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in status



10 µg	Lot: 0021111	Exp: 11/14
200 µg/ml	Store at -20°C	

Description: pKYB1 is an *E. coli* expression vector for use with the IMPACT[™] Kit (1,2). It is designed for in-frame insertion of a target gene into the polylinker, upstream of the Sce VMA intein/chitin binding domain (55 kDa)(1,3). pKYB1 carries the kanamycin resistance gene (Kn) from Tn 903. This double stranded vector is 8,393 bp in length.

Source: pKYB1 is isolated from an *E. coli* strain (r-m-) by a standard plasmid purification procedure.



N6706S

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Source: pKYB1 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.

Features of this vector:

- A pBR322 derivative
- The Ndel site in the polylinker contains an ATG sequence for translation initiation.
- Use of the Sapl site allows for cloning of the target protein adjacent to the intein, resulting in purification of the target protein without any additional amino acids at its C-terminus.

Polylinker Region:

pKYB1 T7 Universal Primer \rightarrow												
5´CGG GG	A TCT CGA	TCC CGC	GAA AT <u>T</u>	AAT ACG	ACT CAC TA	<u>at ag</u> g g <u>ga</u>	ATT GTG AGC					
				T7 P	romoter	lac	<i>lac</i> operator					
<u>GGA TA</u>	A CAA TTC	<u>CC</u> C <u>TCT</u>	<u>AGA</u> AAT	AAT TTT	GTT TAA C	it taa <u>gaa</u>	<u>gga g</u> at ata					
XbaI Shine Dalgarno												
							Gly Ser Ser GGC TCT TCC	Cys1				
NdeI	NheI	NruI	Sall	NotI	EcoR	XhoI	SapI					
TTT GC			GTT TTA	ATG GCG	GAT GGG TO	CT ATT GAA	TGT ATT					
KpnI												
GAA AA	C ATT GAG	GTT GGT	AAT AAG	GTC ATG	<u>GGT</u> 3´							
← Intein Reverse Primer												

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5´ .	.CGG	GGA	тст	CGA	тсс	CGC	GAA	AT <u>T</u>	AAT	ACG	ACT	CAC	TAT	<u>AG</u> G	G <u>GA</u>	ATT	GTG	AGC		
		T7 P								romoter /				la	<i>ac</i> operator					
	<u>GGA</u>	TAA	CAA	TTC	<u>CC</u> C	TCT	AGA	AAT	AAT	TTT	GTT	TAA	СТТ	TAA	<u>GAA</u>	GGA	<u>G</u> AT	ATA		
	XbaI									Shine Dalgarno						\rightarrow				
	<u>CAT</u>								~							~		Ser <u>TC</u> C	Cys1 TGC	
	Nc	leI	Nh	eI	Nr	uI	Sal	I	ľ	lotI		Есо	RI	Xh	IoI	5	SapI			
	TTT	GCC	AAG		ACC onI	AAT	GTT	TTA	ATG	GCG	GAT	GGG	TCT	ATT	GAA	TGT	ATT			

GAA AAC ATT GAG GTT GGT AAT AAG GTC ATG <u>GGT</u> ...3⁺ ← Intein Reverse Primer

- Unique sites are in bold. Nhel and Nrul are not unique.
- Expression of the fusion gene is under the stringent control of the T7 promoter (4) and is regulated by IPTG due to the presence of a *lac* I gene.
- Origin of DNA replication from bacteriophage M13, which allows for the production of singlestranded DNA by helper phage superinfection of cells bearing the plasmid (M13K07 Helper Phage, NEB #N0315)

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gene.

- Kanamycin resistant
- Compatible restriction sites for subcloning a fusion gene from other IMPACT vectors.
- 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.

References:

- 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.
- 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.

(see other side) CERTIFICATE OF ANALYSIS

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- 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.
- 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.

Additional information such as vector sequences and frequently asked questions, are available at www. neb.com.

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

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