

Biomarker-MOAKS\_BMLnum\_Pat

High

Feature value

Low

S\_HA

U\_CTXIALPHA\_CREA

S\_hsCRP

S\_RE\_C1M

S\_hsPRO\_C2

-0.25 -0.20 -0.15 -0.10 -0.05 0.00 0.05 0.10 0.15

SHAP value (impact on model output)

