

Biomarker-MOAKS\_BMLnum\_wholejoint

High

Feature value

Low

U\_CTXII\_CREA

S\_CTXI

S\_COLL2\_1NO2

S\_COMP

S\_C3M

-0.05

0.00

0.05

0.10

0.15

0.20

SHAP value (impact on model output)

