

Biomarker-REG_MOAKS_BMLnum_Pat

High

Feature value

Low

S_HA

S_C10C

S_COLL2_1

U_CTXIALPHA_CREA

S_hsCRP

-0.10 -0.05 0.00 0.05 0.10 0.15 0.20

SHAP value (impact on model output)

