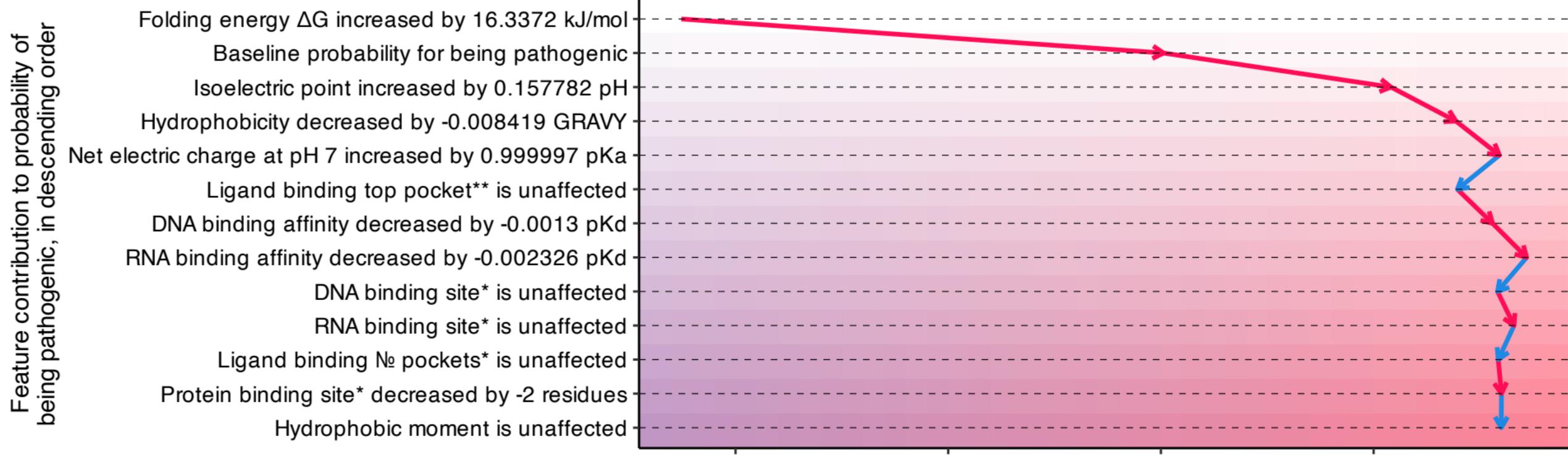


DAVE1 decision plot for GA338R in gene C1RL (Q9NZP8, ENST00000266542), GRCh38 12:7096843C>T

SHAP values do not correlate with feature values, but instead capture feature contributions based on the interactions among all features uniquely for this prediction



* = if any, one or more ** = if any

Cumulative probability heuristic for SHAP values
Final probability for being pathogenic is 0.96, variant is estimated to be pathogenic