

Biomarker-KIDA_OP_wholejoint

High

Feature value

Low

U_CTXII_CREA

S_COMP

S_hsPRO_C2

S_HA

S_NMID

-0.25

-0.20

-0.15

-0.10

-0.05

0.00

0.05

0.10

SHAP value (impact on model output)

