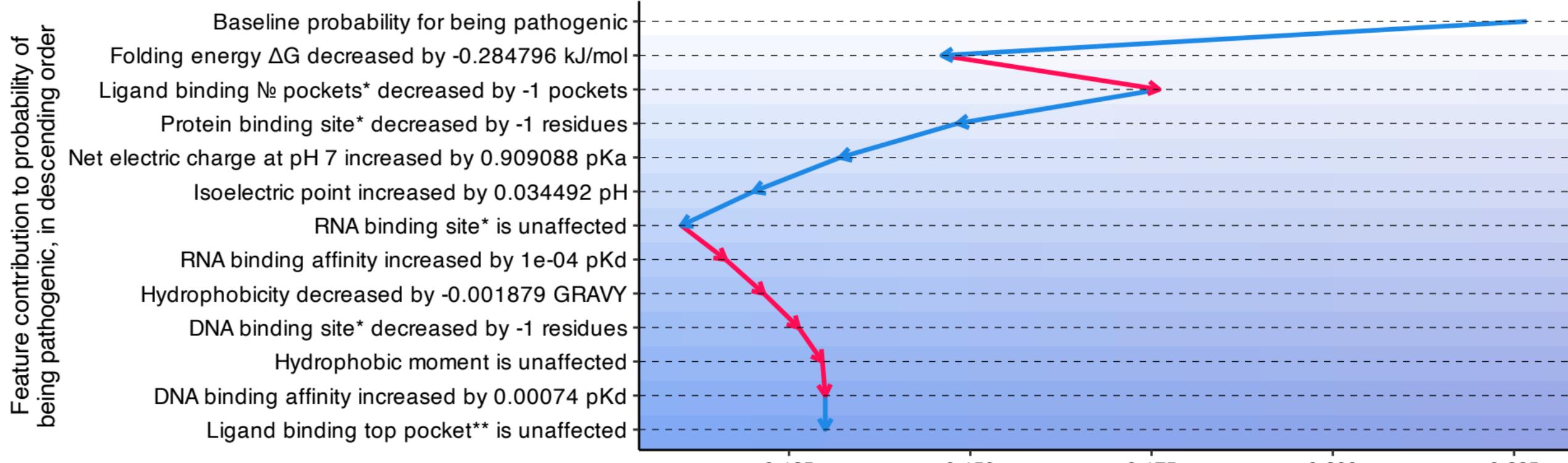


SHAP decision plot for HA553R in gene PCSK9 (Q8NBP7), GRCh38 1:55059640A>G, VKGL April 2024: VUS

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 0.13, variant is estimated to be benign