

Product datasheet for SC201421

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

GABPB2 (GABPB1) (NM 016655) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: GABPB2 (GABPB1) (NM_016655) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: GABPB1

Synonyms: BABPB2; E4TF1; E4TF1-47; E4TF1-53; E4TF1B; GABPB; GABPB-1; GABPB2; NRF2B1; NRF2B2

ACCN: NM_016655

Insert Size: 166 bp

Insert Sequence: >SC201421 3'UTR clone of NM_016655

The sequence shown below is from the reference sequence of NM_016655. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AGCCCTCAATAAATAAATATTGAATGAA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeg: NM 016655.5





GABPB2 (GABPB1) (NM_016655) Human 3' UTR Clone - SC201421

Summary: This gene encodes the GA-binding protein transcription factor, beta subunit. This protein

forms a tetrameric complex with the alpha subunit, and stimulates transcription of target genes. The encoded protein may be involved in activation of cytochrome oxidase expression and nuclear control of mitochondrial function. The crystal structure of a similar protein in mouse has been resolved as a ternary protein complex. Multiple transcript variants encoding

distinct isoforms have been identified for this gene. [provided by RefSeq, Jul 2008]

Locus ID: 2553

MW: 6.3