

Biomarker-REG\_KIDA\_JSWmean\_MAC

High

Feature value

Low

S\_RE\_C1M

U\_CTXII\_CREA

S\_COMP

S\_HA

S\_COLL2\_1

-0.3

-0.2

-0.1

0.0

0.1

SHAP value (impact on model output)

