

# G562= — WFS1 Molecular Atlas Card

**Variant type:** Synonymous (silent)

**Codon:** position 562 (Glycine, G) — amino acid unchanged

**Domain context:** Lumenal loop 4

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

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No amino-acid change (G562 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.1686C>G
- **ClinVar accession:** VCV003004613
- **Last evaluated:** 2023/11/15 00:00
- **Submissions:** 1

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Card generated by *wolfram-atlas-batch* (synonymous pipeline) on 2026-06-08T02:54:40.125866Z.

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