

TMC01

Transmembrane and coiled-coil domains 1 (TMC01) is a recently discovered [transmembrane protein](#) of [endoplasmic reticulum](#) (ER), which plays a critical role in maintaining [calcium](#) homeostasis. TMC01 dysfunction has been proved to be closely related to a variety of human diseases, including [glaucoma](#), deformities, [mental retardation](#), and tumorigenesis.

Xin et al. identified an autosomal recessive condition in 11 individuals in the Old Order Amish of northeastern Ohio. The syndrome was characterized by distinctive craniofacial dysmorphism, skeletal anomalies, and mental retardation. The typical craniofacial dysmorphism included brachycephaly, highly arched bushy eyebrows, synophrys, long eyelashes, low-set ears, microdontism of primary teeth, and generalized gingival hyperplasia, whereas Sprengel deformity of scapula, the fusion of the spine, rib abnormalities, pectus excavatum, and pes planus represented skeletal anomalies. The genome-wide homozygosity mapping using six affected individuals localized the disease gene to a 3.3-Mb region on chromosome 1q23.3-q24.1. Candidate gene sequencing identified a homozygous frameshift mutation, c.139_140delAG, in the transmembrane and coiled-coil domains 1 (TMC01) gene, as the pathogenic change in all affected members of the extended pedigree. This mutation is predicted to result in a severely truncated protein (p.Ser47Ter) of only one-fourth the original length. The TMC01 gene product is a member of DUF841 superfamily of several eukaryotic proteins with unknown function. The gene has highly conserved amino acid sequence and is universally expressed in all human tissues examined. The high degree of conservation and the ubiquitous expression pattern in human adult and fetal tissues suggest a critical role for TMC01. This report shows a TMC01 sequence variant being associated with a genetic disorder in human. We propose "TMC01 defect syndrome" as the name of this condition ¹⁾.

The role of TMC01 in gliomas remains unclear. The purpose of this study was to detect the role of TMC01 in the pathogenesis and progression of gliomas. This study demonstrated that TMC01 was upregulated in gliomas and its overexpression predicted poor prognosis. We also revealed that the expression of TMC01 was associated with the World Health Organization (WHO) grade of gliomas. Knockdown of TMC01 inhibited the [proliferation](#) and induced [apoptosis](#) of U87 and U251 cells. In addition, TMC01 induced Glioblastoma [cell migration](#) and invasion by promoting epithelial-mesenchymal transition (EMT). This data collectively proved the crucial role of TMC01 as a novel prognostic factor and underlying therapeutic target for glioma patients ²⁾.

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Xin B, Puffenberger EG, Turben S, Tan H, Zhou A, Wang H. Homozygous frameshift mutation in TMC01 causes a syndrome with craniofacial dysmorphism, skeletal anomalies, and mental retardation. Proc Natl Acad Sci U S A. 2010 Jan 5;107(1):258-63. doi: 10.1073/pnas.0908457107. Epub 2009 Dec 14. PMID: 20018682; PMCID: PMC2806776.

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Gao L, Ye Z, Liu JH, Yang JA, Li Y, Cai JY, Wang YX, Tong SA, Deng G, Zhang S, Chen QX. TMC01 expression promotes cell proliferation and induces epithelial-mesenchymal transformation in human gliomas. Med Oncol. 2022 May 15;39(5):90. doi: 10.1007/s12032-022-01687-y. PMID: 35568751.

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