

Biomarker-REG\_MOAKS\_CartAny\_MAC

High

Feature value

Low

U\_CTXIALPHA\_CREA

S\_hsCRP

S\_ARGS\_LUND

S\_HA

S\_COMP

S\_C3M

S\_COLL2\_1

S\_COLL2\_1NO2

S\_NMID

U\_CTXII\_CREA

-0.10 -0.05 0.00 0.05 0.10

SHAP value (impact on model output)

