

Biomarker-REG\_MOAKS\_BMLnum\_MAC

High

S\_HA

S\_C10C

U\_CTXIALPHA\_CREA

S\_COLL2\_1NO2

S\_COMP

Feature value

Low

-0.4

-0.3

-0.2

-0.1

0.0

0.1

SHAP value (impact on model output)

