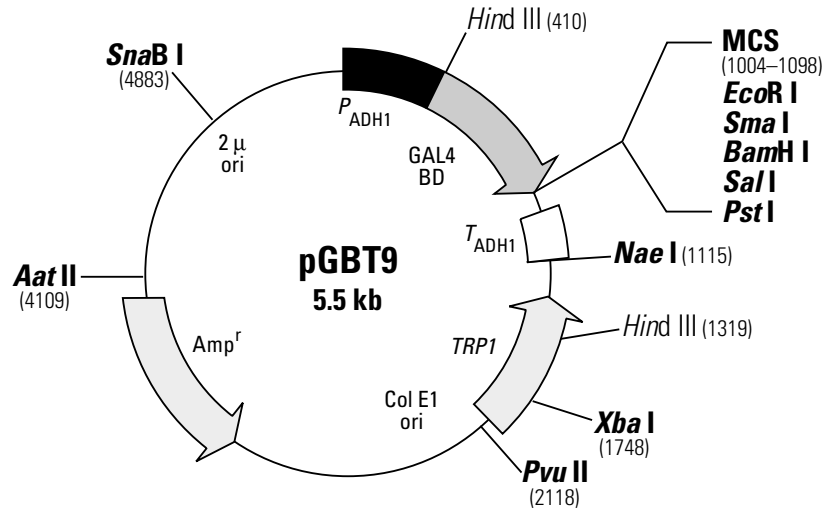


pGBT9 DNA-BD Vector Information

PT1021-5

GenBank Accession #: U07646

Catalog #K1605-A



MATCHMAKER 5' DNA-BD Insert
Screening Amplimer &
GAL4 DNA-BD Sequencing Primer

827
TCA TCG GAA GAG AGT AGT AAC AAA GGT CAA AGA CAG TTG ACT GTA TCG CCG

878
GAA TTC CCG GGG ATC CGT CGA CCT GCA GCC AAG CTA ATT CCG GGC GAA TTT
EcoR I Sma I BamH I Sal I Pst I STOP (ORF 2)

929
CTT ATG ATT TAT GAT TTT TAT TAT TAA ATA AGT TAT AAA AAA AAT AAG TGT ATA

983
CAA ATT TTA AAG TGA CTC TTA GGT TTT AAA ACG

MATCHMAKER 3' DNA-BD Insert
Screening Amplimer

Restriction map and multiple cloning site (MCS) of ppGBT9 DNA-BD Vector. Unique restriction sites are in bold.

Description:

pGBT9 generates a hybrid protein that contains the sequences for the GAL4 DNA-binding domain (DNA-BD; a.a. 1–147). For the construction of a hybrid protein, the gene encoding the protein of interest is ligated into the MCS in the correct orientation and in the correct reading frame such that a fusion protein is generated. The fusion protein is expressed in yeast host cells from the constitutive *ADH1* promoter; transcription is terminated at the *ADH1* transcription termination signal. The hybrid protein is targeted to the yeast nucleus by nuclear localization sequences that are an intrinsic part of the GAL4 DNA-BD (2). pGBT9 is a shuttle vector that replicates autonomously in both *E. coli* and *S. cerevisiae*. It carries the *bla* gene (for ampicillin resistance in *E. coli*) and the *TRP1* nutritional marker that allow yeast auxotrophs carrying pGBT9 to grow on limiting synthetic medium lacking Trp.



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(PR81583)

Location of features

- Promoter
Fragment containing the *S. cerevisiae ADH1* promoter: 10–406
- GAL4 DNA-binding domain polypeptide
Start codon: 434–436; stop codon: 953–955
GAL4 codons 1–147: 434–874
- Multiple cloning site: 878–905
- Transcription termination signal
Fragment carrying the *S. cerevisiae ADH1* terminator: 921–1112
- *TRP1* coding sequence
Start codon: 1835–1833; stop codon: 1163–1161
- Col E1 origin of plasmid replication: 2322–2965
- Ampicillin resistance gene
Promoter: –35 region: 4043–4038; –10 region: 4020–4015
Transcription start point: 4008
Ribosome binding site: 3985–3981
 β -lactamase coding sequences
Start codon: 3973–3971; stop codon: 3115–3113
Signal peptide: 3973–3905
Mature protein: 3904–3116
- Fragment containing the 2 μ origin of replication: 1–1348

Primer locations:

- MATCHMAKER DNA-BD 5' Insert Screening Amplimer (#5417-1) or GAL4 BD Sequencing Primer (#6474-1): 827–843
- MATCHMAKER DNA-BD 3' Insert Screening Amplimer (#5417-1): 1015–994

Propagation in *E. coli*

- Suitable host strains: DH5 α and other general purpose strains.
- Selectable marker: plasmid confers resistance to ampicillin (50 μ g/ml) to *E. coli* hosts.
- *E. coli* replication origin: Col E1
- Copy number: 15–20

Propagation in *S. cerevisiae*

- Suitable host strains: Y187(α), Y190(a), HF7c(a), CG1945(a), PJ69-2A, PJ69-4A, and SFY526(a)
- Selectable marker: *TRP1*
- *S. cerevisiae* replication origin: 2 μ
- Copy number: multiple copy

References

1. Bartel, P. L., *et al.* (1993) In *Cellular Interactions in Development: A Practical Approach* (Oxford University Press, Oxford) pp. 135–179.
2. Silver, P. A., *et al.* (1984) *Proc. Natl. Acad. Sci. USA* **91**:5951–5955.

Note: The attached sequence file has been compiled from information in the sequence databases, published literature, and other sources, together with partial sequences obtained by CLONTECH. This vector has not been completely sequenced.