

Biomarker-REG\_KIDA\_OP\_MAC

High

Feature value

Low

U\_CTXII\_CREA

S\_hsPRO\_C2

S\_NMID

S\_COMP

S\_COLL2\_1

-0.15

-0.10

-0.05

0.00

0.05

0.10

0.15

SHAP value (impact on model output)

