

all-boneMRI_ShapeFemur

High

Feature value

Low

KL_Score_K_LAS

MOSFLC_K

KL_Score_K_ML

T_subchEpi_Lat_Global_Inhomo_K

2D_Shape_Mode_12

T_subchEpi_Tot_Global_Inhomo_K

U_CTXII_CREA

cMT.ThCtAB.aMe

MT.ThCtAB.aMav

F_Cort_Med_BMD_K

-0.15 -0.10 -0.05 0.00 0.05 0.10 0.15

SHAP value (impact on model output)

