

Biomarker-REG_KIDA_JSWmean_wholejoint

High

Feature value

Low

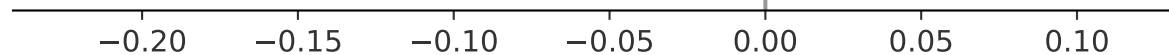
S_COMP

U_CTXII_CREA

S_COLL2_1NO2

S_COLL2_1

U_CTXIALPHA_CREA



SHAP value (impact on model output)