RARERESEARCH.AI · WFS1 MOLECULAR ATLAS · MISSENSE

M657V — WFS1 Molecular Atlas Card

Variant type: Missense

Substitution: Methionine (M) → Valine (V) at position 657

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

ALPHAMISSENSE

- **Pathogenicity score:** 0.2272
- **Class:** likely benign

ALPHAFOLD CONFIDENCE

- **pLDDT at residue 657:** 51.97

> **DynaMut2 ΔΔG:** not yet computed for this variant — AlphaMissense + AlphaFold

> confidence shown above. Stability ΔΔG and the wild-type/mutant structural

> comparison backfill behind this note.

CLINICAL EVIDENCE

- **Classification:** Uncertain significance/Uncertain risk allele
- **Review status:** criteria provided, multiple submitters, no conflicts
- **Associated conditions:** WFS1-Related Spectrum Disorders; Cataract 41; Autosomal dominant nonsyndromic hearing loss 6; Type 2 diabetes mellitus; Wolfram syndrome 1; Wolfram-like syndrome; Spastic ataxia
- **cDNA change:** c.1969A>G
- **ClinVar accession:** VCV000215365
- **Last evaluated:** 2026/01/07 00:00
- **Submissions:** 1

Card generated by `wolfram-atlas-batch` (missense AlphaMissense mint) on 2026-06-08T02:27:33.687174Z.

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