BUILT WITH CLAUDE · LIFE SCIENCES

Read the evidence. Draft the verdict.

Norn is a variant-interpretation copilot. It drafts ACMG/AMP evidence for a human curator to confirm.

Norn

NOT FOR CLINICAL USE

WHAT NORN IS

Gather, weigh, decree.

Paste one variant. Norn gathers public genomics evidence, adjudicates each ACMG criterion with Claude, and computes a transparent classification.

Urðr *what was*

GATHER

Consequence, frequency, and neighbor evidence from Ensembl VEP, gnomAD v4, and ClinVar.

Verðandi *what is*

WEIGH

Claude adjudicates each criterion against code-computed signals, with reasoning.

Skuld *what shall be*

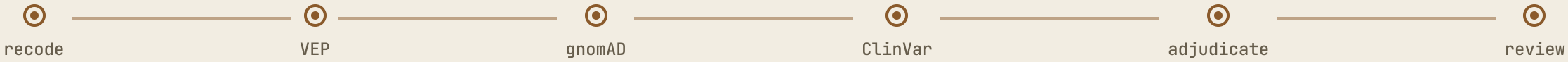
DECREE

The ClinGen points framework combines the verdicts in code. The engine owns the label.

ONE PRINCIPLE

The model justifies. The engine decides.

Claude returns a per-criterion verdict with reasoning; it never returns the final label. The classification is always computed in code, so the same evidence yields the same call.



Norn

A DRAFTED INTERPRETATION

Transparent, sourced, yours to confirm.

`BRCA1:c.5266dupC`

Likely Pathogenic

ACMG POINT AGGREGATION

+9 pts

Benign | Likely benign | VUS | **Likely path.** | Pathogenic

PVS1 (+8) frameshift, PM2 (+1) absent from gnomAD. PM2 is applied at supporting strength, so a classic loss-of-function variant lands at Likely Pathogenic on automated evidence alone.

Spend your time confirming, not gathering.

Live demo: norn-five.vercel.app · Repo: github.com/vignesh-nagarajan-vn/Norn

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