

Biomarker-REG_MOAKS_BMLnum_MAC

High

Feature value

Low

S_COMP

U_CTXII_CREA

U_CTXIALPHA_CREA

S_C10C

S_COLL2_1NO2

S_NMID

S_CTXI

S_HA

S_C3M

S_CRPM

-0.05

0.00

0.05

0.10

0.15

0.20

SHAP value (impact on model output)

