

Biomarker-structural_REG_all

High

Feature value

Low

U_CTXII_CREA

S_C10C

S_NMID

S_COLL2_1

S_hsPRO_C2

-0.20

-0.15

-0.10

-0.05

0.00

0.05

0.10

SHAP value (impact on model output)

