

HALLMARK\_INFLAMMATORY\_RESPONSE

0.528

UplnMDS\_StrictCutoff

0.477

ASXL1\_Mutated

0.219

Monocyte.like

0.196

HSC.like

0.191

Myeloid.like

0.154

TP53\_Mutated

0.150

cDC.like

0.140

HSC.Prog.like

0.134

Progenitor.like

0.133

FLT3\_ITD\_Mutated

0.123

RUNX1\_Mutated

0.109

CBFB\_MYH11\_Present

0.057

SHAP value (impact on model output)

Feature value

Low

High

