

# Gene Section

## Short Communication

# TALDO1 (transaldolase 1)

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## Abstract

Short communication on TALDO1, with data on DNA/RNA, on the protein encoded and where the gene is implicated.

## Identity

**Other names:** TAL, TAL-H, TALDOR, TALH

**HGNC (Hugo):** TALDO1

**Location:** 11p15.5

## DNA/RNA

### Note

Starts at 747432 bp from pter and ends at 765024 bp from pter according to hg19-Feb\_2009.

Size: 17593 bases; Orientation: TALDO1 is on the plus strand.

### Description

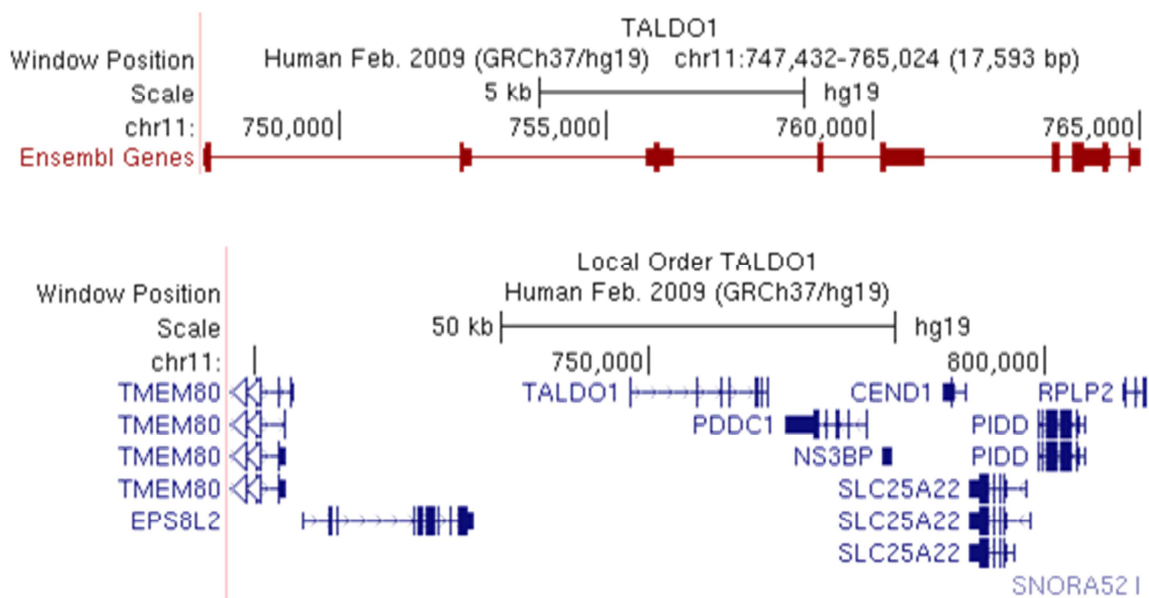
Exons 2 and 3 of TALDO1 contain retrotransposable elements (Banki et al., 1994).

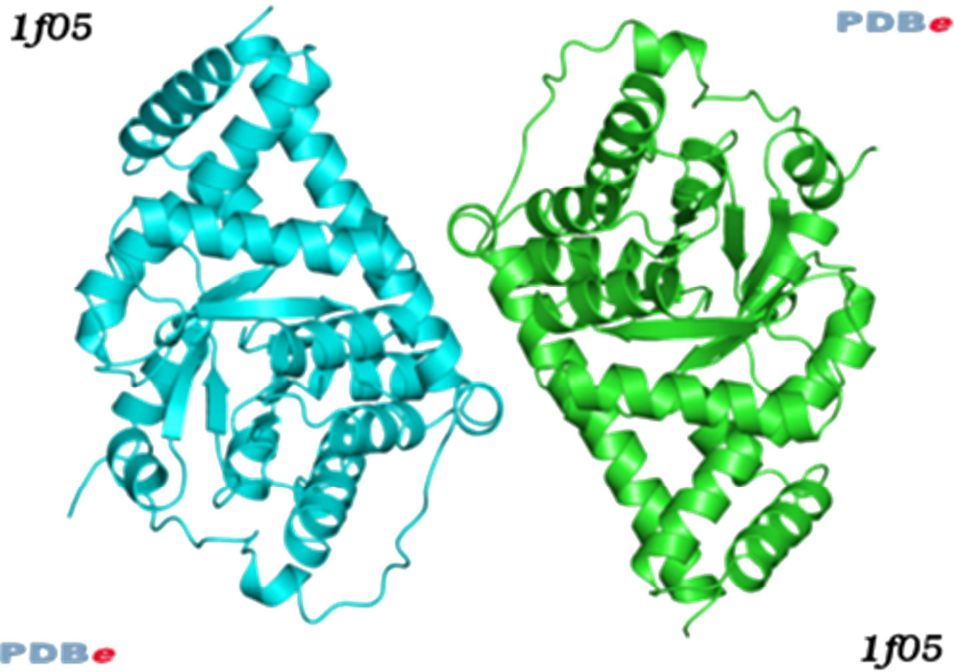
### Transcription

TALDO1 has 8 exons and its mRNA is composed of 1319 bp.

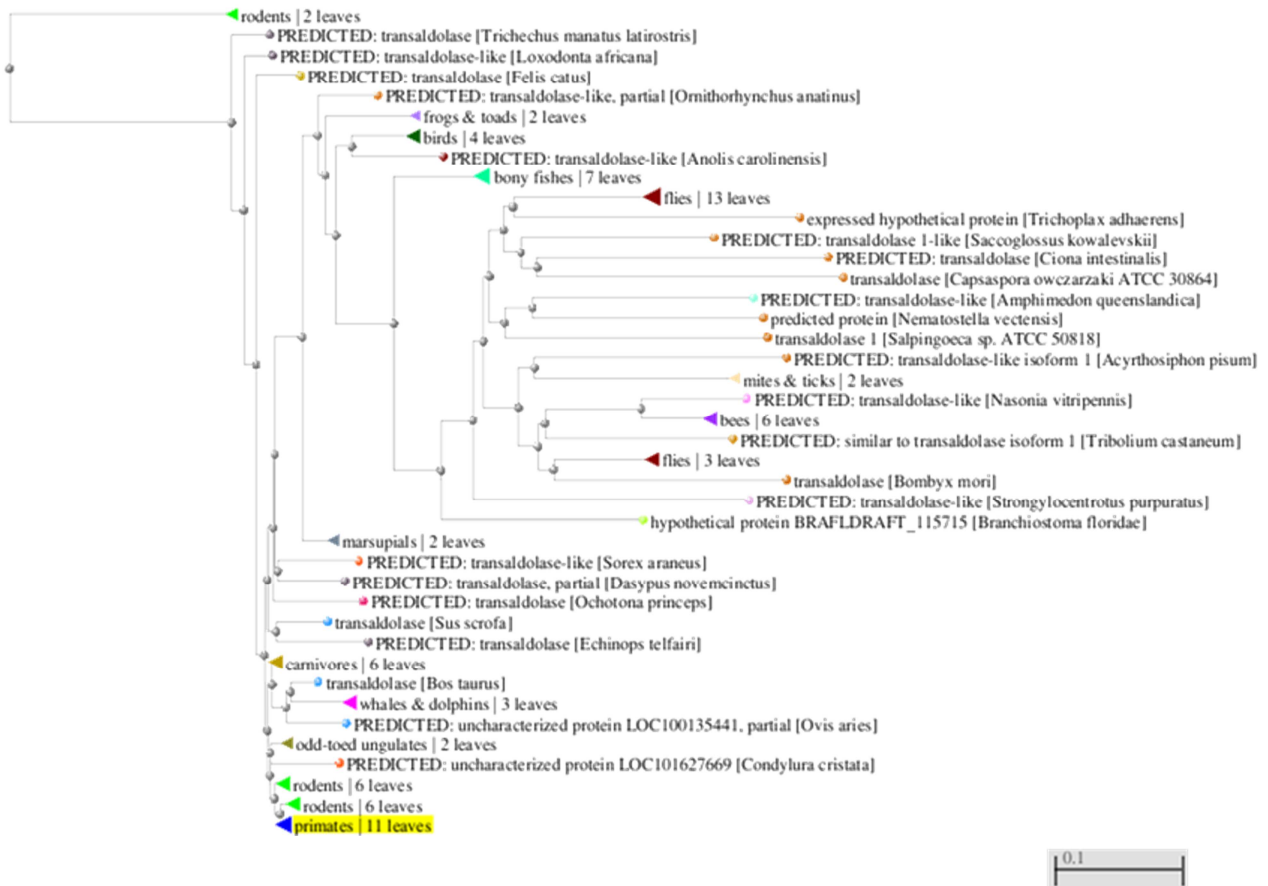
### Pseudogene

TALDO1P1 (Transaldolase 1 pseudogene 1).





The 3-dimensional structure of transaldolase from Thorell et al., 2000. Image downloaded from <http://www.ebi.ac.uk>.



From blast of reference proteins (Refseq) against human Transaldolase (NP\_006746.1).

## Protein

### Description

Transaldolase (TAL) is a 337 amino acid protein in the non-oxidative phase of the pentose phosphate pathway (PPP) with a predicted mass of 37.55kDa. TAL has an  $\alpha/\beta$  barrel and that includes lysine 142 which is responsible for generating the Schiff base intermediate during sugar phosphate metabolism (Thorell et al., 200). A mass spectrometry based investigation of the acetylome identified TAL acetylation at lysines 286, 269, 321, 219 (Choudhary et al., 2009).

It has also been proposed that TAL activity may be affected by phosphorylation (Lachaise et al., 2001).

### Expression

TALDO1 is ubiquitously expressed, except in erythrocytes.

### Localisation

Cytosol and nucleus (Colombo et al., 1997).

### Function

The reversible reaction carried out by TAL is: erythros-4-phosphate+fructose-6-phosphateglyceraldehydes-3-phosphate+sedoheptulose-7-phosphate. TAL has been proposed as the rate limiting enzyme in the non-oxidative PPP (Banki et al., 1996; Heinrich et al., 1976; Perl, 2007; Wood, 1985).

### Homology

Using the blastp function within Homo sapiens, the only protein to share homology within humans (a paralog) was sorting nexin 32 (SNX32) with an identity of 36%.

## Mutations

### Germinal

Homozygous deletion of Serine 171 due to the loss of 3bp in the TALDO1 sequence results in liver cirrhosis and subsequent carcinogenesis.

## Implicated in

### Hepatocellular carcinoma

#### Cytogenetics

Deletion of nucleotides 512-514 in TALDO1 resulted in the loss of serine 171 in the TAL protein and subsequent TAL deficiency (Verhoeven et al., 2001; Valayannopoulos et al., 2006). TAL

deficiency results in the accumulation of sedoheptulose-7-phosphate and polyols. Further studies into TAL deficiency determined that the deletion of S171 resulted in a complete loss of enzymatic activity and rapid degradation in the proteasome (Grossman et al., 2004). Missense mutations at arginine 192, in which the arginine is mutated to either a histidine or cysteine, also results in loss of TAL activity and liver damage in patients (Verhoeven et al., 2005; Wamelink et al., 2008). In addition to liver damage, renal and cardiac complications are also present in these patients (Verhoeven et al., 2001; Valayannopoulos et al., 2006; Verhoeven et al., 2005; Wamelink et al., 2008).

In a mouse model of TAL deficiency, sperm dysmotility and subsequent male infertility are present (Perl et al., 2006). Furthermore, TAL deficiency results in the development of hepatosteatosis, cirrhosis, and hepatocellular carcinoma in both homozygous TAL knockouts and heterozygous mice relative to C57Bl/6 wild type mice (Hanczko et al., 2009).

The pathogenic mechanism of liver damage in TAL deficiency is linked to depletion of NADPH, oxidative stress, and mitochondrial dysfunction (Perl et al., 2011).

In TAL deficiency, it has been proposed that oxidative stress is exacerbated by increased aldose reductase activity which generates polyols and depletes NADPH (Perl et al., 2011). Low NADPH diminishes the cells ability to regulate cellular redox and polyols can induce proliferation through JNK/c-Jun (Perl et al., 2011). Thus, TAL deficiency and insufficiency predispose to oxidative stress which promotes liver damage, increased proliferation, and hepatocellular carcinoma.

### Squamous cell carcinoma of the head and neck

#### Cytogenetics

3 SNPs in the TALDO1 gene have been associated with different squamous cell carcinoma of the head and neck risk.

The conversion of cytosine to either guanine or thymine at 490bp upstream of the origin of replication (rs10794338) was protective against tumorigenesis (Basta et al., 2008).

In contrast, the mutation of thymine to adenine at position 1874 (rs3901233) and adenine to cytosine at position 2187 (rs4963163) increase the risk of squamous cell cancer of the head and neck (Basta et al., 2008).



From blastp of NP\_006746.1 against all non-redundant protein sequences.

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