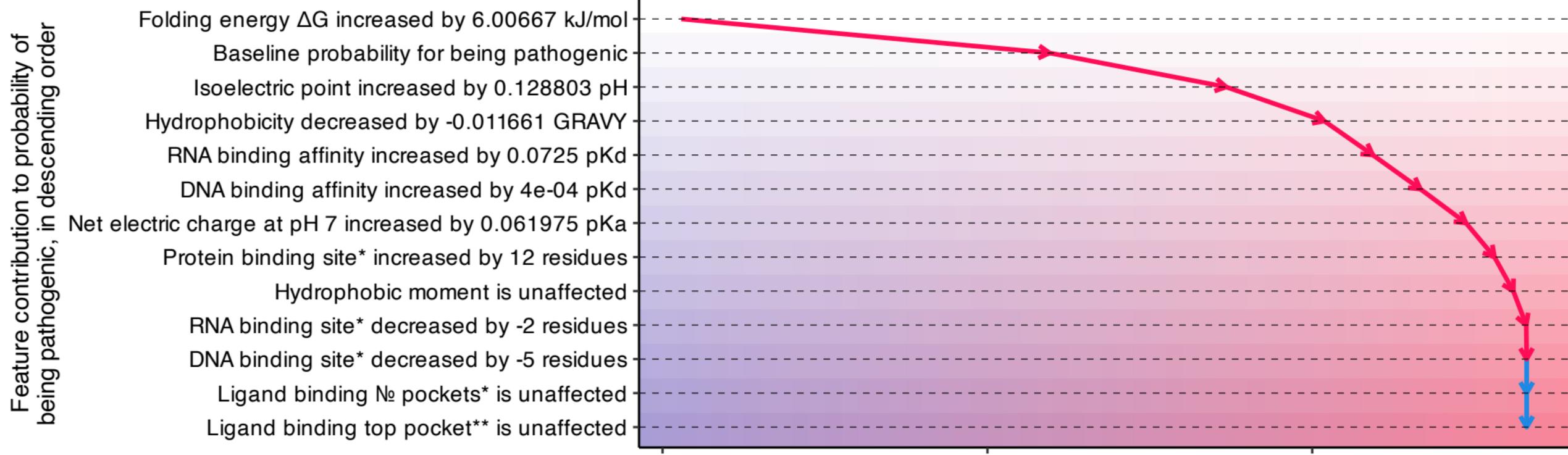


DAVE1 decision plot for CA60S in gene GJB1 (P08034, ENST00000361726), GRCh38 X:71223885T>A

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 0.932, variant is estimated to be pathogenic