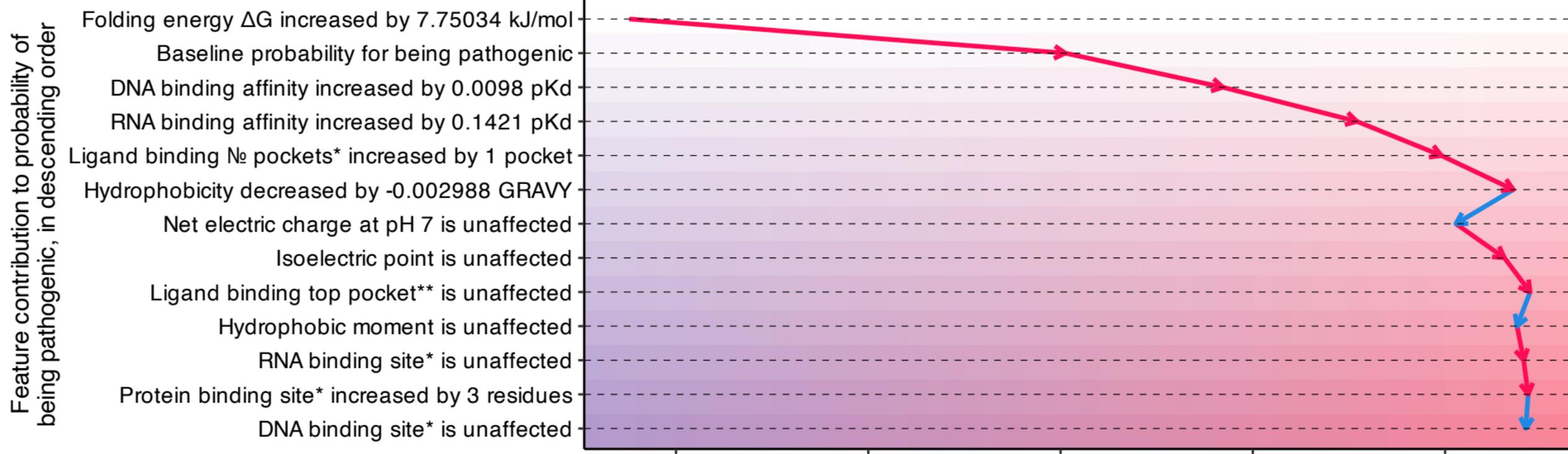


DAVE1 decision plot for LA105P in gene TSC2 (P49815), GRCh38 16:2053430T>C, VKGL April 2024: VUS

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 0.942, variant is estimated to be pathogenic