

Product datasheet for SC200146

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H2BC8 (NM_003518) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: H2BC8 (NM_003518) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: H2BC8

Synonyms: dJ221C16.8; H2B.1A; H2B/a; H2BC4; H2BC6; H2BC7; H2BC10; H2BFA; HIST1H2BG

ACCN: NM_003518

Insert Size: 93 bp

Insert Sequence: >SC200146 3'UTR clone of NM_003518

The sequence shown below is from the reference sequence of NM_003518. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GCTGTCACCAAGTATACAAGCTCCAAGTAAATGTGTGCTTAGGTGCTTTAAAAACTCAAAGGCTCTTTTC

AGAGCCACTCAAGTCTCACATAAA

 ${\tt CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG}$

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 003518.4







Summary:

Histones are basic nuclear proteins that are responsible for the nucleosome structure of the chromosomal fiber in eukaryotes. Nucleosomes consist of approximately 146 bp of DNA wrapped around a histone octamer composed of pairs of each of the four core histones (H2A, H2B, H3, and H4). The chromatin fiber is further compacted through the interaction of a linker histone, H1, with the DNA between the nucleosomes to form higher order chromatin structures. The protein has antibacterial and antifungal antimicrobial activity. This gene is intronless and encodes a replication-dependent histone that is a member of the histone H2B family. Transcripts from this gene lack polyA tails; instead, they contain a palindromic termination element. This gene is found in the large histone gene cluster on chromosome 6p22-p21.3. [provided by RefSeq, Aug 2015]

Locus ID: 8339 **MW:** 3.4