

Summary Table

#Genes	#Samples
2515	89
2515	74
2515	105

Methods for Meta KTSP

Mean score ▾

Max number of top scoring pairs (K)

9 ▾


Number of cores for parallel computing

2 ▾

Please select TWO labels to cluster


Please select studies for training

Please select ONE study for testing

 Train model

Number of top scoring pairs (K)

9 ▾

 Predict

Gene pair table

K diagnostic plot