

# V624= — WFS1 Molecular Atlas Card

**Variant type:** Synonymous (silent)

**Codon:** position 624 (Valine, V) — amino acid unchanged

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

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## SCHEMA CATEGORY: SILENT — SILENT — NO AMINO-ACID CHANGE

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No amino-acid change (V624 is unchanged): the codon is altered but the protein sequence is identical to wild-type. No structural, stability or AlphaMissense effect applies. Synonymous variants are typically benign unless they affect splicing or regulatory elements; this one is not adjacent to an exon boundary.

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## CLINICAL EVIDENCE

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- **Classification:** Likely benign
  - **Review status:** criteria provided, single submitter
  - **cDNA change:** c.1872G>T
  - **ClinVar accession:** VCV004751625
  - **Last evaluated:** 2025/03/10 00:00
  - **Submissions:** 1
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*Card generated by wolfram-atlas-batch (synonymous pipeline) on 2026-06-08T02:55:13.119726Z.*

*WFS1: UniProt O76024, AlphaFold v6. Synonymous variants carry no protein-structural effect.*