

Biomarker-REG_MOAKS_BMLnum_MAC

High

Feature value

Low

S_HA

S_C10C

U_CTXIALPHA_CREA

S_COLL2_1NO2

S_COMP

-0.1 0.0 0.1 0.2 0.3 0.4

SHAP value (impact on model output)

