Working Directory
/Users/Masaki/Dropbo

Active Study
Demo

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

inv(16) t(8;21)

Please select studies for training

study1.csv study2.csv

Please select ONE study for testing

study3.csv

🚀 Train model

Number of top scoring pairs (K)

9

🚀 Predict

Gene pair table

GeneIndex1	GeneIndex2	Gene1	Gene2	Score_overall	Score_study1	Score_study2
102	531	RNASE3	KDM4B	-1.97	-0.97	-1.00
256	328	ANPEP	P2RX5	-1.94	-0.94	-1.00
126	432	LST1	TM7SF3	-1.94	-0.94	-1.00
25	326	VEGFA	ZBTB18	1.94	0.97	0.96
281	833	RPUSD3	SLC43A3	1.94	0.97	0.96
14	239	CD9	HSPG2	-1.91	-0.91	-1.00
151	216	RGS10	AIF1	1.91	0.91	1.00
213	481	GLIPR2	SORD	-1.90	-0.94	-0.96
109	169	CD96	KCNK17	1.90	0.97	0.93

K diagnostic plot