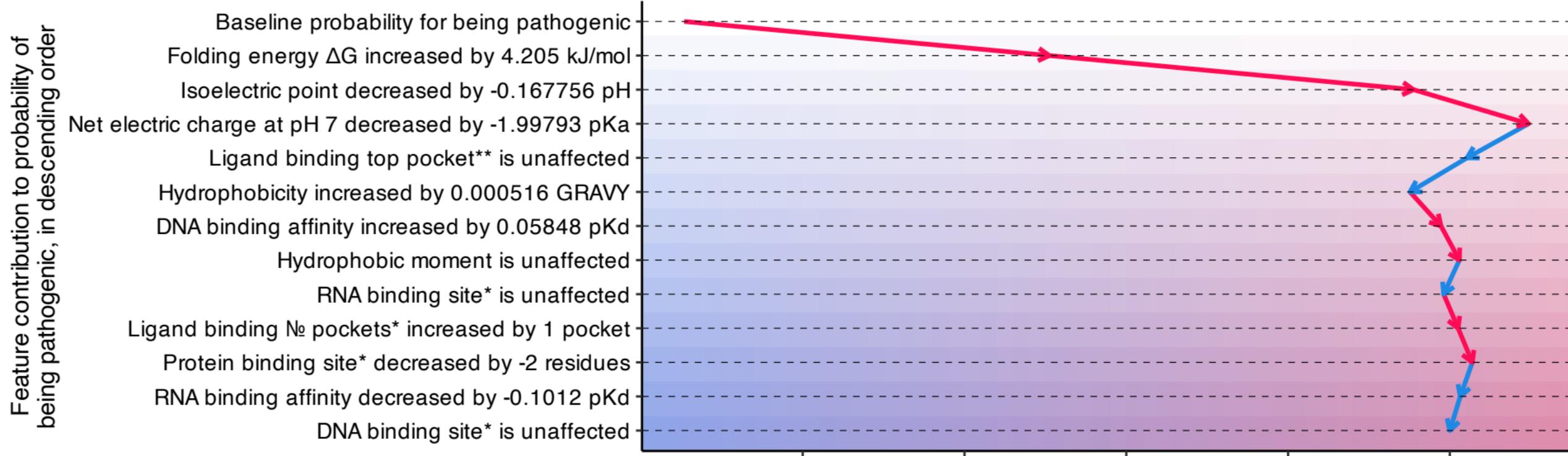


# DAVE1 decision plot for KA771E in gene ECEL1 (O95672, ENST00000304546), GRCh38 2:232480170T>C

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

Cumulative probability heuristic for SHAP values  
Final probability for being pathogenic is 0.7, variant is estimated to be pathogenic