

Biomarker-MOAKS_CartFull_MAC

High

Feature value

Low

S_NMID

S_CTXI

U_CTXII_CREA

S_COLL2_1NO2

S_C3M

-0.20

-0.15

-0.10

-0.05

0.00

0.05

0.10

0.15

SHAP value (impact on model output)

