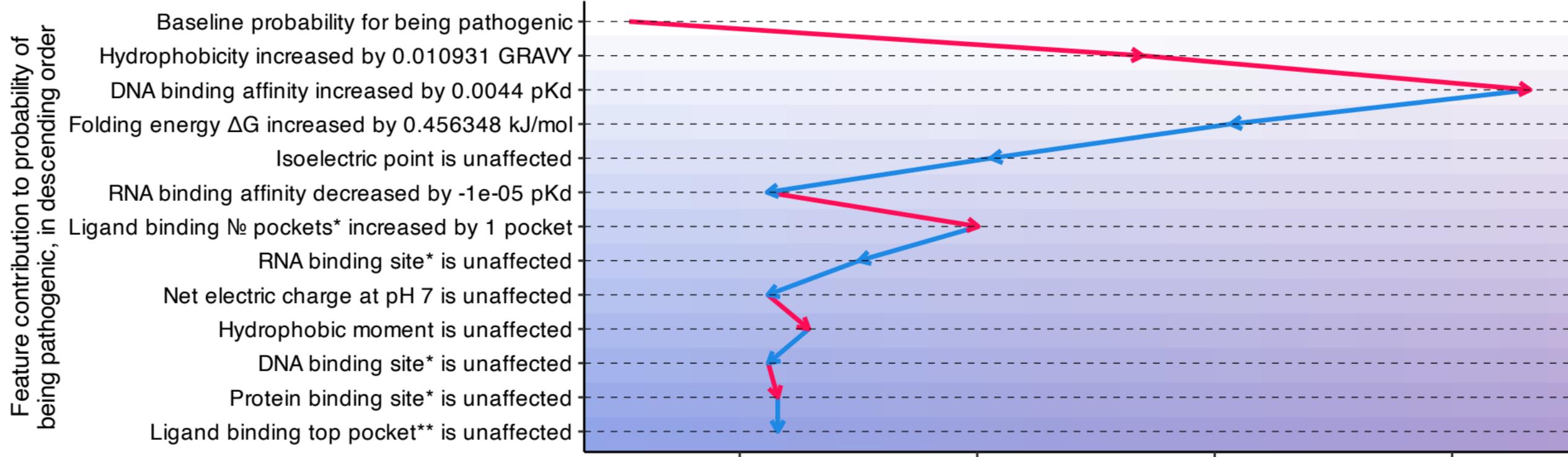


# DAVE1 decision plot for NA54S in gene PRSS1 (P07477, ENST00000311737), GRCh38 7:142750675A>G

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 0.258, variant is estimated to be benign