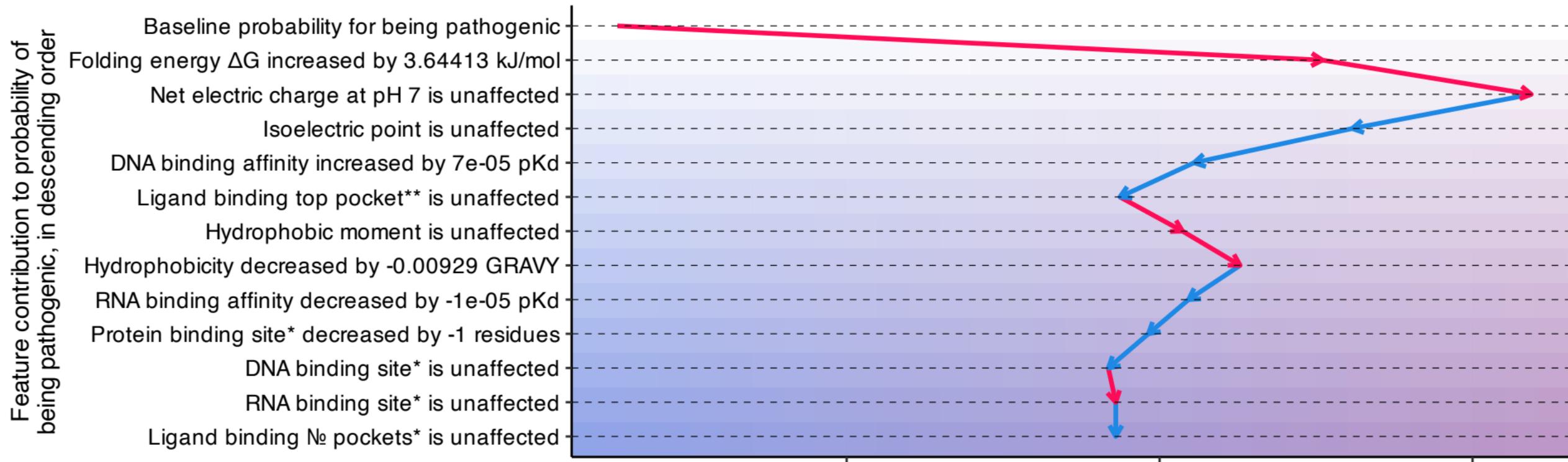


# SHAP decision plot for AA302P in gene APOA5 (Q6Q788), GRCh38 11:116790325C>G, VKGL April 2024: VUS

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 0.386, variant is estimated to be pathogenic