

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)

