

Biomarker-REG\_MOAKS\_BMLnum\_Pat

High

Feature value

Low

S\_HA

S\_C3M

S\_hsPRO\_C2

U\_CTXII\_CREA

S\_COMP

-0.15

-0.10

-0.05

0.00

0.05

0.10

0.15

0.20

SHAP value (impact on model output)

