

E231K — WFS1 Molecular Atlas Card

Variant type: Missense

Substitution: Glutamic acid (E) → Lysine (K) at position 231

Domain context: N-terminal cytoplasmic (intrinsically disordered)

ALPHAMISSENSE

- **Pathogenicity score:** 0.7027
- **Class:** likely pathogenic

ALPHAFOLD CONFIDENCE

- **pLDDT at residue 231:** 75.94

> **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.691G>A
 - **ClinVar accession:** VCV001078282
 - **Last evaluated:** 2024/10/26 00:00
 - **Submissions:** 1
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Card generated by `wolfram-atlas-batch` (missense AlphaMissense mint) on 2026-06-08T02:27:33.411475Z.

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