

Biomarker-KIDA_BD_MAC

High

Feature value

Low

S_hsPRO_C2

U_CTXII_CREA

S_COLL2_1

S_COLL2_1NO2

S_NMID

-0.2

-0.1

0.0

0.1

0.2

SHAP value (impact on model output)

