

Biomarker-MOAKS_BMLnum_wholejoint

High

Feature value

Low

U_CTXII_CREA

S_CTXI

S_COLL2_1NO2

S_COMP

S_C3M

-0.20

-0.15

-0.10

-0.05

0.00

0.05

SHAP value (impact on model output)

