.

2. Product Components and Storage Conditions

PRODUCT	CAT.#
TNT® T7 Quick Coupled Transcription/Translation System	**L1170**
TNT® SP6 Quick Coupled Transcription/Translation System	**L2080**

Each system contains sufficient reagents to perform approximately 40 × 50µl translation reactions. Includes:

- 1.6ml TNT® Quick Master Mix (8 × 200µl)
- 5µg SP6 or T7 Luciferase Control DNA (0.5µg/µl)
- 100µl T7 TNT® PCR Enhancer (L1170 only)
- 50µl Methionine, 1mM
- 250µl Luciferase Assay Reagent
- 1.25ml Nuclease-Free Water

2. Product Components and Storage Conditions (continued)

PRODUCT	CAT.#
TNT® T7 Quick Coupled Transcription/Translation System, Trial Size	L1171
TNT® SP6 Quick Coupled Transcription/Translation System, Trial Size	L2081

Each system contains sufficient reagents to perform approximately 5 × 50µl translation reactions. Includes:

- 200µl TNT® Quick Master Mix
- 5µg SP6 or T7 Luciferase Control DNA (0.5µg/µl)
- 100µl T7 TNT® PCR Enhancer (L1171 only)
- 50µl Methionine, 1mM

Storage and Stability: Store all components at –70°C. Product components are sensitive to CO_2 (avoid prolonged exposure), frequent temperature fluctuations and multiple freeze-thaw cycles, which can adversely affect stability, activity and performance. Luciferase Assay Reagent (LAR) is stable for at least 12 months if stored and handled properly.

Note: See Note 5, Section 4.C, for details on how to refreeze the lysate.

Note that the systems are shipped in foil packaging because the system is sensitive to carbon dioxide released from dry ice. If storing the system in a freezer containing dry ice, keep system components sealed in foil packaging for best results. **DO NOT** store the unfoiled lysate in the presence of dry ice. Prolonged exposure to dry ice causes significant loss of activity. The expiration date for the TNT® Quick Master Mix is listed on the product vial. **Do not freeze-thaw the Master Mix more than two times.**

3. General Considerations

3.A. DNA Template Considerations

DNA Expression Elements

1. In addition to circular plasmid DNA, PCR-generated DNA templates can be transcribed/translated using the T7 System. For maximal expression from such templates, we recommend that approximately 11bp be present upstream of the T7 RNA polymerase promoter for efficient promoter binding. A stop codon (usually UAA) is important for truncated gene products in order to prevent ribosomes from stalling at the ends of RNAs without stop codons. This can be done through appropriate primer design (11). The best transcription/translation results are obtained when the fragment contains the T7 RNA polymerase promoter. **We do not recommend using linear DNA with the SP6 System because of reduced transcription efficiencies.**

   Note: For coupled transcription/translation from PCR-generated templates, Promega offers TNT® T7 Quick for PCR DNA (Cat.# L5540).

2. While Rabbit Reticulocyte Lysate-based systems are less sensitive to 5´-untranslated region (UTR) secondary structure than other systems, it is still important to avoid strong hairpin secondary structure in the 5´-UTR region, because this can impair translation efficiency (12).
3. We have observed enhanced translation of proteins when using DNA constructs containing a poly(A) sequence downstream of the gene of interest. Poly(A) sequences are important for mRNA stability and can play a role in translation initiation in Rabbit Reticulocyte Lysate (13). For example, we have observed a two- to fivefold increase in the production of luciferase when the gene is cloned into the pSP64 Poly(A) Vector (Cat.# P1241).

Plasmid DNA

1. Residual ethanol should be removed from DNA preparations before they are added to the TNT® Quick Master Mix.
2. Linearized templates produced by restriction enzyme digestion should be cleaned up either by using the Wizard® PCR Preps DNA Purification System or by phenol:chloroform extraction, followed by ethanol precipitation, before use in the TNT® Quick reaction.
3. Plasmid DNA can be purified using the Wizard® *Plus* SV Minipreps DNA Purification System or the PureYield™ Plasmid Midiprep System. DNA prepared by the standard alkaline lysis method described by Sambrook, Fritsch and Maniatis (14) is also sufficiently clean for use in the TNT® Quick Coupled Transcription/Translation System. For most constructs, optimal results are obtained when 1µg of plasmid DNA template is used. However, we have used 0.2–2.0µg of DNA template and obtained satisfactory levels of translation. The use of more than 1µg of plasmid does not necessarily increase the amount of protein produced.
4. If linearizing plasmid DNA for use with the T7 System, avoid the use of restriction enzymes that yield 3´-overhangs (PstI, KpnI, SacI, SacII, BstXI, NsiI, ApaI and AatII), as aberrant transcription products can be produced (15). If no alternative enzyme is available, the 3´-overhang can be removed by adding T4 DNA polymerase.

   Note: If you are using a linearized plasmid as a template, include 1µl of the T7 TNT® PCR Enhancer in each 50µl reaction.
5. Check the sequence of the DNA template for the presence of additional upstream start codons. During translation, the ribosome is thought to scan from the 5´ end of the RNA and begin translation at the first AUG encountered. Thus, any AUGs within the transcribed portion of the vector or untranslated sequence of the insert may cause translation initiation to occur prior to the desired start codon and result in a shift in the reading frame or production of a larger protein than expected.

3.A. DNA Template Considerations (continued)

PCR-Generated DNA Templates

1. Because PCR DNA templates are usually much smaller than plasmid templates, the amount of DNA necessary for optimal expression is often less than for inserts cloned into plasmid vectors (e.g., for a 500bp PCR product, use 100–800ng for each 50µl TNT® Quick reaction).

   Note: For coupled transcription/translation from PCR-generated templates, Promega offers TNT® T7 Quick for PCR DNA (Cat.# L5540).

2. PCR products (5–7µl) can be used directly from the amplification reaction.

   Note: If you are using a PCR-generated template, include 1µl of the T7 TNT® PCR Enhancer in each 50µl reaction.

3.B. Creating a Ribonuclease-Free Environment

To reduce the chance of RNase contamination, gloves should be worn when setting up experiments, and microcentrifuge tubes and pipette tips should be RNase-free. It is not necessary to add Recombinant RNasin® Ribonuclease Inhibitor to the TNT® Quick reactions to prevent degradation of RNA, because it is already present in the TNT® Quick Master Mix.

3.C. Handling of Lysate

Except for the actual transcription/translation incubation, all handling of the TNT® Quick Master Mix should be done at 4°C. Any unused Master Mix should be refrozen as soon as possible after thawing to minimize loss of translational activity (see Note 5, Section 4.C). Do not freeze-thaw the Master Mix more than two times.

4. Translation Procedure

The following is a general guideline for setting up a transcription/translation reaction. Also provided are examples of standard reactions using [^{35}S]methionine (radioactive), Transcend™ Non-Radioactive Detection System (colorimetric or chemiluminescent) or FluoroTect™ Green$_{Lys}$ Systems (fluorescent). Using the Transcend™ Systems, biotinylated lysine residues are incorporated into nascent proteins during translation. This biotinylated lysine is added to the transcription/translation reaction as a precharged ε-labeled, biotinylated lysine-tRNA complex (Transcend™ tRNA) rather than a free amino acid. For more information on the Transcend™ Systems, request Technical Bulletin #TB182. The FluoroTect™ System uses a charged lysine tRNA labeled with the fluorophore, BODIPY®-FL, to incorporate fluorescently labeled lysine residues into the in vitro translation product. For more information on the FluoroTect™ System, request Technical Bulletin #TB285.

Note: Technical Manuals and Bulletins are available online at: **www.promega.com/protocols/** or by request from Technical Services.

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4.A. General Protocol for TNT® Quick Coupled Transcription/Translation Reactions Using Plasmid DNA

Materials to Be Supplied by the User

- Nuclease-Free Water (Cat.# P1193)
- Radiolabeled amino acid (for radioactive detection; Note 4, Section 4.C) or Transcend™ tRNA (Cat.# L5061) or Transcend™ Colorimetric (Cat.# L5070) or Chemiluminescent (Cat.# L5080) Translation Detection System (for non-radioactive detection) or FluoroTect™ $Green_{Lys}$ in vitro Translation Labeling System (for fluorescent detection; Cat.# L5001).

1. Remove the reagents from storage at –70°C. Rapidly thaw the TNT® Quick Master Mix by hand-warming and place on ice. The other components can be thawed at room temperature and then stored on ice.
2. Following the example below, assemble the reaction components in a 0.5ml or 1.5ml microcentrifuge tube. After adding all of the components, gently mix by pipetting. If necessary, centrifuge briefly to return the reaction to the bottom of the tube. For additional information on performing a TNT® Quick reaction, see Notes 1–9 in Section 4.C.
3. We recommend including a control reaction containing no added DNA. This reaction allows measurement of any background incorporation of labeled amino acids.

Examples of TNT® Quick Reactions Using Plasmid DNA

		Labeled		
Components	**Unlabeled**	**[^{35}S]methionine**	**Transcend™ tRNA**	**FluoroTect™ $Green_{Lys}$ tRNA**
TNT® Quick Master Mix (see Note 3, Section 4.C)	40µl	40µl	40µl	40µl
Methionine, 1mM (mix gently prior to use)	1µl	–	1µl	1µl
[^{35}S]methionine (1,000Ci/mmol at 10mCi/ml) (see Note 4, Section 4.C)	–	2µl	–	–
plasmid DNA template(s) (0.5µg/µl) (see Note 6, Section 4.C)	2µl	2µl	2µl	2µl
Transcend™ Biotin-Lysyl-tRNA (see Note 9, Section 4.C)	–	–	1–2µl	–
FluoroTect™ $Green_{Lys}$ tRNA (see Note 9, Section 4.C)	–	–	–	1–2µl
Nuclease-Free Water to a final volume of	**50µl**	**50µl**	**50µl**	**50µl**

Note: Small-scale reactions may be performed by reducing the recommended volumes proportionally.

4.A. General Protocol for TNT® Quick Coupled Transcription/Translation Reactions Using Plasmid DNA (continued)

4. Incubate the reaction at 30°C for 60–90 minutes.
5. Analyze the results of translation. A protocol for Western Blot analysis is provided in Section 7.A. Procedures for determining radiolabel incorporation (Section 7.B) and SDS-PAGE analysis of translation products (Section 7.C) are provided. If using FluoroTect™ $Green_{Lys}$ tRNA, see Section 7.D; for Transcend™ tRNA reactions, see Section 7.E.

4.B. General Protocol for TNT® T7 Quick Coupled Transcription/Translation Reactions Using PCR-Generated DNA

Materials to Be Supplied by the User

- Nuclease-Free Water (Cat.# P1193)
- Radiolabeled amino acid (for radioactive detection; Note 4, Section 4.C) or Transcend™ tRNA (Cat.# L5061) or Transcend™ Colorimetric (Cat.# L5070) or Chemiluminescent (Cat.# L5080) Translation Detection System (for non-radioactive detection) or FluoroTect™ $Green_{Lys}$ in vitro Translation Labeling System (for fluorescent detection; Cat.# L5001).

1. Remove the reagents from storage at –70°C. Rapidly thaw the TNT® Quick Master Mix by hand-warming and place on ice. The other components can be thawed at room temperature and then stored on ice.
2. Following the example below, assemble the reaction components in a 0.5ml or 1.5ml microcentrifuge tube. After adding all of the components, gently mix by pipetting. If necessary, centrifuge briefly to return the reaction to the bottom of the tube. For additional information on performing a TNT® Quick reaction, see Notes 1–9 in Section 4.C.
3. We recommend including a control reaction containing no added DNA. This reaction allows measurement of any background incorporation of labeled amino acids.

Components	Unlabeled	Labeled		
		[^{35}S]methionine	Transcend™ tRNA	FluoroTect™ $Green_{Lys}$ tRNA
TNT® T7 Quick Master Mix (see Note 3, Section 4.C)	40µl	40µl	40µl	40µl
Methionine, 1mM (mix gently prior to use)	1µl	–	1µl	1µl
[^{35}S]methionine (1,000Ci/mmol at 10mCi/ml) (see Note 4, Section 4.C)	–	2µl	–	–
PCR-generated DNA template(s) (see Note 1, Section 4.C)	2.5–5µl	2.5–5µl	2.5–5µl	2.5–5µl
T7 TNT® PCR Enhancer (see Note 2, Section 4.C)	1µl	1µl	1µl	1µl
Transcend™ Biotin-Lysyl-tRNA (see Note 9, Section 4.C)	–	–	1–2µl	–
FluoroTect™ $Green_{Lys}$ tRNA (see Note 9, Section 4.C)	–	–	–	1–2µl
Nuclease-Free Water to a final volume of	**50µl**	**50µl**	**50µl**	**50µl**

Note: Small-scale reactions may be performed by reducing the recommended volumes proportionally.

4. Incubate the reaction at 30°C for 60–90 minutes.
5. Analyze the results of translation. A protocol for Western Blot analysis is provided in Section 7.A. Procedures for determining radiolabel incorporation (Section 7.B) and SDS-PAGE analysis of translation products (Section 7.C) are provided. If using FluoroTect™ $Green_{Lys}$ tRNA, see Section 7.D; for Transcend™ tRNA reactions, see Section 7.E.

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4.C. Notes

1. PCR-generated templates can be used directly from the amplification reaction. We recommend using 2.5–5µl from the amplification reaction, but up to 7µl can be used in a 50µl reaction. For PCR-generated DNA that has been purified following amplification, we recommend using 100–800ng of the purified product for each reaction.
2. We recommend using 1µl of the T7 TNT® PCR Enhancer in a 50µl reaction to increase transcription/translation when using PCR-generated DNA, linear plasmid or viral-enhanced plasmids.
3. The TNT® Quick Master Mix is designed to give the highest expression for most expression constructs. However, we have observed that certain gene constructs may differ in the Mg^{2+} and K^{+} concentrations required for optimal expression in the coupled reaction. For example, some viral leaders will increase translation efficiency and fidelity if additional magnesium acetate and potassium chloride are added to the TNT® Quick reaction. If using a construct with a viral leader, we suggest adding 1–2µl of the T7 TNT® PCR Enhancer.
4. We recommend using a grade of [^{35}S]methionine, PerkinElmer EasyTag™ L-[^{35}S]methionine (PerkinElmer Cat.# NEG709A), which does not cause the background labeling of the rabbit reticulocyte lysate 42kDa protein. Background labeling of the 42kDa protein can occur using other grades of label (16).

   Between 10–40µCi (1–4µl) of [^{35}S]methionine can be added to the TNT® Quick reactions, depending upon the balance between labeling efficiency and cost. For gene constructs that express well and contain several methionines, the 10µCi level (1µl) is sufficient for adequate detection.
5. Except for the actual transcription/translation incubation, all handling of the TNT® Quick System components should be done at 4°C or on ice. Optimum results are obtained when any unused Master Mix is quick-frozen with liquid nitrogen as soon as possible after thawing to minimize loss of translational activity.
6. For most plasmid constructs, optimal results are obtained when 1µg of plasmid DNA template is used. We recommend using 0.2–2.0µg of plasmid DNA in TNT® Quick reactions. The use of more than 1µg of plasmid does not necessarily increase the amount of protein produced.
7. Avoid adding calcium to the transcription/translation reaction. Calcium may reactivate the micrococcal nuclease used to destroy endogenous RNA in the Master Mix and result in degradation of DNA or RNA templates.
8. The TNT® Quick Master Mix contains roughly 100–200mg/ml of endogenous protein.
9. The level of added Transcend™ tRNA and FluoroTect™ $Green_{Lys}$ tRNA can be increased (1–4µl) to allow more sensitive detection of proteins that contain few lysines or are poorly expressed.

5. Positive Control Translation Reactions Using Luciferase

The assay for firefly luciferase activity is extremely sensitive, rapid and easy to perform. It is a good control for in vitro translations because only full-length luciferase is active. Additionally, luciferase is a monomeric protein (61kDa) that does not require post-translational processing or modification for enzymatic activity. The Luciferase Assay System is a substantial improvement over conventional methods in both sensitivity and simplicity (17).

The control reaction can be performed with or without the addition of radiolabeled amino acids.

5.A. Radioactive Luciferase Control Reaction

1. The following example contains [^{35}S]methionine:

TNT® Quick Master Mix (see Note 3, Section 4.C)	40µl
[^{35}S]methionine (1,000Ci/mmol at 10mCi/ml) (see Note 4, Section 4.C)	2µl
Appropriate Luciferase Control DNA (0.5µg/µl) (see Section 11.B)	2µl
Nuclease-Free Water to a final volume of	50µl

2. Incubate the reaction at 30°C for 60–90 minutes (see Note 3, Section 4.C).
3. Analyze the results of translation by measuring direct incorporation of radiolabel (Section 7.B) and/or gel analysis of translation products (Section 7.C).
4. The Luciferase Control reactions can be stored at –20°C for up to 2 months or at –70°C for up to 6 months with little loss of luciferase activity.

5.B. Non-Radioactive Luciferase Control Reaction

1. The following example contains Methionine:

TNT® Quick Master Mix (see Note 3, Section 4.C)	40µl
Methionine, 1mM	1µl
Appropriate Luciferase Control DNA (0.5µg/µl) (see Section 11.B)	2µl
Nuclease-Free Water to a final volume of	50µl

2. Incubate the translation reaction at 30°C for 60–90 minutes.
3. Test for the synthesis of functional luciferase using the standard luciferase assay (see Section 8.A).
4. The Luciferase Control reactions can be stored at –20°C for up to 2 months or at –70°C for up to 6 months with little loss of luciferase activity.

6. Cotranslational Processing Using Canine Pancreatic Microsomal Membranes

Microsomal vesicles are used to study cotranslational and initial post-translational processing of proteins. Processing events such as signal peptide cleavage, membrane insertion, translocation and core glycosylation can be examined by the translation of the appropriate gene in vitro in the presence of these membranes. To ensure consistent performance with minimal background, Canine Pancreatic Microsomal Membranes (Cat.# Y4041) have been isolated so that they are free from mRNA.

For assistance in troubleshooting Microsomal Membrane translation reactions, contact Promega Technical Services. E-mail: **techserv@promega.com**

6.A. General Protocol for Translation with Microsomal Membranes

Materials to Be Supplied by the User

- Canine Pancreatic Microsomal Membranes (Cat.# Y4041)
- [^{35}S]methionine (1,000Ci/mmol at 10mCi/ml)

1. Remove the reagents from the freezer and allow them to thaw on ice.

   Note: The storage buffer for Canine Pancreatic Microsomal Membranes is 50mM triethanolamine, 2mM DTT and 250mM sucrose.

2. Mix the following components on ice, in the order given, in a sterile 1.5ml microcentrifuge tube:

T7 TNT® Quick Master Mix	20µl
[^{35}S]methionine (1,000Ci/mmol at 10mCi/ml) (see Note 4, Section 4.C)	2.0µl
plasmid DNA, 0.5µg/µl	0.5µl
Canine Pancreatic Microsomal Membranes (see Note 1, below)	0.3–1.8µl
Nuclease-Free Water to a final volume of	25µl

3. Incubate at 30°C for 60–90 minutes.

4. Analyze the results of translation and processing. Procedures for Western Blot analysis (Section 7.A), incorporation assays (Section 7.B) and SDS-PAGE analysis of translation products (Section 7.C) are provided.

   Note: TNT® Quick Coupled Transcription/Translation Systems are not tested for performance with Canine Microsomal Membranes.

Notes:

1. We **do not** recommend using Canine Microsomal Membranes when using SP6 TNT® Quick Coupled Transcription/Translation Systems, because SP6 polymerase is sensitive to salts. Transcription may be inhibited as much as 70% by the presence of Canine Microsomal Membranes in the reaction.
2. The amount of Canine Microsomal Membranes used in the reaction may need to be titrated. While these reaction conditions will be suitable for most applications, the efficiency of processing using membranes may vary. Thus, reaction parameters may need to be altered to suit individual requirements. In general, increasing the amount of membranes in the reaction increases the proportion of polypeptides processed but reduces the total amount of polypeptides synthesized.
3. For reactions using the TNT® Quick Coupled Transcription/Translation System, the Canine Microsomal Membranes will inhibit transcription. We do not recommend exceeding 1.8µl of Canine Microsomal Membranes. Transcription/Translation may be inhibited by as much as 50% with 0.6µl of Canine Microsomal Membranes.
4. The amount of protein produced in TNT® Quick reactions using Canine Pancreatic Microsomal Membranes will be less than the amount produced in TNT® Quick reactions alone. Depending on the construct used, protein synthesis efficiency can be expected to drop between 10–50% in the presence of Microsomal Membranes.
5. In some cases, it is difficult to determine if efficient processing or glycosylation has occurred by gel analysis alone. Other assays, such as various protection assays (18), may be required to determine if processing events have taken place.

7. Post-Translational Analysis

Materials to Be Supplied by the User

(Solution compositions are provided in Section 11.A.)

- 1M NaOH/2% H_2O_2
- 25% TCA/2% casamino acids (Difco brand, Vitamin Assay Grade)
- 5% TCA
- Whatman GF/A glass fiber filter (Whatman Cat.# 1820 021)
- acetone
- Whatman 3MM filter paper
- 30% acrylamide solution
- separating gel 4X buffer
- stacking gel 4X buffer
- SDS sample buffer
- SDS polyacrylamide gels
- 50mM DTT
- Blot-Qualified BSA (Cat.# W3841)
- PVDF membrane
- iBlot
- SP-antibody
- TBST buffer

7.A. Western Blot Analysis

1. Add 1µl of the standard, unlabeled translation reaction to 19µl of 1X SDS loading dye with 50mM DTT.
   Note: Include a no template control on the gel to identify background bands.
2. Incubate at 95°C for 5 minutes. Centrifuge briefly to collect the contents in the bottom of the tube.
3. Load 20µl onto a 4–20% gradient Tris-glycine SDS polyacrylamide gel.
4. Following electrophoresis, remove the gel and place it in water.
5. Transfer the proteins to a PVDF membrane using a Western blotting system (e.g., iBlot® System; Invitrogen Cat.# IB1001).
6. Block the membrane using 15ml of 5% Blot-Qualified BSA in TBST (1X TBS + 0.1% Tween® 20). Incubate for 1 hour with gentle shaking.
7. Dilute your primary antibody in 1X TBST.
   Note: We recommend that you titrate your primary antibody dilutions to determine what dilution produces the best results for your protein.
8. Following incubation, remove the blocking solution from the membrane, and add 15ml of diluted primary antibody.
9. Incubate the membrane with the primary antibody at room temperature for 1 hour with gentle shaking.
10. Remove the primary antibody solution, and wash the membrane with 15ml of 1X TBST for 5 minutes with gentle shaking.
11. Repeat the wash 5 more times for a total of six washes.
12. Dilute your secondary antibody 1:2,500 in 1X TBST.
13. Following that last wash, remove buffer from the membrane and add 15ml of diluted secondary antibody.
14. Incubate the membrane with the secondary antibody for 1 hour with gentle shaking.
15. Following the incubation, remove the secondary antibody solution, and wash the membrane with 15ml of 1X TBST for five minutes. Repeat for a total of six washes.
16. Proceed to the detection method appropriate for your secondary antibody. If using FluoroTect™ $Green_{Lys}$ tRNA, see Section 7.D; for Transcend™ tRNA reactions, see Section 7.E.

7.B. Determination of Percent Incorporation of Radioactive Label

1. After the 50µl translation reaction is complete, remove 2µl from the reaction, and add it to 98µl of 1M NaOH/ 2% H_2O_2.
2. Vortex briefly and incubate at 37°C for 10 minutes.
3. At the end of the incubation, add 900µl of ice-cold 25% TCA/2% casamino acids to precipitate the translation product. Incubate on ice for 30 minutes.
4. Wet a Whatman GF/A glass fiber filter with a small amount of ice-cold 5% TCA. Collect the precipitated translation product by vacuum filtering 250µl of the TCA reaction mix. Rinse the filter 3 times with 1–3ml of ice-cold 5% TCA. Rinse once with 1–3ml of acetone. Allow the filter to dry at room temperature or under a heat lamp for at least 10 minutes.
5. For determination of ^{35}S incorporation, put the filter in the appropriate scintillation cocktail, invert to mix and count in a liquid scintillation counter.
6. To determine total counts present in the reaction, spot a 5µl aliquot of the TCA reaction mix directly onto a filter. Dry the filter for 10 minutes. Count in a liquid scintillation counter as in Step 5.
7. To determine background counts, remove 2µl from a 50µl translation reaction containing no DNA and proceed as described in Steps 1–5.
8. Perform the following calculation to determine percent incorporation:

$$\frac{\text{cpm of washed filter (Step 5)}}{\text{cpm of unwashed filter (Step 6)} \times 50} \times 100 = \text{percent incorporation}$$

9. Perform the following calculation to determine the fold stimulation over background:

$$\frac{\text{cpm of washed filter (Step 5)}}{\text{cpm of ``no DNA control reaction'' filter (Step 7)}} = \text{fold stimulation}$$

7.C. Denaturing Gel Analysis of Radioactively Labeled Translation Products

Precast polyacrylamide gels are available from a number of manufacturers. For protein analysis, Invitrogen Corporation and Bio-Rad Laboratories, Inc., offer a variety of precast mini-gels, which are compatible with their vertical electro-phoresis and blotter systems. These companies offer Tris-Glycine, Tricine and Bis-Tris gels for resolution of proteins under different conditions and over a broad spectrum of protein sizes. The Invitrogen Novex® 4–20% Tris-Glycine gradient gels (Cat.# EC6025BOX or EC60355BOX) and the Bio-Rad Ready Gel® 4–20% Tris-Glycine Gel, 10-well (Cat.# 161-0903) are convenient for resolving proteins over a wide range of molecular weights. In addition to convenience and safety, precast gels provide consistent results.

1. Once the 50µl translation reaction is complete (or at any desired timepoint), remove a 1–5µl aliquot, and add it to 20µl of SDS sample buffer. The remainder of the reaction may be stored at –20°C, or at –70°C for long-term storage.

7.C. Denaturing Gel Analysis of Radioactively Labeled Translation Products (continued)

2. Cap the tube and heat at 100°C for 2 minutes to denature the proteins. This may cause protein aggregation. Incubation at a lower temperature (e.g., 20 minutes at 60°C, 10 minutes at 70°C or 3–4 minutes at 80–85°C) may be more appropriate.
3. A small aliquot (5–10µl) of the denatured sample can then be loaded onto an SDS-polyacrylamide gel or stored at –20°C. It is not necessary to separate labeled polypeptides from free amino acids by acetone precipitation.
4. Typically, electrophoresis is carried out at a constant current of 15mA in the stacking gel and 30mA in the separating gel (or 30mA for a gradient gel). Electrophoresis is usually performed until the bromophenol blue dye has run off the bottom of the gel. Disposal of unincorporated label may be easier if the gel is stopped while the dye front remains in the gel, as the dye front also contains the unincorporated labeled amino acids. If transferring the gel to a membrane filter for Western blotting, proceed to Step 7.
5. Place the polyacrylamide gel in a plastic box, and cover the gel with fixing solution (as prepared in Section 11.A) for 30 minutes. Agitate slowly on an orbital shaker. Pour off the fixing solution. Proceed to Step 6 (gel drying prior to film exposure).

   Optional: Labeled protein bands in gels may be visualized by autoradiography or fluorography. Fluorography dramatically increases the sensitivity of detection of ^{35}S-, ^{14}C- and ^{3}H-labeled proteins and is recommended for the analysis of in vitro translation products. The increased detection sensitivity of fluorography is obtained by infusing an organic scintillant into the gel. The scintillant converts the emitted energy of the isotope to visible light and increases the proportion of energy that may be detected by X-ray film. Commercial reagents, such as Amplify® Reagent (GE Healthcare Bio-sciences), can be used for fluorographic enhancement of signal. Alternatively, the fixed gel can be exposed to a phosphorimaging screen. These systems provide greater sensitivity, greater speed and the ability to quantitate the radioactive bands.
6. Dry the gel before exposure to film as follows: Soak the gel in 10% glycerol for 5 minutes to prevent the gel from cracking during drying. Place the gel on a sheet of Whatman 3MM filter paper, cover with plastic wrap and dry at 80°C for 30–90 minutes under a vacuum using a conventional gel dryer; dry completely. The gel also may be dried overnight using the Gel Drying Kit (Cat.# V7120). To decrease the likelihood of cracking gradient gels, dry them with the wells pointing down. Expose the gel on Kodak X-OMAT® AR film for 1–6 hours at –70°C (with fluorography) or 6–15 hours at room temperature (with autoradiography).
7. For Western blot analysis of proteins, transfer (immobilize) the protein from the gel onto nitrocellulose or PVDF membrane (19,20). Usually Western blots are made by electrophoretic transfer of proteins from SDS-polyacrylamide gels. Detailed procedures for electrophoretic blotting usually are included with commercial devices and can be found in references 19, 21, 22 and 23. A general discussion of Western blotting with PVDF membranes is found in reference 24. PVDF membranes must be prewet in methanol or ethanol before equilibrating in transfer buffer. The blot may then be subjected to immunodetection analysis. For more information, refer to the Promega *Protocols and Applications Guide*, Online Edition (25).

7.D. Denaturing Gel Analysis of Translation Products Labeled with the FluoroTect™ Green$_{Lys}$ in vitro Translation Labeling System

The fluorescent translation product should be resolved by SDS-PAGE and then visualized by placing the gel on a laser-based fluorescence scanning device.

Note: The use of gel systems other than Tris-Glycine may cause different migration patterns for the expressed and background bands.

Denaturing Gel Analysis

1. Once the translation reaction is complete (or at any desired time point), remove a 5µl aliquot, and add it to 20µl of 1X SDS gel-loading buffer. Store the remainder of the translation reaction at –20°C. The FluoroTect™ tRNA fluorophore is sensitive to extreme heating. If heating to denature the proteins, do not exceed 70°C for more than 2–3 minutes.
2. Load the sample from Step 1 on an SDS-PAGE gel.
3. Perform electrophoresis using standard conditions for your apparatus. Typically, electrophoresis is carried out at a constant current of 20mA. Electrophoresis usually is performed until the bromophenol blue dye has run to the bottom of the gel.

Fluorescent Detection

Materials to Be Supplied by the User

- Fluorescent Imaging Instrument (i.e., FluorImager® SI or FluorImager® 595 [Molecular Dynamics], both with a 499 argon laser; the Typhoon® 8600 [Molecular Dynamics], with a 532nm excitation, or the FMBIO® II [Hitachi], with a 505 channel)

Note: The Storm® instrument (Molecular Dynamics) is not recommended for use with the FluoroTect™ System due to reduced sensitivity.

After electrophoresis is completed, immediately place the gel in water, then complete fluorescent scanning.

Use gloves when handling the gels.

Notes:

1. Fixing polyacrylamide gels does not interfere with the detection of FluoroTect™ Green$_{Lys}$-labeled in vitro **translation products**, although the signal intensity may be somewhat decreased.
2. Drying fixed polyacrylamide gels in cellophane does not interfere with the detection of FluoroTect™ Green$_{Lys}$-labeled in vitro translation products, although signal intensity may be somewhat decreased.
3. Fixing and/or drying gels may decrease the signal intensity of **prestained molecular weight markers**, making them difficult to detect with fluorescent scanners.

7.E. Denaturing Gel Analysis of Translation Products Labeled with the Transcend™ Non-Radioactive Translation Detection Systems

Biotinylated protein standards (Bio-Rad Cat.# 161-0319) can be used to determine the apparent molecular weight of the translated biotinylated protein. Alternatively, fluorescently labeled size standards can be observed after transfer and marked with a pencil under UV irradiation. The positions of unlabeled size standards also can be determined by staining the blot after transfer (see *Transcend™ Non-Radioactive Translation Detection Systems Technical Bulletin* #TB182).

1. Once the 50µl translation reaction is complete (or at any desired time point), remove a 1µl aliquot and add it to 15µl of SDS sample buffer. The remainder of the reaction may be stored at –20°C.
2. Close the tube and heat at 90–100°C for 2 minutes to denature the proteins.

   Note: In some cases, high molecular weight complexes are formed at 100°C, and denaturation may need to be performed at lower temperatures (e.g., 20 minutes at 60°C, 10 minutes at 70°C or 3–4 minutes at 80–85°C).
3. Load the denatured sample on an SDS-polyacrylamide gel. (Protocols for SDS polyacrylamide gel electrophoresis may be found in the *Protocols and Applications Guide,* Online Edition [25]).
4. Perform electrophoresis using standard conditions for your apparatus. Typically, electrophoresis is carried out at a constant current of 20mA. Electrophoresis usually is performed until the bromophenol blue dye has run off the bottom of the gel.

   Note: If a gene product is weakly expressed or contains few lysines, up to 2µl of the translation reaction (Reticulocyte Lysate) can be loaded on an SDS gel without the loss of resolution observed with autoradiography. However, loading more of the translation reaction can result in high background on the blot.

Electroblotting of Proteins to Membrane

For colorimetric detection, see Section 5.C of the *Transcend™ Non-Radioactive Translation Detection Systems Technical Bulletin* #TB182. The translation products can be blotted from the SDS-polyacrylamide gel to (in decreasing order of preference) PVDF, nitrocellulose or another membrane using any standard apparatus and protocol, including semi-dry systems. Detailed procedures for electrophoretic blotting are usually included with commercial devices. We routinely transfer at a constant voltage of 100V for 60 minutes using a minigel-size electroblotting unit or 15 minutes using a semi-dry system. PVDF membrane must be pre-wet in methanol before it is equilibrated in transfer buffer.

Instructions for chemiluminescent detection of products are found in Section 5.D of the *Transcend™ Non-Radioactive Translation Detection Systems Technical Bulletin* #TB182.

8. Positive Control Luciferase Assays

Light intensity is a measure of the rate of catalysis by luciferase and is therefore dependent upon temperature. The optimum temperature for luciferase activity is approximately room temperature (20–25°C). **It is important** that the Luciferase Assay Reagent be fully equilibrated to room temperature before beginning measurements. To ensure temperature equilibration, place a thawed aliquot of the Luciferase Assay Reagent in a sealed tube into a water bath maintained at ambient temperature, and equilibrate for at least 30 minutes. The sample to be assayed should also be at ambient temperature.

Either a luminometer or a scintillation counter can be used for quantitation. (There is usually insufficient light output for qualitative visual detection.) A luminometer can measure as little as 10^{-20} moles (0.001pg) of luciferase, whereas a scintillation counter typically has a less sensitive detection limit. However, the limits of sensitivity may vary depending upon the particular instrument used. The assay should be linear in some portion of the detection range of the instrument. Please consult your instrument operator's manual for general operating instructions.

8.A. Using a Luminometer

1. Dispense 50µl of the Luciferase Assay Reagent into luminometer tubes, one tube per sample.
2. Program the luminometer to perform a 2-second measurement delay followed by a 10-second measurement read for luciferase activity. The read time may be shortened if sufficient light is produced.
3. Add 2.5µl of cell lysate to a luminometer tube containing the Luciferase Assay Reagent. Mix by pipetting 2–3 times or vortex briefly.
4. Place the tube in the luminometer and initiate reading.
5. If the luminometer is not connected to a printer or computer, record the reading.

8.B Using a Scintillation Counter

Ideally, the coincidence circuit of the scintillation counter should be turned off. Usually, this is achieved through an option of the programming menu or by a switch within the instrument. Consult the user's manual or the manufacturer of the scintillation counter. If the circuit cannot be turned off, a linear relationship between luciferase concentration and cpm still can be produced by calculating the square root of measured counts per minute (cpm) minus background cpm (i.e., $[\text{sample} - \text{background}]^{1/2}$). To measure background cpm, use water or Luciferase Assay Reagent as a blank.

Use the same protocol as luciferase assays using a luminometer (Section 7.B). The sample may be placed directly in the scintillation vial if it completely covers the bottom of the vial (clear or translucent vials are acceptable). **Do not** add scintillant because it will inactivate luciferase. Alternatively, place the sample in a microcentrifuge tube, and then place the tube in the scintillation vial. To ensure consistency when working with multiple samples, place each microcentrifuge tube at the same relative position within the scintillation vial.

For consistency in measuring luciferase activity, use the scintillation counter in manual mode. Initiate each sample reaction immediately before measurement, and read the samples one at a time. Because the enzymatic reaction produces light at all wavelengths, read the samples with all channels open (open window). To reduce background counts, it may be necessary to wait 10–30 seconds before counting. Read individual samples for 1–5 minutes.

9. Troubleshooting

For questions not addressed here, please contact your local Promega Branch Office or Distributor. Contact information available at: **www.promega.com.** E-mail: **techserv@promega.com**

Symptoms	Causes and Comments
The control reaction produces no luciferase	Loss of reaction component(s) activity. The lysate should not be used after more than two freeze-thaw cycles. Do not use reagents after the expiration date.
	Ethanol or salt present in the reaction may inhibit translation.
Low translation efficiency	Certain gene constructs may require different Mg^{2+} and K^{+} concentrations for optimal expression. Add 1–3µl of the T7 TNT® PCR Enhancer.
	Calcium is present in the translation reaction. Avoid adding calcium to the translation reaction. Calcium may reactivate the micrococcal nuclease used used to destroy endogenous mRNA in the lysate and result in degradation of the DNA or mRNA template.
	Ethanol present in the translation reaction. Residual ethanol should be removed from template DNA preparations and amino acids before they are added to the translation reaction.
	Incubation of the reaction at 37°C causes decreased protein synthesis. Incubate the reaction at 30°C, the optimal temperature.
Unexpected bands present at higher molecular weights or bands stuck in stacking gel	Denaturing temperature too high. Denature sample at a lower temperature (e.g., 60–80°C) for 10–15 minutes.
Unexpected bands present on the gel	Proteolysis of translation product. Add protease inhibitors, such as α2-macroglobulin, leupeptin or chymostatin (0.5–1µg/ml).
	More than one peptide is translated from the template. Leaky scanning for translation initiation can result in translation initiating at internal methionines. Optimizing the Mg^{2+} or K^{+} concentration can increase fidelity (26).
	The [^{35}S]methionine used is not translational grade or beyond its expiration date. There are reports of a 42kDa band with some grades of [^{35}S]methionine (16). We recommend Perkin-Elmer EasyTag™ L-[^{35}S]methionine (Perkin-Elmer Cat.# NEG709A) to avoid this 42kDa band.
	Globin may appear on the autoradiogram or stained gel. It appears as a broad band migrating at 10–15kDa.

Symptoms	Causes and Comments
Unexpected bands present on the gel (continued)	Aminoacyl tRNAs may produce background bands (~25kDa). Add RNase A to the lysate reaction (after completion) to a final concentration of 0.2mg/ml. Incubate for 5 minutes at 30°C.
	Oxidized β-mercaptoethanol is present or not enough SDS in the loading buffer. Use a loading buffer that contains 2% SDS and 100mM DTT.
Unexpected bands present when isolating polyhistidine-tagged protein	A nickel-based resin is used to purify polyhistidine-tagged proteins. Hemoglobin present in the rabbit reticulocyte lysate will bind to the nickel and co-elute with the polyhistidine-tagged protein. Use the MagZ™ Protein Purification System (Cat.# V8830) or an alternate purification tag to isolate the protein from the TNT® lysate and avoid this problem.
Smearing on the gel	Gel not clean. Gel must be washed before placing onto film. Once gel electrophoresis is complete, soak the gel in either a standard Coomassie® destaining solution (50% methanol, 7.5% glacial acetic acid) or in water for 15–30 minutes prior to drying.
	Too much protein loaded on the gel. Check the amount of sample loaded on the gel and the amount of loading buffer. Too much protein loaded can cause smearing.
	Acrylamide concentration in the gel is too low. Acrylamide concentration can be increased to 12%.
	Sample contains ethanol, which can cause gel smearing.
High background levels when performing Western Blots.	Primary antibody concentration too high.Increase the dilution of the primary antibody.

10. References

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11. Appendix

11.A. Composition of Buffers and Solutions

acrylamide solution, 30% (37.5:1)

30g	acrylamide
0.8g	bisacrylamide

Add water to a final volume of 100ml. Store at 4°C.

fixing solution

50%	methanol
10%	glacial acetic acid
40%	water

1X SDS gel-loading buffer

50mM	Tris-HCl (pH 6.8)
100mM	dithiothreitol
2%	SDS
0.1%	bromophenol blue
10%	glycerol

1X SDS gel-loading buffer lacking dithiothreitol can be stored at room temperature. **Dithiothreitol should be added from a 1M stock just before the buffer is used.**

SDS polyacrylamide running 10X buffer

30g	Tris base
144g	glycine
100ml	10% SDS

Add deionized water to a final volume of 1 liter. Store at room temperature.

separating gel 4X buffer

18.17g	Tris base
4ml	10% SDS

Bring the volume to approximately 80ml with deionized water. Adjust to pH 8.8 with 12N HCl and add deionized water to a final volume of 100ml. Store at room temperature.

stacking gel 4X buffer

6.06g	Tris base
4ml	10% SDS

Bring the volume to approximately 80ml with deionized water. Adjust to pH 6.8 with 12N HCl and add deionized water to a final volume of 100ml. Store at room temperature.

T7 TNT® PCR Enhancer

0.5M	KCl
12.5mM	$Mg(OAc)_2$

In nanopure water.

TBST

50mM	Tris HCl (pH7.4)
150mM	NaCl
0.1%	Tween®-20

11.B. Luciferase SP6/T7 Control DNAs

The Luciferase SP6/T7 Control DNAs are used as functional controls in the TNT® Quick Coupled Transcription/ Translation System. The Control DNAs contain the gene for luciferase under transcriptional control of a phage RNA polymerase promoter. The constructs carry a 30bp poly[d(A)/d(T)] tail following the luciferase gene. The maps of the Luciferase SP6 Control DNA and T7 Control DNA are shown in Figures 3 and 4, respectively. Please note that these vectors are intended for use as control luciferase expression vectors only. They are not intended for use as cloning vectors.

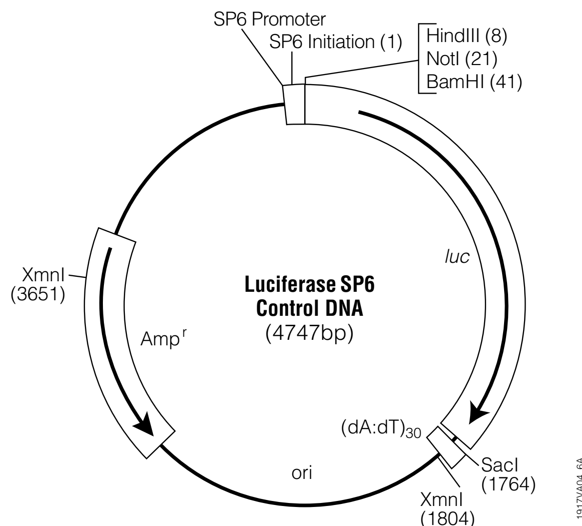

Figure 3. Luciferase SP6 Control DNA circle map and sequence reference points. Additional description: Amp^r, β-lactamase gene (resistant to ampicillin); ori, origin of plasmid replication.

Sequence reference points:

SP6 RNA polymerase initiation	1
GLPrimer2	49–71
Luciferase gene	48–1700
Poly(A) $(dA)_{30}$	1767–1796
pUC/M13 reverse primer (17mer)	1833–1817
pUC/M13 reverse primer (22mer)	1838–1817
β-lactamase gene (Amp^r)	3838–2978
SP6 RNA polymerase promoter primer	4731–1
SP6 RNA polymerase promoter	4731–3

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11.B. Luciferase SP6/T7 Control DNAs (continued)

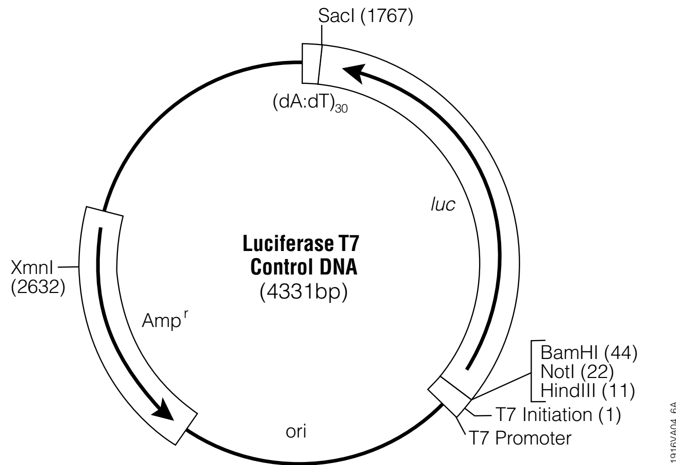

Figure 4. Luciferase T7 Control DNA circle map and sequence reference points. Additional description: Amp^r, β-lactamase gene (resistant to ampicillin); ori, origin of plasmid replication.

Sequence reference points:

T7 RNA polymerase initiation	1
GLPrimer2	52–74
Luciferase gene	51–1700
Poly(A) $(dA)_{30}$	1770–1799
β-lactamase gene (Amp^r)	2444–3304
T7 RNA polymerase promoter	4315–3
T7 RNA polymerase promoter primer	4315–3

11.C. Related Products

The in vitro synthesis of proteins is a popular method in biological research. Among other applications, translation systems are used to rapidly characterize plasmid clones, study structural mutations and examine translational signals.

Two basic approaches to in vitro protein synthesis are available: 1) systems programmed with RNA (translation systems), or 2) systems programmed with DNA (coupled transcription/translation systems). Several general considerations to assist you in selecting the appropriate Promega product(s) are discussed in this section.

Translation Systems

A number of cell-free translation systems have been developed for the translation of mRNA isolated from tissue or generated in vitro. Promega offers several Rabbit Reticulocyte Lysate and Wheat Germ Extract Systems. All are reliable, convenient and easy-to-use systems to initiate translation and produce full-size polypeptide products. Rabbit Reticulocyte Lysate is appropriate for the translation of larger mRNA species and generally is recommended when microsomal membranes are to be added for cotranslational processing of translation products. Flexi® Rabbit Reticulocyte Lysate is recommended where optimization of translation of particular RNAs through adjustments to salt and DTT concentrations is required. Wheat Germ Extract is recommended for translation of RNA preparations containing low concentrations of double-stranded RNA (dsRNA) or oxidized thiols, which are inhibitory to reticulocyte lysate.

Coupled Transcription/Translation Systems

DNA sequences cloned in plasmid vectors also may be expressed directly using either TNT® Coupled Reticulocyte Lysate Systems, Wheat Germ Extract Systems or *E. coli* S30 Extract Systems. The TNT® Systems are used to direct eukaryotic translation, whereas the S30 Systems are under prokaryotic translational controls. The TNT® Systems require plasmid constructs containing a phage RNA polymerase promoter (SP6, T3 or T7) for the initiation of transcription, but translation in this system is under eukaryotic controls. Optimal translation will occur if the AUG initiation codon is in a "Kozak consensus" context (A/GCCAUGG) (27) in the absence of inhibiting secondary structure. The template DNA to be expressed in the S30 Systems must contain *E. coli* promoter sequences or a phage T7 promoter sequence and prokaryotic ribosome binding sites (GGAGG) for translation. The TNT® and *E. coli* S30 Systems can use either circular or linear DNA templates.

TNT® Coupled Reticulocyte Lysate Systems

Product	Size	Cat.#
TNT® SP6 Coupled Reticulocyte Lysate System	40 reactions	L4600
TNT® T7 Coupled Reticulocyte Lysate System	40 reactions	L4610
TNT® T3 Coupled Reticulocyte Lysate System	40 reactions	L4950
TNT® T7/T3 Coupled Reticulocyte Lysate System	40 reactions	L5010
TNT® T7/SP6 Coupled Reticulocyte Lysate System	40 reactions	L5020
TNT® T7 Quick for PCR DNA	40 reactions	L5540
TNT® SP6 Coupled Reticulocyte Lysate System, Trial Size	8 reactions	L4601
TNT® T7 Coupled Reticulocyte Lysate System, Trial Size	8 reactions	L4611

11.C. Related Products (continued)

TnT® Coupled Wheat Germ Extract Systems

Product	Size	Cat.#
TnT® SP6 High-Yield Wheat Germ Protein Expression System	4 × 300µl	L3260
	1 × 300µl	L3261
TnT® SP6 Coupled Wheat Germ Extract System	40 reactions	L4130
TnT® T7 Coupled Wheat Germ Extract System	40 reactions	L4140
TnT® T7/SP6 Coupled Wheat Germ Extract System	40 reactions	L5030

Rabbit Reticulocyte Lysate

Product	Size	Cat.#
Rabbit Reticulocyte Lysate, Nuclease-Treated	5 × 200µl	L4960
Rabbit Reticulocyte Lysate, Untreated	1ml	L4151

Bulk Rabbit Reticulocyte Lysate is available from Promega.

Flexi® Rabbit Reticulocyte Lysate System

Product	Size	Cat.#
Flexi® Rabbit Reticulocyte Lysate System	5 × 200µl	L4540

Bulk Flexi® Rabbit Reticulocyte Lysate is available from Promega.

Wheat Germ Extract

Product	Size	Cat.#
Wheat Germ Extract	5 × 200µl	L4380

Rabbit Reticulocyte Lysate/Wheat Germ Extract Combination System

Product	Size	Cat.#
Rabbit Reticulocyte/Wheat Germ Extract Combination System	12 reactions each	L4330

E. coli S30 Extract Systems

Product	Size	Cat.#
E. coli S30 Extract System for Linear Templates	30 reactions	L1030
E. coli S30 Extract System for Circular DNA	30 reactions	L1020
E. coli T7 S30 Extract System for Circular DNA	30 reactions	L1130

Transcend™ Non-Radioactive Translation Detection Systems

Product	Size	Cat.#
Transcend™ Colorimetric Translation Detection System	30 reactions	L5070
Transcend™ Chemiluminescent Translation Detection System	30 reactions	L5080
Transcend™ Biotinylated tRNA	30µl	L5061

FluoroTect™ Green$_{Lys}$ in vitro Translation Labeling System

Product	Size	Cat.#
FluoroTect™ Green$_{Lys}$ in vitro Translation Labeling System	40 reactions	L5001

Canine Pancreatic Microsomal Membranes

Product	Size	Cat.#
Canine Pancreatic Microsomal Membranes	50µl	Y4041

Protein:Protein Interactions

Product	Size	Cat.#
MagneGST™ Pull-Down System	80 reactions	V8870

Plasmid Purification

Product	Size	Cat.#
PureYield™ Plasmid Midiprep System	25 preps	A2492
	100 preps	A2495

12. Summary of Change

The following change was made to the 5/17 revision of this document:

1. Removed expired patent statements.

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