

MRI-KIDA_BD_wholejoint

High

Feature value

Low

LFTC.S.T2.Me

eMT.ThCtAB.aMe

cLF.ThCtAB.aMav

b_score

MT.S.T2.Me

-0.15 -0.10 -0.05 0.00 0.05

SHAP value (impact on model output)

