

Biomarker-REG_MOAKS_Effusion

High

Feature value

Low

S_C3M

S_COMP

S_hsPRO_C2

S_RE_C1M

S_COLL2_1

-0.15

-0.10

-0.05

0.00

0.05

0.10

0.15

0.20

0.25

SHAP value (impact on model output)

