

Biomarker-REG_MOAKS_BMLnum_Pat

High

Feature value

Low

S_HA

S_C10C

S_COLL2_1

U_CTXIALPHA_CREA

S_hsCRP

-0.20

-0.15

-0.10

-0.05

0.00

0.05

0.10

SHAP value (impact on model output)

