

Biomarker-REG_MOAKS_BMLsize_wholejoint

High

Feature value

Low

S_HA

U_CTXII_CREA

S_COLL2_1

S_C2M

S_C10C

S_COLL2_1NO2

S_COMP

S_hsCRP

S_NMID

S_C3M

-0.15

-0.10

-0.05

0.00

0.05

0.10

SHAP value (impact on model output)

