

Biomarker-REG\_MOAKS\_BMLsize\_MAC

High

Feature value

Low

S\_HA

U\_CTXII\_CREA

S\_COLL2\_1

S\_COLL2\_1NO2

S\_C3M

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

SHAP value (impact on model output)

