

Biomarker-REG_MOAKS_OP_wholejoint

High

Feature value

Low

U_CTXIALPHA_CREA

U_CTXII_CREA

S_C10C

S_hsCRP

S_NMID

-0.15 -0.10 -0.05 0.00 0.05 0.10 0.15 0.20 0.25

SHAP value (impact on model output)

