

Biomarker-boneMRI\_ShapeFermur

High

Feature value

Low

U\_CTXII\_CREA

S\_C10C

S\_hsCRP

S\_HA

S\_COLL2\_1

-0.20

-0.15

-0.10

-0.05

0.00

0.05

0.10

0.15

SHAP value (impact on model output)

