

# WFS1 A43V — Wolframin

Ala→Val p43 IDR AM=0.08 ddg=-1.11 pLDDT=28. ClinVar Conflicting evidence.  
Atlas mechanism: see structural analysis.

## IDENTITY

Variant	A43V (p.Alanine43Valine)
DNA change	c.128C>T
Gene · Protein	WFS1 · Wolframin (890 aa)
UniProt	O76024 · WFS1_HUMAN
ClinVar accession	VCV000166568
Amino acid change	conservative volume increase

## STRUCTURAL CONTEXT

AlphaFold model	AF-O76024-F1, v6
pLDDT at residue 43	<b>28.23</b> <span>BELOW IDR THRESHOLD</span>
Domain	N-terminal intrinsically disordered region (1-86)
Position context	N-terminal IDR
IDR flag	YES — pLDDT 28.23 is below 50 threshold (route to Cat 5)

Position analysis: ARG42 (2.5 Å — R42Q!), PRO44 (2.5 Å), PRO41 (4.9 Å).  
pLDDT 28 deep IDR. Substantial  $\Delta\Delta G$  but untrustworthy. The Atlas's neighbor extraction surfaces this variant's contacts and connects them to the broader multi-variant target landscape.

## COMPUTATIONAL PREDICTIONS

ALPHAMISSENSE

**0.077**am\_class: **LBen** —  
threshold > 0.564DYNAMUT2  $\Delta\Delta G$ **-1.11** kcal/

mol

Destabilising · Job  
177992522666

PLDDT (ALPHAFOLD)

**28.23**

BELOW IDR THRESHOLD

## CLINICAL EVIDENCE

ClinVar classification

**CONFLICTING CLASSIFICATIONS OF PATHOGENICITY**

Review status

criteria provided, conflicting classifications

Last evaluated

2013/11/06 00:00

Inheritance

Conflicting ClinVar classifications.

WFS1 variant landscape

A43V is 1 of ~326 pathogenic-spectrum variants in WFS1 (out of 2,243 in ClinVar)

- (no specific conditions catalogued)

## RESEARCH PATH DECISION TREE

$\Delta\Delta G < 2$  + binding site affected → CATEGORY 3 – docking experiments  $\Delta\Delta G$  2–4 → CATEGORY 2 – pharmacological chaperones  $\Delta\Delta G > 4$  → CATEGORY 1 – gene therapy pLDDT < 50 → CATEGORY 5 – IDR, experimental only Stable fold + functional site hit → CATEGORY 4 – site-specific docking

**Cat 5 IDR — see structural prose.** AlphaMissense below threshold (AM under-call class) but mechanism is structurally identified. Therapeutic strategy: site-directed at contacts identified above, or wet-lab validation if pLDDT borderline/below 50.

Deep IDR adjacent to R42Q.