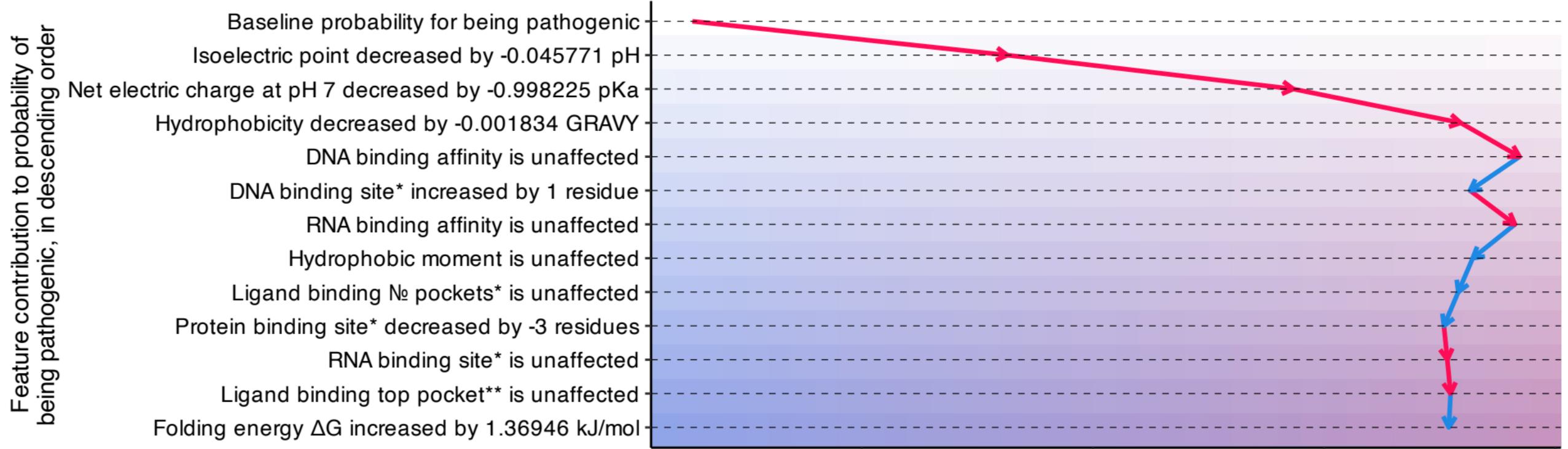


DAVE1 decision plot for GA816E in gene COL4A4 (P53420, ENST00000396625), GRCh38 2:227057537C>T

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 0.554, variant is estimated to be pathogenic