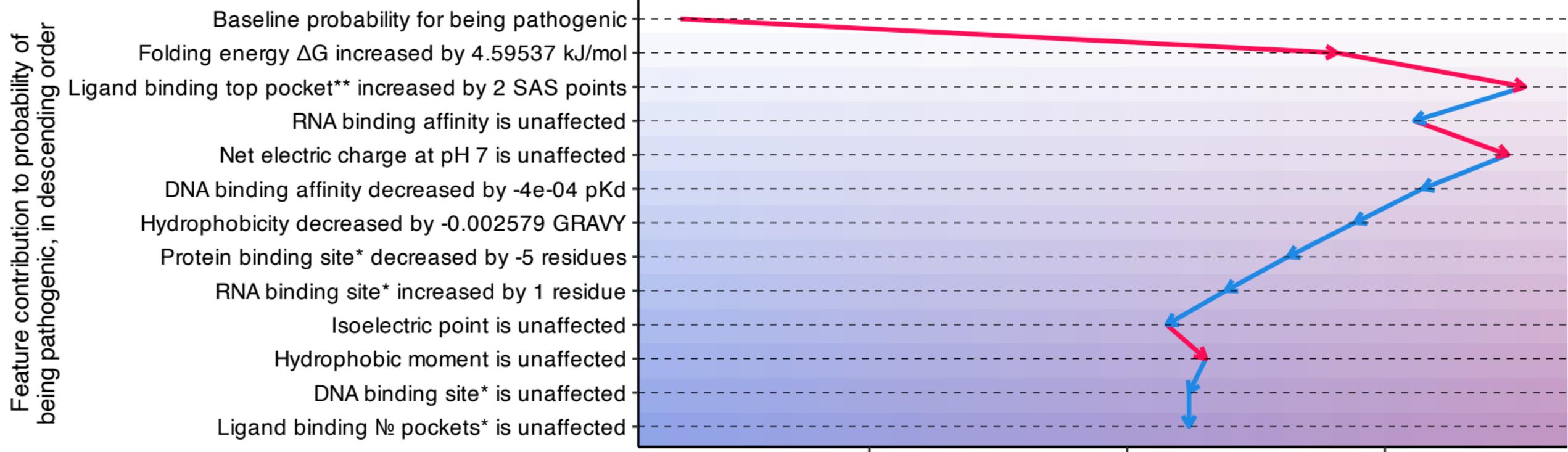


DAVE1 decision plot for TA111P in gene WNT7B (P56706, ENST00000339464), GRCh38 22:45931337T>G

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, benign/path. threshold is 0.286
Final probability for being pathogenic is 0.424, variant is estimated to be pathogenic