

pTYB3 Vector



1-800-632-7799
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www.neb.com



N6703S 003111113111

N6703S

10 µg Lot: **0031111** Exp: **11/13**
200 µg/ml Store at **-20°C**

Description: pTYB3 is an *E. coli* expression vector (7,477 bp) used in the IMPACT™ Kit (1). This C-terminal fusion vector is designed for the in-frame insertion of a target gene into a polylinker upstream of an intein tag (the *Sce* VMA intein/chitin binding domain, 55 kDa) (1,2). This results in the fusion of the C-terminus of the target protein to the N-terminus of the intein tag. Thiol-induced self-cleavage of the intein releases the target protein from the chitin-bound intein tag, resulting in a single column purification of the target protein (3).

For the fusion of the N-terminus of the target protein to the intein tag, use pTYB11 (NEB #N6901) or pTYB12 (NEB #N6902).

Source: pTYB3 is isolated from an *E. coli* strain (r-m-) by a standard plasmid purification procedure.

Supplied in: 10 mM Tris-HCl (pH 8.0), 1 mM EDTA.

Polylinker Region:

T7 Universal Primer →
5'...CGG GGA TCT CGA TCC CGC GAA ATT AAT ACG ACT CAC TAT AGG GGA ATT GTG AGC
T7 Promoter lac operator
GGA TAA CAA TTC CCC TCT AGA AAT AAT TTT GTT TAA CTT TAA GAA GGA GAT ATA
XbaI Shine Dalgarno Intein →
Met Ala Ser Ser Arg Val Asp Gly Gly Arg Glu Phe Leu Glu Gly Ser Ser Cys1
ACC ATG GCT AGC TCG CGA GTC GAC GGC GGC CGC GAA TTC CTC GAG GGC TCT TCC TGC
NcoI NheI NruI Sall NotI EcoRI XhoI SapI
TTT GCC AAG GGT ACC AAT GTT TTA ATG GCG GAT GGG TCT ATT GAA TGT ATT
KpnI
GAA AAC ATT GAG GTT GGT AAT AAG GTC ATG GGT ...3'
← Intein Reverse Primer

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TTT GCC AAG GGT ACC AAT GTT TTA ATG GCG GAT GGG TCT ATT GAA TGT ATT
KpnI
GAA AAC ATT GAG GTT GGT AAT AAG GTC ATG GGT ...3'
← Intein Reverse Primer

Features of pTYB3:

- The NcoI site in the polylinker contains an ATG sequence for translation initiation.
- Use of the SapI site allows cloning of the target protein adjacent to the intein, resulting in cleavage of the target protein without any additional amino acids at its C-terminus.
- Unique sites are indicated in **bold**.
- Expression of the fusion gene is under the control of an IPTG-inducible T7 promoter (4).

- Origin of DNA replication from bacteriophage M13, which allows for the production of single-stranded DNA by helper phage superinfection of cells bearing the plasmid (M13K07 Helper Phage, NEB #N0315)
- A pBR322 derivative with a ColE1 replication origin.
- Ampicillin resistance
- Other IMPACT vectors are available which allow for fusion of a target gene to N- or C- terminus of an intein. The cleavage reaction may be induced by thiol reagent or temperature/pH shift.
- Companion vectors (pTYB1, pTYB2, pTYB4) differ only in the sites present in the polylinker.
- A wide range of *E. coli* host strains: T7 Express Competent *E. coli* (High Efficiency) (NEB #C2566) or BL21(DE3) Competent *E. coli* (NEB #C2527) and derivatives.

(see other side)

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NcoI NheI NruI Sall NotI EcoRI XhoI SapI
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(see other side)

CERTIFICATE OF ANALYSIS

References:

1. Chong, S., Mersha, F.B., Comb, D.G., Scott, M. E., Landry, D., Vence, L.M., Perler, F.B., Benner, J., Kucera, R.B., Hirvonen, C.A., Pelletier, J.J., Paulus, H. and Xu, M. -Q. (1997). Single-column purification of free recombinant proteins using a self-cleavable affinity tag derived from a protein splicing element. *Gene* 192, 271–281.
2. Watanabe, T., Ito, Y., Yamada, T., Hashimoto, M., Sekine, S. and Tanaka, H. (1994). The role of the C-terminal domain and type III domains of chitinase A1 from *Bacillus circulans* WL–12 in chitin degradation. *J. Bacteriol.* 176, 4465–4472.
3. Chong, S., Shao, Y., Paulus, H. Benner, J., Perler F.B. and Xu, M.-Q. (1996). Protein splicing involving the *Saccharomyces cerevisiae* VMA intein: the steps in the splicing pathway, side reactions leading to protein cleavage, and establishment of an *in vitro* splicing system *J. Biol. Chem.* 271, 22159–22168.
4. Dubendorff, J.W. and Studier, F.W. (1991). Controlling basal expression in an inducible T7 expression system by blocking the target T7 promoter with *lac* repressor *J. Mol. Biol.* 219, 45–59.

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U.S. Patent Nos. 5,496,714, 5,834,247

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