T7 Express Competent E. coli (High Efficiency)

C2566H

20 x 0.05 ml/tube Lot: 0221408
Store at –80°C

CAUTION: This product contains DMSO, a hazardous material. Review the MSDS before handling.

Description: Chemically competent E. coli cells suitable for high efficiency transformation and protein expression.

Features:
- Transformation efficiency: 0.6–1 x 10^9 cfu/µg pUC19
- Enhanced BL21 derivative for T7 expression
- T7 RNA Polymerase in the lac operon - no lambda prophage
- Deficient in proteases Lon and OmpT
- Resistant to phage T1
- Resistant to methylated DNA (McrA–, McrBC–, EcoBr–m–, Mrr–)
- Does not restrict methylated DNA (McrA, McrBC, EcoBr m, Mrr)
- B Strain
- Free of animal products

Reagents Supplied:
- 20 x 0.05 ml/tube of chemically competent T7 Express Competent E. coli cells (Store at –80°C)
- 20 ml of SOC Outgrowth Medium (Store at room temperature)
- 0.025 ml of 50 pg/µl pUC19 Control DNA (Store at –20°C)

Quality Control Assays
Transformation Efficiency: 100 pg of pUC19 plasmid DNA was used to transform one tube of T7 Express Competent E. coli following the high efficiency protocol provided. 0.6–1 x 10^9 colonies formed/µg after an overnight incubation on LB-ampicillin plates at 37°C.
Untransformed cells were also tested for resistance to phage ø80, a standard test for resistance to phage T1, and sensitivity to ampicillin, chloramphenicol, kanamycin, spectinomycin, streptomycin and tetracycline.

High Efficiency Transformation Protocol
Perform steps 1–7 in the tube provided.
1. Thaw a tube of T7 Express Competent E. coli cells on ice for 10 minutes.
2. Add 1–5 µl containing 1 pg–100 ng of plasmid DNA to the cell mixture. Carefully flick the tube 4–5 times to mix cells and DNA. Do not vortex.
3. Place the mixture on ice for 30 minutes. Do not mix.
5. Place on ice for 5 minutes. Do not mix.
6. Pipette 950 µl of room temperature SOC into the mixture.
7. Place at 37°C for 60 minutes. Shake vigorously (250 rpm) or rotate.
8. Warm selection plates to 37°C.
9. Mix the cells thoroughly by flicking the tube and inverting, then perform several 10-fold serial dilutions in SOC.
10. Spread 50–100 µl of each dilution onto a selection plate and incubate overnight at 37°C. Alternatively, incubate at 30°C for 24–36 hours or at 25°C for 48 hours.

5 Minute Transformation Protocol
A shortened transformation protocol resulting in approximately 10% efficiency compared to the standard protocol may be suitable for applications where a reduced total number of transformants is acceptable. Follow the High Efficiency Transformation Protocol above with the following changes:
1. Steps 3 and 5 are reduced to 2 minutes.
2. Omit outgrowth (step 7) completely for ampicillin-resistant plasmids or reduce the outgrowth time for other selective media as appropriate.

Protocol for Expression Using T7 Express
1. Transform expression plasmid into T7 Express. Plate on antibiotic selection plates and incubate overnight at 37°C.
2. Resuspend a single colony in 10 ml liquid culture with antibiotic.
3. Incubate at 37°C until OD600 reaches 0.4–0.8.
4. Induce with 40 µl of a 100 mM stock of IPTG (final concentration of 0.4 mM) and induce for 2 hours at 37°C.
5. Check for expression either by Coomassie stained protein gel, Western Blot or activity assay. Check expression in both the total cell extract (soluble + insoluble) and the soluble fraction alone.
6. For large scale, inoculate 1 L of liquid medium (with antibiotic) with a freshly grown colony or 10 ml of freshly grown culture. Incubate at 37°C until OD600 reaches 0.4–0.8. Add IPTG to 0.4 mM. Induce 2 hours at 37°C or 15°C overnight.

Transformation Protocol Variables
Thawing: Cells are best thawed on ice and DNA added as soon as the last bit of ice in the tube disappears. Cells can also be thawed by hand, but warming above 0°C will decrease the transformation efficiency.

Incubation of DNA with Cells on Ice: For maximum transformation efficiency, cells and DNA should be incubated together on ice for at least 10 minutes. Expect a 2-fold loss in transformation efficiency for every 15 minutes this step is shortened.

Heat Shock: Both the temperature and the timing of the heat shock step are important and specific to the transformation volume and vessel. Using the transformation tube provided, 10 seconds at 42°C is optimal.

Outgrowth: Outgrowth at 37°C for 1 hour is best for cell recovery and for expression of antibiotic resistance. Expect a 2-fold loss in transformation efficiency for every 15 minutes this step is shortened. SOC gives 2-fold higher transformation efficiency than LB medium; and incubation with shaking or rotating the tube gives 2-fold higher transformation efficiency than incubation without shaking.

Plating: Selection plates can be used warm or cold, wet or dry without significantly affecting the transformation efficiency. However, warm, dry plates are easier to spread and allow for the most rapid colony formation.

Transformation of a toxic mammalian clone into E. coli hosts. A T7 expression plasmid and the same plasmid containing a gene encoding a toxic mammalian protein were transformed into each host. Comparison of the relative transformation efficiencies demonstrates that the T7 Express hosts provide the levels of control necessary for transformation of potentially toxic clones. BL21(DE3) could not be transformed with the toxic clone.

CERTIFICATE OF ANALYSIS
DNA Contaminants to Avoid

<table>
<thead>
<tr>
<th>Contaminant</th>
<th>Removal Method</th>
</tr>
</thead>
<tbody>
<tr>
<td>Detergents</td>
<td>Ethanol precipitate</td>
</tr>
<tr>
<td>Phenol</td>
<td>Extract with chloroform and ethanol precipitate</td>
</tr>
<tr>
<td>Ethanol or isopropanol</td>
<td>Dry pellet before resuspending</td>
</tr>
<tr>
<td>PEG*</td>
<td>Column purity or phenol/chloroform extract and ethanol precipitate</td>
</tr>
<tr>
<td>DNA binding proteins* (e.g. Ligase)</td>
<td>Column purity or phenol/chloroform extract and ethanol precipitate</td>
</tr>
</tbody>
</table>

*Ideally, DNA for transformation should be purified and resuspended in water or TE. However, up to 10 µl of DNA directly from a ligation mix can be used with only a two-fold loss of transformation efficiency. Where it is necessary to maximize the number of transformants (e.g. a library), a purification step, either a spin column or phenol/chloroform extraction and ethanol precipitation should be added.

Troubleshooting T7 Protein Expression

No colonies or no growth in liquid culture: Even though T7 expression is tightly regulated, there may be a low level of basal expression in the T7 Express host. If toxicity of the expressed protein is likely, transformation of the expression plasmid should be carried out in one of the following strains:

- T7 Express E::T7 RNA polymerase
- T7 Express 1ysY::T7 RNA polymerase, reducing basal expression of the target protein. Upon induction, newly made T7 RNA polymerase titrates out the lysozyme and results in expression of the target protein
- T7 Express 1ysY/1ompT combines both above effects.

Incubation at 30°C or room temperature may also alleviate toxicity issues. In addition, check antibiotic concentration (test with control plasmid).

No protein visible on gel or no activity: Check for toxicity - no protein may mean the cells have eliminated or deleted elements in the expression plasmid.

- Culture cells for protein induction. Just before induction, plate a sample on duplicate plates with and without antibiotic selection. If toxicity is an issue, there will be a significant difference between the number of colonies on the plates. Fewer colonies will be seen on plates containing antibiotic (indicating that the plasmid has been lost) compared to plates without antibiotic.
- If toxicity is the problem test the above E and 1ysY hosts to reduce basal level expression.

Induced protein is insoluble: Check for insolubility - this is important because T7 expression often leads to very high production of protein that can result in the target protein becoming insoluble. Potential solutions for this are:

- Induce at lower temperatures (as low as 15°C overnight)
- Reduce IPTG concentration to 0.1 mM and 0.01 mM
- Induce for less time (as little as 15 minutes)
- Induce earlier in growth (OD<sub>600</sub> = 0.3 or 0.4)

Antibiotics for Plasmid Selection

<table>
<thead>
<tr>
<th>Antibiotic</th>
<th>Working Concentration</th>
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<tbody>
<tr>
<td>Ampicillin</td>
<td>100 µg/ml</td>
</tr>
<tr>
<td>Carbenicillin</td>
<td>100 µg/ml</td>
</tr>
<tr>
<td>Chloramphenicol</td>
<td>33 µg/ml</td>
</tr>
<tr>
<td>Kanamycin</td>
<td>30 µg/ml</td>
</tr>
<tr>
<td>Streptomycin</td>
<td>25 µg/ml</td>
</tr>
<tr>
<td>Tetracycline</td>
<td>15 µg/ml</td>
</tr>
</tbody>
</table>

Genotype: tnuA2 lacZ::T7 gene1 [lon] ompT gal sulA11 R(mcr-car73::miniTn10--Te)<sup>2</sup> [dcm] R(zgb-210::MiniTn10--Tet<sup>2</sup> endA1 Δ(mcrC-mrr)<sup>114::IS10</sup>]

Solutions/Recipes

<table>
<thead>
<tr>
<th>SOB:</th>
<th>SOC:</th>
</tr>
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<tbody>
<tr>
<td>2% Vegetable peptone (or Tryptone)</td>
<td>SOB + 20 mM Glucose</td>
</tr>
<tr>
<td>0.5% Yeast Extract</td>
<td>LB agar:</td>
</tr>
<tr>
<td>10 mM NaCl</td>
<td>2.5 mM KCl</td>
</tr>
<tr>
<td>10 mM MgCl₂</td>
<td>1% Tryptone</td>
</tr>
<tr>
<td>10 mM MgSO&lt;sub&gt;4&lt;/sub&gt;</td>
<td>0.5% Yeast Extract</td>
</tr>
<tr>
<td>0.17 M NaCl</td>
<td>1.5% Aga</td>
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Strain Properties

The properties of this strain that contribute to its usefulness as a protein expression strain are described below. The genotypes underlying these properties appear in parentheses.

T7 RNA Polymerase (lacZ::T7 gene1): T7-Express has the T7 RNA polymerase gene inserted into the lac operon on the E. coli chromosome and is expressed under the control of the lac promoter. This configuration provides controlled induction of the polymerase and consequently, inducible control of transcription of genes downstream of the T7 promoter. This system provides potential advantages over strains such as BL21(DE3), that carry the T7 RNA polymerase on a lysogenic prophage. Although DE3 is normally dormant in the host chromosome, the induction of the SOS cascade can occur as the result of expressing proteins that damage the E. coli chromosome, either directly or indirectly. This may lead to cell lysis.

Protease Deficient (lon ompT): E. coli B strains are "naturally" deficient in the lon protease which in K-12 strains serves to degrade misfolded proteins and to prevent some cell cycle-specific proteins from accumulating. The OmpT protease resides at the surface of wild type E. coli in both K-12 and B strains, presumably helping the cells to derive amino acids from their external environment. Cells deficient in both these proteases are much more amenable to the production of proteins from cloned genes. Mutations of other genes can help to ameliorate the sometimes-deleterious effects of these protease defects (e.g. sulA, below).

Endonuclease I Deficient (endA1): The periplasmic space of wild type E. coli cells contains a nonspecific endonuclease. Extreme care must be taken to avoid degradation of plasmids prepared from these cells. The endA1 mutation deletes this endonuclease and can significantly improve the quality of plasmid preparations.

Restriction Deficient (Δ(mcrC-mrr)<sup>114::IS10</sup>): Wild type E. coli B strains carry a Type I restriction endonuclease which cleaves DNA with the site TGA(8/6) tetT. While E. coli DNA is protected from degradation by a cognate methyl-transferase, foreign DNA will be cut at these sites. The deletion described above eliminates both the methylase and the endonuclease.

Methyl Restriction Deficient (Δ(mcrC-mrr)<sup>114::IS10</sup> and R(mcr-car73::MiniTn10--Tet<sup>2</sup> endA1): E. coli has a system of enzymes encoded by mcrA, mcrB, and mrr which will cleave DNA with methylation patterns found in higher eukaryotes, as well as some plant and bacterial strains. All three Mcr enzymes and Mrr have been inactivated in T7 Express allowing the introduction of eukaryotic DNA of genomic origin (e.g. primary libraries) if desired.

Phage Resistant (θA2): T1, an extremely virulent phage requires the fhuA2 gene, isolated from T1 primary libraries if desired.