

Biomarker-MOAKS_BMLsize_TF

High

Feature value

Low

S_HA

U_CTXII_CREA

S_hsCRP

S_CRPM

S_COLL2_1

-0.15

-0.10

-0.05

0.00

0.05

0.10

0.15

SHAP value (impact on model output)

