

Biomarker-KIDA\_OP\_wholejoint

High

Feature value

Low

U\_CTXII\_CREA

S\_COMP

S\_hsPRO\_C2

S\_HA

S\_NMID

-0.10 -0.05 0.00 0.05 0.10 0.15 0.20 0.25

SHAP value (impact on model output)

