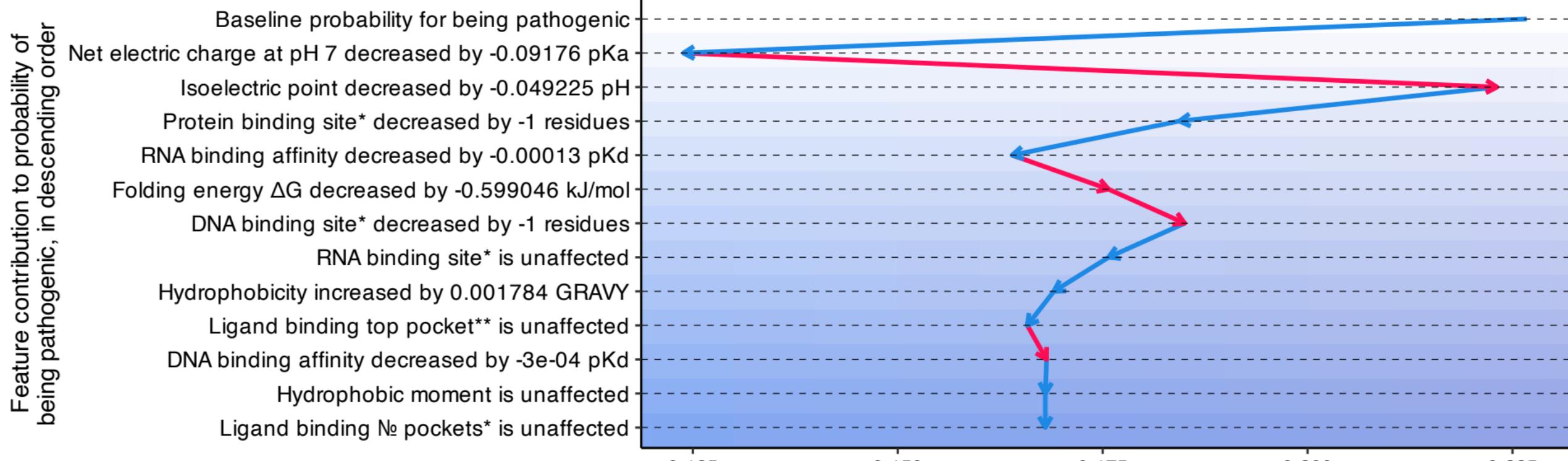


# SHAP decision plot for HA108Y in gene CP (P00450), GRCh38 3:149212523G>A, VKGL April 2024: VUS

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

Final probability for being pathogenic is 0.168, variant is estimated to be benign