

Biomarker-MOAKS\_BMLsize\_wholejoint

High

Feature value

Low

S\_HA

U\_CTXII\_CREA

S\_hsCRP

S\_CRPM

S\_COLL2\_1

-0.15

-0.10

-0.05

0.00

0.05

0.10

0.15

SHAP value (impact on model output)

