

Biomarker-REG_MOAKS_BMLsize_MAC

High

Feature value

Low

S_HA

U_CTXII_CREA

S_COLL2_1

S_COLL2_1NO2

S_C3M

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

SHAP value (impact on model output)

