

Biomarker-REG_KIDA_BD_MAC

High

Feature value

Low

U_CTXII_CREA

S_COLL2_1

S_COLL2_1NO2

S_C10C

S_HA

S_NMID

S_CRPM

S_C3M

S_COMP

S_hsPRO_C2

-0.05

0.00

0.05

0.10

0.15

SHAP value (impact on model output)

