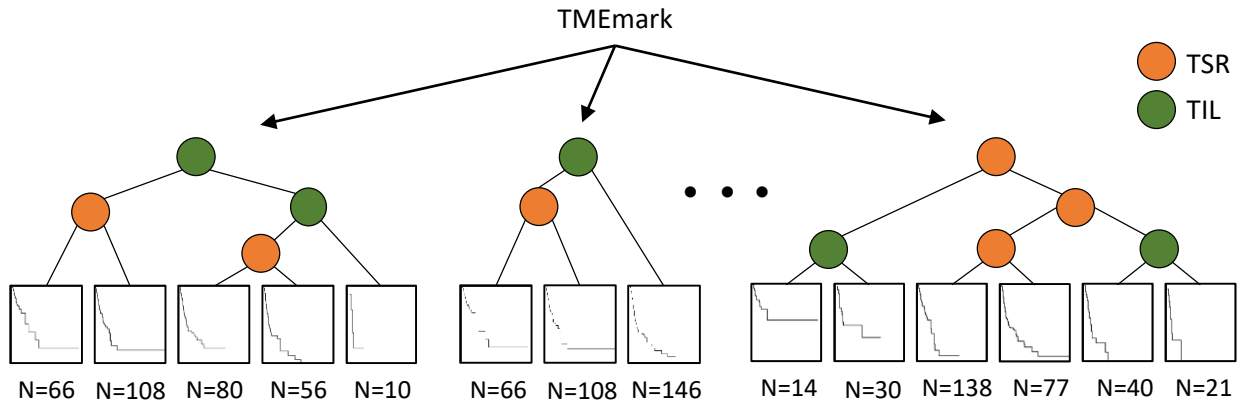


Survival random forest using continuous TSR and TIL
(Each of 1,000 trees with maximum depth of 3)



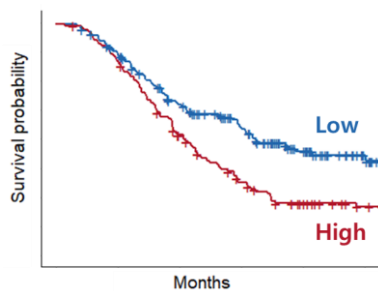
Identify all candidate cutoffs for each TMemark
corresponding to all root and internal nodes

Choose Top 5 frequently identified
candidate cutoffs for each TMemark

Evaluate C-index for all single TMemark-based subtypes
defined by the chosen cutoffs
using univariable cox regression model

Select the single TMemark-based subtype
with highest C-index

TSR-based subtype



Single-TIL-based subtype

