

T487I — WFS1 Molecular Atlas Card

Variant type: Missense

Substitution: Threonine (T) → Isoleucine (I) at position 487

Domain context: Luminal loop 3

ALPHAMISSENSE

- **Pathogenicity score:** 0.4261
- **Class:** ambiguous

ALPHAFOLD CONFIDENCE

- **pLDDT at residue 487:** 73.94

> **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:** Likely benign
 - **Review status:** criteria provided, single submitter
 - **cDNA change:** c.1460C>T
 - **ClinVar accession:** VCV001652195
 - **Last evaluated:** 2024/12/02 00:00
 - **Submissions:** 1
-

Card generated by `wolfram-atlas-batch` (missense AlphaMissense mint) on 2026-06-08T02:27:33.570050Z.

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