

Biomarker-KIDA_BD_wholejoint

High

Feature value

Low

U_CTXII_CREA

S_COLL2_1NO2

S_COLL2_1

S_CTXI

S_COMP

S_HA

S_C10C

S_hsPRO_C2

S_NMID

S_C3M

-0.100 -0.075 -0.050 -0.025 0.000 0.025 0.050 0.075 0.100

SHAP value (impact on model output)

