

Biomarker-MOAKS_OP_wholejoint

High

Feature value

Low

U_CTXII_CREA

S_C10C

S_COLL2_1

S_COMP

S_NMID

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

SHAP value (impact on model output)

