

## **Product datasheet for SC202814**

## OriGene Technologies, Inc.

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## beta glucuronidase (GUSB) (NM\_000181) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

**Product Name:** beta glucuronidase (GUSB) (NM\_000181) Human 3' UTR Clone

**Symbol:** beta glucuronidase

**Synonyms:** BG; MPS7

Mammalian Cell Neomycin

Selection:

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_000181

**Insert Size:** 243 bp

Insert Sequence: >SC202814 3'UTR clone of NM\_000181

The sequence shown below is from the reference sequence of NM\_000181. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

CAATGTTTGGAAAACAGCCTGTTTACTTGAGCAAGACTGATACCACCTGCGTGTCCCTTCCTCCCCGAG TCAGGGCGACTTCCACAGCAGCAGAACAAGTGCCTCCTGGACTGTTCACGGCAGACCAGAACGTTTCTG GCCTGGGTTTTGTGGTCATCTATTCTAGCAGGGAACACTAAAGGTGGAAATAAAAGATTTTCTATTATG

GAAATAAAGAGTTGGCATGAAAGTGGCTACTGAAAA

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

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.

**RefSeq:** <u>NM 000181.4</u>





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Summary: This gene encodes a hydrolase that degrades glycosaminoglycans, including heparan sulfate,

dermatan sulfate, and chondroitin-4,6-sulfate. The enzyme forms a homotetramer that is localized to the lysosome. Mutations in this gene result in mucopolysaccharidosis type VII. Alternative splicing results in multiple transcript variants. There are many pseudogenes of this

locus in the human genome.[provided by RefSeq, May 2014]

**Locus ID:** 2990

**MW:** 9.2