

all-Bone

High

Feature value

Low

KL\_Score\_K\_LAS

b\_score

Sumscore\_K

JSW\_Min\_K

F\_midEpi\_Lat\_Global\_Inhomo\_K

F\_Juxtaphy\_Lat\_Global\_Inhomo\_K

KOOS\_ADL

cLF.D.T2.Sd

T\_MidEpi\_Lat\_BMD\_N\_Menis\_K

LFTC.D.T2.Me

KOOS\_Total

LT.D.T2.Sd

Density\_Mean\_Femur\_K

cMT.ThCtAB.aMe

MT.ThCtAB.aMav

-0.10 -0.05 0.00 0.05 0.10

SHAP value (impact on model output)

