

## **Product datasheet for SC313601**

## HNF6 (ONECUT1) (NM\_004498) Human Untagged Clone

## **Product data:**

**Product Type:** Expression Plasmids

Product Name: HNF6 (ONECUT1) (NM\_004498) Human Untagged Clone

Tag: Tag Free Symbol: HNF6

Synonyms: HNF-6; HNF6; HNF6A

Mammalian Cell None

Selection:

Vector:

pCMV6-XL5

E. coli Selection: Ampicillin (100 ug/mL)

Fully Sequenced ORF: >OriGene ORF sequence for NM\_004498 edited

ATGAACGCGCAGCTGACCATGGAAGCGATCGGCGAGCTGCACGGGGTGAGCCATGAGCCG GTGCCCGCCCTGCCGACCTGCTGGGCGCGCAGCCCCCACGCGCGCAGCTCCGTGGCGCAC CGCGGCAGCCACCTGCCCCCGCGCACCCGCGCTCCATGGGCATGGCGTCCCTGCTGGAC GGCGGCAGCGGCGGGGGAGATTACCACCACCACCACCGGGCCCCTGAGCACAGCCTGGCC GGCCCCTGCATCCCACCATGACCATGGCCTGCGAGACTCCCCCAGGTATGAGCATGCCC ACCACCTACACCACCTTGACCCCTCTGCAGCCGCTGCCTCCCATCTCCACAGTCTCGGAC CTGGCGGCAACGTGAGCGTAGCTTCACGCTCATGCGGGATGAGCGCGGGCTGGCCTCC ATGAATAACCTCTATACCCCCTACCACAAGGACGTGGCCGGCATGGGCCAGAGCCTCTCG CCCCTCTCCAGCTCCGGTCTGGGCAGCATCCACAACTCCCAGCAAGGGCTCCCCCACTAT GCCCACCGGGGGCCGCCATGCCCACCGACAGATGCTCACCCCCAACGGCTTCGAAGCC ATGGTGCCCATCAACGGCCTTCCTCCGCACCATCCCCACGCCCACCTGAACGCCCAGGGC CACGGGCAACTCCTGGGCACAGCCCGGGAGCCCAACCCTTCGGTGACCGGCGCGCAGGTC AGCAATGGAAGTAATTCAGGTCAGATGGAAGAGATCAATACCAAAGAGGTGGCGCAGCGT ATCACCACCGAGCTCAAGCGCTACAGCATCCCACAGGCCATCTTCGCGCAGAGGGTGCTC TGCCGCTCCCAGGGGACCCTCTCGGACCTGCTGCGCAACCCCAAACCCTGGAGCAAACTC AAATCCGGCCGGGAGACCTTCCGGAGGATGTGGAAGTGGCTGCAGGAGCCGGAGTTCCAG CGCATGTCCGCGCTCCGCTTAGCAGCATGCAAAAGGAAAGAACAAGAACATGGGAAGGAT AGAGGCAACACCCAAAAAGCCCAGGTTGGTCTTCACAGATGTCCAGCGTCGAACTCTA CATGCAATATTCAAGGAAAATAAGCGTCCATCCAAAGAATTGCAAATCACCATTTCCCAG CAGCTGGGGTTGGAGCTGAGCACTGTCAGCAACTTCTTCATGAACGCAAGAAGGAGGAGT CTGGACAAGTGGCAGGACGAGGCAGCTCCAATTCAGGCAACTCATCTTCATCAAGC

ACTTGTACCAAAGCATGA

**Restriction Sites:** Please inquire



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## HNF6 (ONECUT1) (NM\_004498) Human Untagged Clone - SC313601

**ACCN:** NM\_004498

**Insert Size:** 1500 bp

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

**OTI Annotation:** The open reading frame of this TrueClone was fully sequenced and found to be a perfect

match to the protein associated to this reference.

**Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

**Reconstitution Method:** 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of

shipping when stored at -20°C.

RefSeq: NM 004498.1, NP 004489.1

RefSeq Size: 1925 bp
RefSeq ORF: 1398 bp
Locus ID: 3175
UniProt ID: Q9UBC0

Cytogenetics: 15q21.3

**Protein Families:** ES Cell Differentiation/IPS, Transcription Factors

**Protein Pathways:** Maturity onset diabetes of the young

**Gene Summary:** This gene encodes a member of the Cut homeobox family of transcription factors. Expression

of the encoded protein is enriched in the liver, where it stimulates transcription of liver-expressed genes, and antagonizes glucocorticoid-stimulated gene transcription. This gene may influence a variety of cellular processes including glucose metabolism, cell cycle

regulation, and it may also be associated with cancer. Alternative splicing results in multiple

transcript variants. [provided by RefSeq, Dec 2012]

Transcript Variant: This variant (1) represents the longer transcript and encodes the functional protein. Sequence Note: This RefSeq record was created from transcript and genomic

sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.