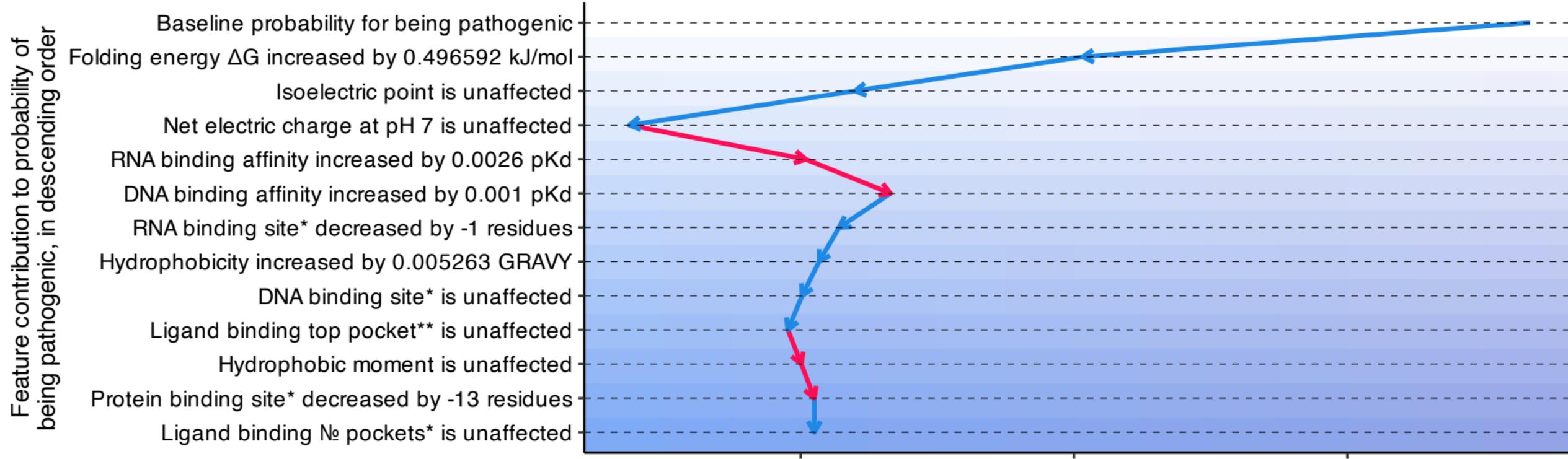


SHAP decision plot for TA379A in gene LPL (P06858), GRCh38 8:19959376A>G, VKGL April 2024: VUS

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 0.122, variant is estimated to be benign