

Product datasheet for SC210351

OriGene Technologies, Inc.

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Collagen I (COL1A2) (NM_000089) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Collagen I (COL1A2) (NM_000089) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: COL1A2

Synonyms: EDSARTH2; EDSCV; OI4

ACCN: NM_000089

Insert Size: 864 bp

Insert Sequence: >SC210351 3'UTR clone of NM_000089

The sequence shown below is from the reference sequence of NM_000089. 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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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 000089.4</u>

Summary: This gene encodes the pro-alpha2 chain of type I collagen whose triple helix comprises two

alpha1 chains and one alpha2 chain. Type I is a fibril-forming collagen found in most connective tissues and is abundant in bone, cornea, dermis and tendon. Mutations in this gene are associated with osteogenesis imperfecta types I-IV, Ehlers-Danlos syndrome type VIIB, recessive Ehlers-Danlos syndrome Classical type, idiopathic osteoporosis, and atypical Marfan syndrome. Symptoms associated with mutations in this gene, however, tend to be less severe than mutations in the gene for the alpha1 chain of type I collagen (COL1A1) reflecting the different role of alpha2 chains in matrix integrity. Three transcripts, resulting from the use of alternate polyadenylation signals, have been identified for this gene.

[provided by R. Dalgleish, Feb 2008]

Locus ID: 1278 MW: 33.9