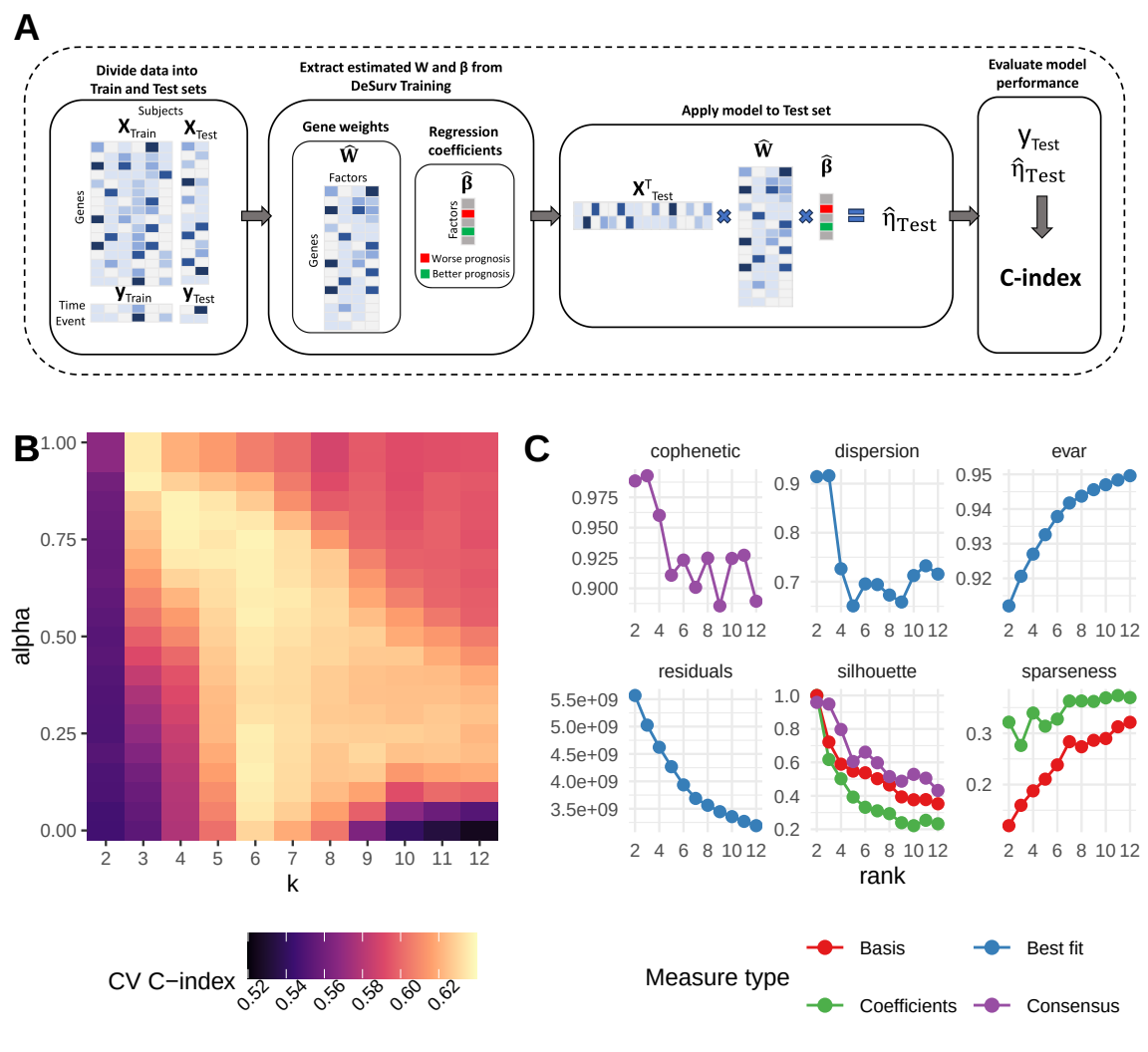
A
Divide data into Train and Test sets
Subjects
Genes
Time Event
Extract estimated W and β from DeSurv Training
Gene weights
Factors
Genes
Regression coefficients
Factors
Worse prognosis
Better prognosis
Apply model to Test set
Evaluate model performance
C-index
B
alpha
k
CV C−index
C
cophenetic
dispersion
evar
residuals
silhouette
sparseness
rank
Measure type
Basis
Best fit
Coefficients
Consensus