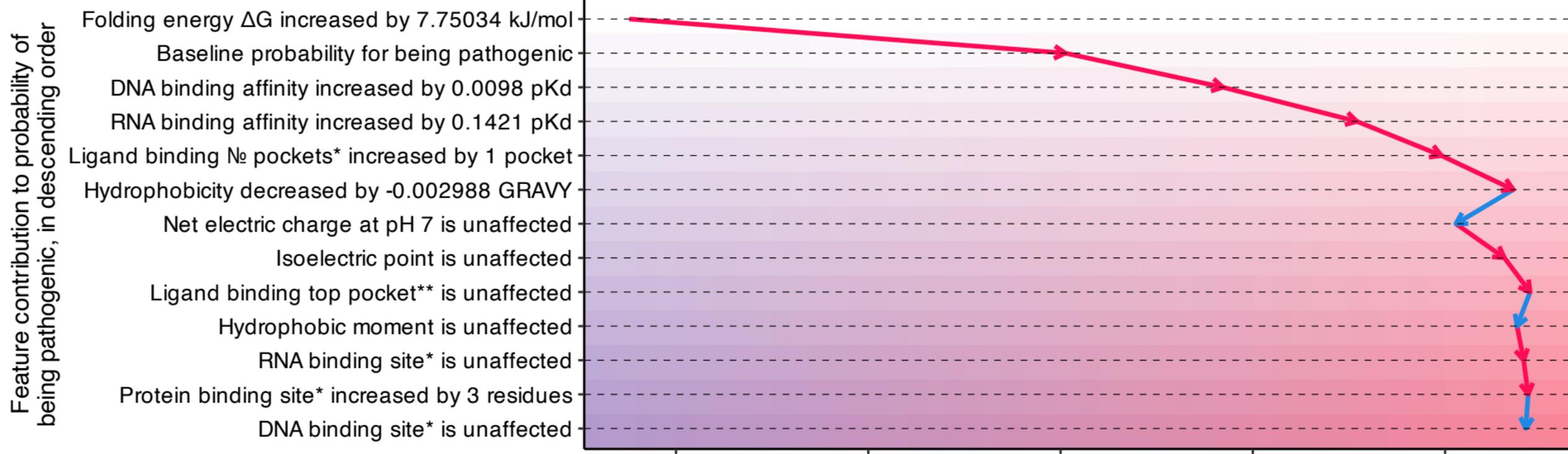


# DAVE1 decision plot for LA105P in gene TSC2 (P49815, ENST00000219476), GRCh38 16:2053430T>C

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 0.942, variant is estimated to be pathogenic