

Tomography-REG_MOAKS_BMLnum_Pat

High

Feature value

Low

T_MidEpi_Med_BMD_N_Menis_K

OA_Sum_Body

T_midEpi_Lat_Entropy_K

T_Juxtaphy_Tot_Global_Inhomo_K

T_SubchEpi_Med_BMD_K

-0.10

-0.05

0.00

0.05

0.10

0.15

0.20

SHAP value (impact on model output)

