

T641M — WFS1 Molecular Atlas Card

Variant type: Missense

Substitution: Threonine (T) → Methionine (M) at position 641

Domain context: C-terminal ER-lumenal (calcium binding, calmodulin, chaperone)

ALPHAMISSENSE

- **Pathogenicity score:** 0.1932
- **Class:** likely benign

ALPHAFOLD CONFIDENCE

- **pLDDT at residue 641:** 86.56

> **DynaMut2 $\Delta\Delta G$:** not yet computed for this variant — AlphaMissense + AlphaFold

> confidence shown above. Stability $\Delta\Delta G$ and the wild-type/mutant structural

> comparison backfill behind this note.

CLINICAL EVIDENCE

- **Classification:** Uncertain significance
- **Review status:** criteria provided, multiple submitters, no conflicts
- **Associated conditions:** Wolfram-like syndrome; Cataract 41; Autosomal dominant nonsyndromic hearing loss 6; Type 2 diabetes mellitus; Wolfram syndrome 1; WFS1-related disorder
- **cDNA change:** c.1922C>T
- **ClinVar accession:** VCV000596878
- **Last evaluated:** 2023/04/26 00:00
- **Submissions:** 1

Card generated by wolfram-atlas-batch (missense AlphaMissense mint) on 2026-06-08T02:27:33.677225Z.

AlphaMissense (Cheng et al. 2023) · AlphaFold model v6 · UniProt O76024.