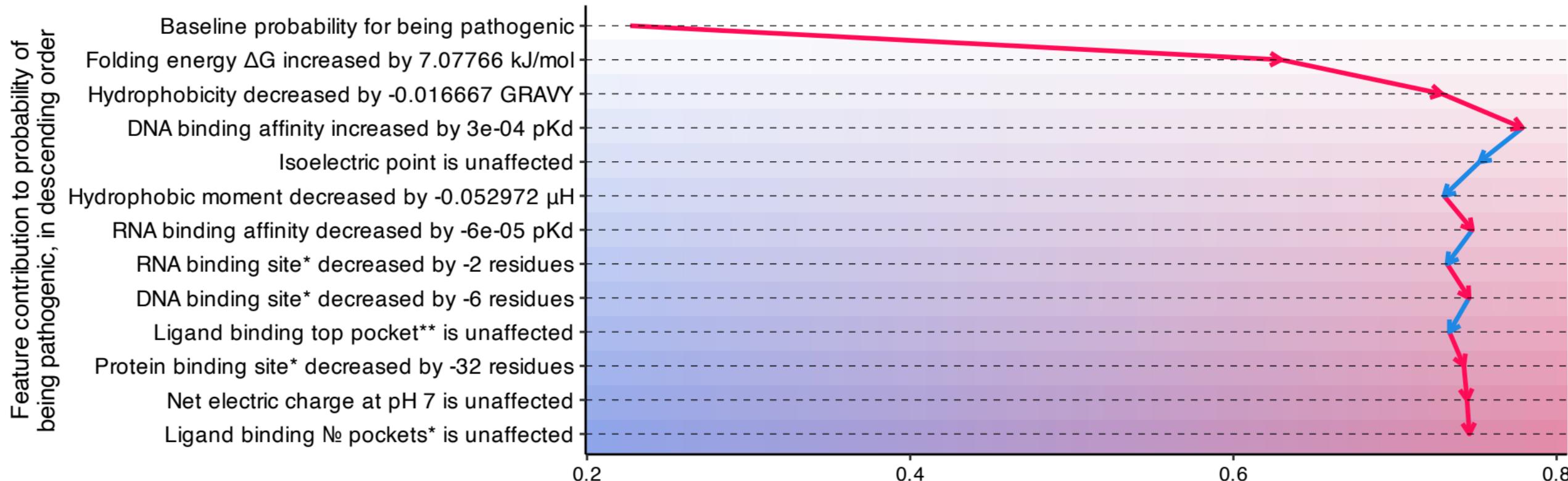


DAVE1 decision plot for LA153P in gene NKX2-5 (P52952, ENST00000329198), GRCh38 5:173233086A>G

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 0.746, variant is estimated to be pathogenic