

Biomarker-REG_MOAKS_CartFull_wholejoint

High

Feature value

Low

S_CTXI

U_CTXII_CREA

S_COMP

U_CTXIALPHA_CREA

S_NMID

S_COLL2_1

S_C3M

S_COLL2_1NO2

S_CRPM

S_hsPRO_C2

-0.08 -0.06 -0.04 -0.02 0.00 0.02 0.04 0.06 0.08

SHAP value (impact on model output)

