

Product datasheet for SC210547

HOXC6 (NM 153693) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: HOXC6 (NM_153693) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: HOXC6

Synonyms: CP25; HHO.C8; HOX3; HOX3C

ACCN: NM_153693

Insert Size: 891 bp

Insert Sequence: >SC210547 3'UTR clone of NM_153693

The sequence shown below is from the reference sequence of NM_153693. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

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).



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HOXC6 (NM_153693) Human 3' UTR Clone - SC210547

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 153693.5</u>

Summary: This gene belongs to the homeobox family, members of which encode a highly conserved

family of transcription factors that play an important role in morphogenesis in all multicellular organisms. Mammals possess four similar homeobox gene clusters, HOXA, HOXB, HOXC and HOXD, which are located on different chromosomes and consist of 9 to 11 genes arranged in tandem. This gene, HOXC6, is one of several HOXC genes located in a cluster on chromosome 12. Three genes, HOXC5, HOXC4 and HOXC6, share a 5' non-coding exon. Transcripts may include the shared exon spliced to the gene-specific exons, or they may

include only the gene-specific exons. Alternatively spliced transcript variants encoding different isoforms have been identified for HOXC6. Transcript variant two includes the shared

exon, and transcript variant one includes only gene-specific exons. [provided by RefSeq, Jul

2008]

Locus ID: 3223

MW: 33.3