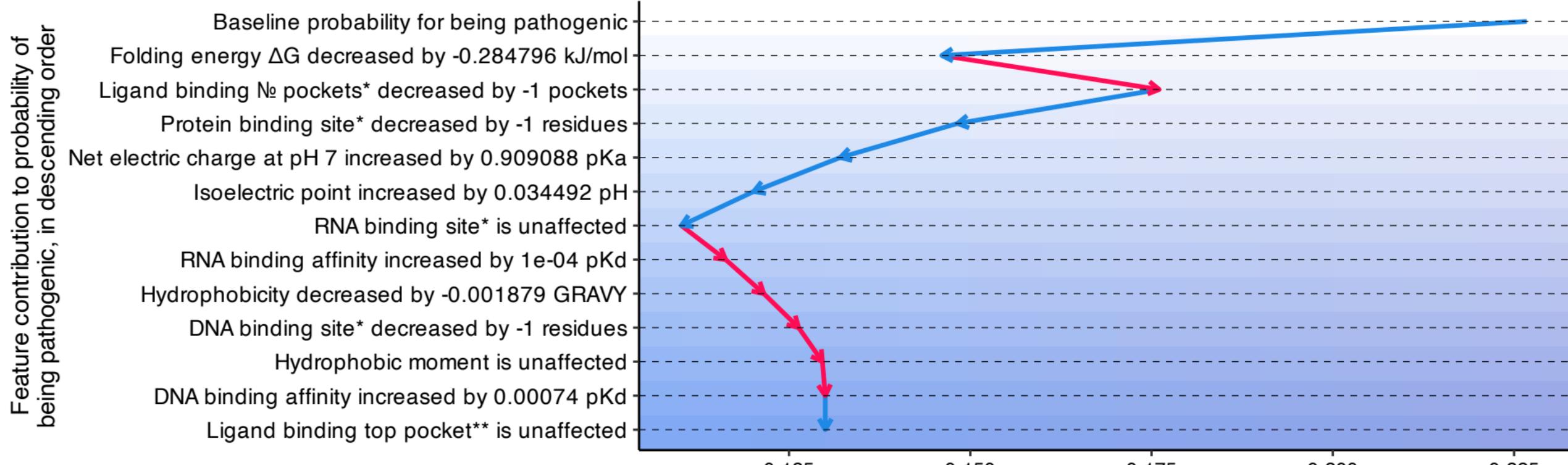


DAVE1 decision plot for HA553R in gene PCSK9 (Q8NBP7, ENST00000302118), GRCh38 1:55059640A>G

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 0.13, variant is estimated to be benign