

Biomarker-REG_MOAKS_CartAny_wholejoint

High

Feature value

Low

S_COLL2_1

S_NMID

S_C10C

S_hsCRP

S_COMP

S_HA

S_RE_C1M

S_CRPM

U_CTXIALPHA_CREA

S_CTXI

-0.05

0.00

0.05

0.10

SHAP value (impact on model output)

