

Biomarker-structural_REG_BML

High

Feature value

Low

U_CTXII_CREA

S_HA

S_COMP

S_COLL2_1NO2

S_hsPRO_C2

-0.15

-0.10

-0.05

0.00

0.05

0.10

0.15

0.20

SHAP value (impact on model output)

