

Biomarker-REG_KIDA_JSWmean_wholejoint

High

Feature value

Low

S_COMP

U_CTXII_CREA

S_COLL2_1NO2

S_COLL2_1

U_CTXIALPHA_CREA

-0.10

-0.05

0.00

0.05

0.10

0.15

0.20

SHAP value (impact on model output)

