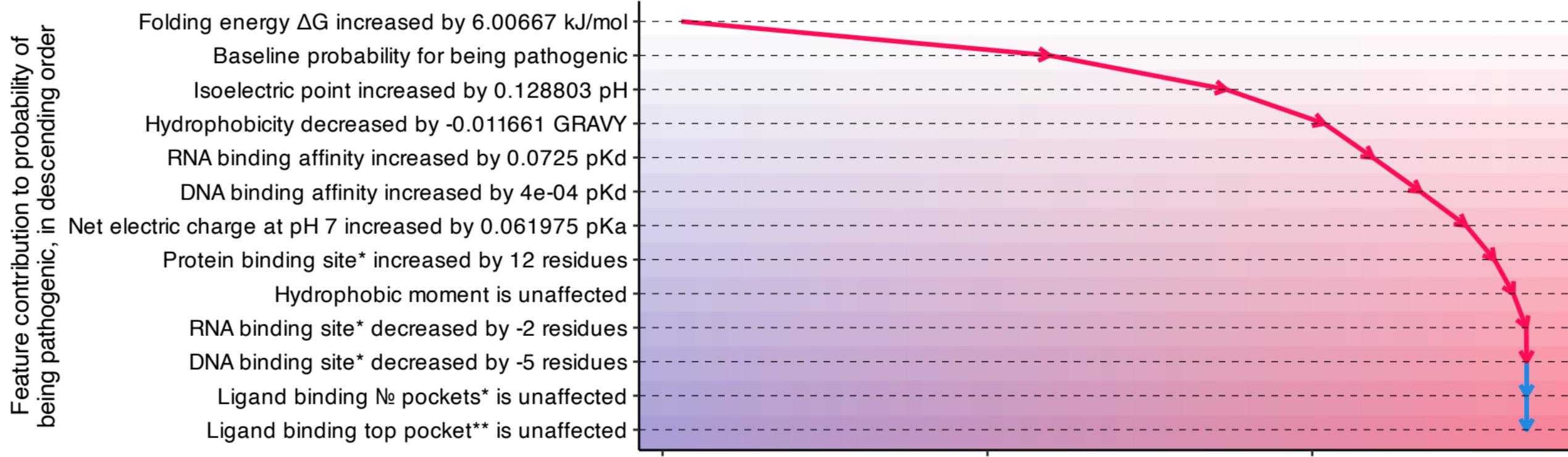


# SHAP decision plot for CA60S in gene GJB1 (P08034), GRCh38 X:71223885T>A, VKGL April 2024: VUS

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 0.932, variant is estimated to be pathogenic