

Biomarker-REG\_MOAKS\_BMLnum\_wholejoint

High

Feature value

Low

S\_HA

U\_CTXIALPHA\_CREA

S\_C10C

S\_CTXI

S\_COLL2\_1NO2

S\_COLL2\_1

S\_COMP

U\_CTXII\_CREA

S\_CRPM

S\_C3M

-0.25 -0.20 -0.15 -0.10 -0.05 0.00 0.05 0.10

SHAP value (impact on model output)

