

Biomarker-REG_KIDA_BD_MAC

High

Feature value

Low

S_CTXI

S_COMP

S_C10C

S_RE_C1M

S_HA

U_CTXII_CREA

S_NMID

S_COLL2_1NO2

S_COLL2_1

S_CRPM

-0.100 -0.075 -0.050 -0.025 0.000 0.025 0.050 0.075

SHAP value (impact on model output)

