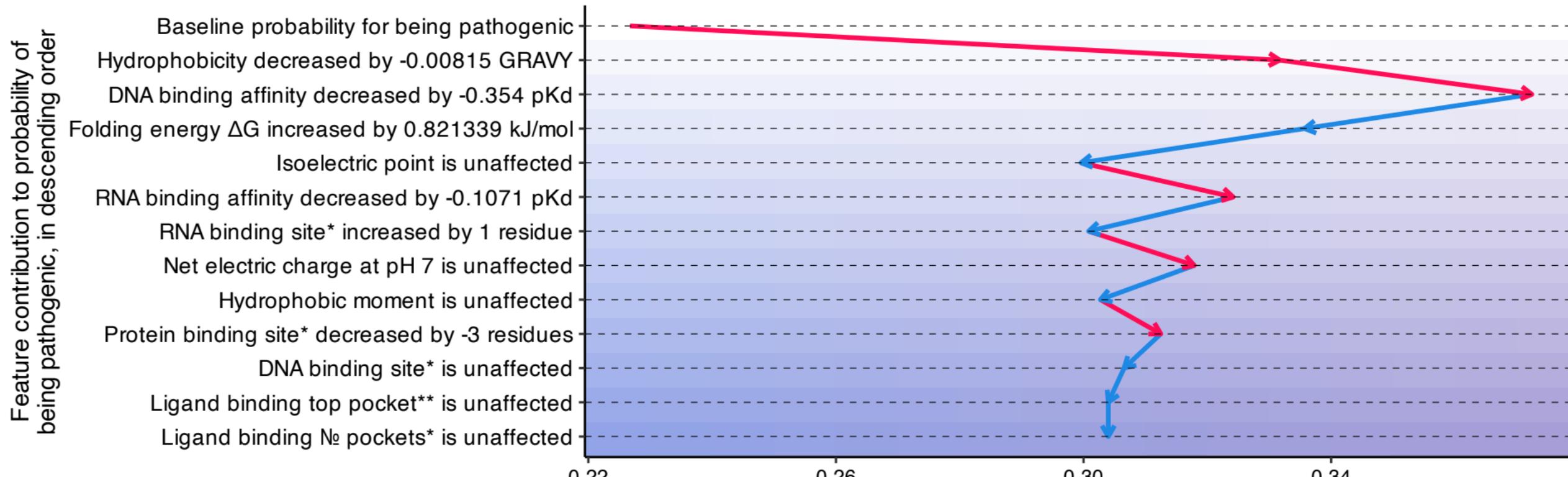


# SHAP decision plot for MA273T in gene ANKRD1 (Q15327), GRCh38 10:90915574A>G, VKGL April 2024: VUS

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 0.304, variant is estimated to be pathogenic