

c.832+1G_A — WFS1 Molecular Atlas Card

Variant type: Splice site

Boundary: donor (5' splice site) · intronic offset +1

Nearest protein position: ~278 (N-terminal cytoplasmic (intrinsically disordered))

SCHEMA CATEGORY: S3 — SPLICEAI COULD NOT SCORE THIS VARIANT — WET-LAB VALIDATION

SpliceAI returned no usable prediction (no genomic coordinate / SpliceAI record). The canonical splice-region call stands; wet-lab RNA validation is required to resolve the isoform.

SPLICE PREDICTION

- **Affected site:** donor (5' splice site), canonical ($\pm 1/\pm 2$ core)
- **SpliceAI:** no genomic coordinate / SpliceAI record

CLINICAL EVIDENCE

Not found in the cached WFS1 ClinVar set ([reference/WFS1clinvarvariants.csv](#))._

Card generated by `wolfram-atlas-batch` (splice pipeline) on 2026-06-08T07:53:13.337833Z.

Schema: `reference/card schemaextension.md` (S1–S3). WFS1: UniProt O76024, AlphaFold v6.