

Biomarker-REG\_MOAKS\_CartFull\_MAC

High

Feature value

Low

S\_NMID

S\_CTXI

S\_C3M

S\_C10C

S\_hsPRO\_C2

-0.10

-0.05

0.00

0.05

0.10

0.15

0.20

0.25

SHAP value (impact on model output)

