

## **Product datasheet for SC209792**

## **DDX11 (NM 152438) Human 3' UTR Clone**

## **Product data:**

**Product Type:** 3' UTR Clones

Product Name: DDX11 (NM\_152438) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: DDX11

Synonyms: CHL1; CHLR1; KRG2; WABS

**ACCN:** NM\_152438

**Insert Size:** 796 bp

Insert Sequence: >SC209792 3'UTR clone of NM\_152438

The sequence shown below is from the reference sequence of NM\_152438. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GGTCCTTAAATAAAAGTGTGCTGTTGGTTTCTGCTGA

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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## DDX11 (NM\_152438) Human 3' UTR Clone - SC209792

**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 152438.2</u>

Summary: DEAD box proteins, characterized by the conserved motif Asp-Glu-Ala-Asp (DEAD), are

putative RNA helicases. They are implicated in a number of cellular processes involving

alteration of RNA secondary structure such as translation initiation, nuclear and

mitochondrial splicing, and ribosome and spliceosome assembly. Based on their distribution

patterns, some members of this family are believed to be involved in embryogenesis, spermatogenesis, and cellular growth and division. This gene encodes a DEAD box protein, which is an enzyme that possesses both ATPase and DNA helicase activities. This gene is a homolog of the yeast CHL1 gene, and may function to maintain chromosome transmission fidelity and genome stability. Alternative splicing results in multiple transcript variants

encoding distinct isoforms. [provided by RefSeq, Jul 2008]

**Locus ID:** 1663

MW: 28.9