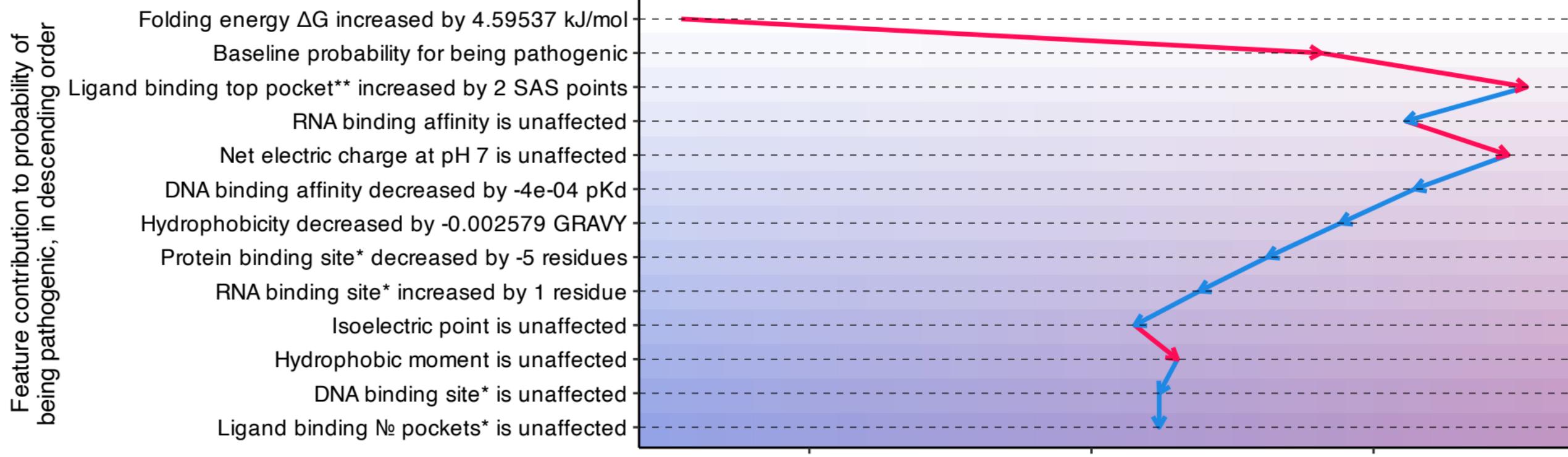


SHAP decision plot for TA111P in gene WNT7B (P56706), GRCh38 22:45931337T>G, VKGL April 2024: VUS

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 0.424, variant is estimated to be pathogenic