SHAP decision plot for LA78F in gene ADAMTSL4 (Q6UY14), GRCh38 1:150553051C>T, 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

Cumulative probability heuristic for SHAP values

Final probability for being pathogenic is 0.03