

Biomarker-REG\_MOAKS\_OP\_MAC

High

Feature value

Low

U\_CTXII\_CREA

S\_CRPM

S\_RE\_C1M

S\_HA

S\_NMID

S\_hsCRP

S\_COMP

S\_C2M

S\_C3M

S\_C10C

-0.05

0.00

0.05

0.10

0.15

0.20

SHAP value (impact on model output)

