Cell Cluster Confusion Matrix, ARI:0.98 Precision: 0.99 Recall: 0.98 9 -8 predicted label 3 -0 -Ó true label Mutation Cluster Confusion Matrix, ARI:0.59 Precision: 0.61 Recall: 0.71 9 -[118] predicted label [37] **[55]** [219] [1] i true label