



	c) ARI scores on bulk RNA-seq datasets	
	BRCA	KIRP
Discover + GMM(*)	0.68	0.17
Discover + HDBSCAN	0.57	0.0
Affinity Propagation	0.02	0.02
Kmeans (*)	0.39	0.0
GMM (*)	0.65	0.01
Spectral Clustering (*)	0.0	0.0
HDBSCAN	0.11	0.07
PCA+ Kmeans (*)	0.27	0.0
PCA + GMM (*)	0.35	0.04
PCA + HDBSCAN	0.0	0.0
Xgboost features + GMM (*)	0.71	0.38
Xgboost features + HDBSCAN	0.3	0.13
MI features + GMM (*)	0.7	0.3
MI features + HDBSCAN	0.53	0.17