Cell Cluster Confusion Matrix, ARI:0.55 Precision: 0.47 Recall: 1.00 [343] 8 predicted label 4 -Ö true label Mutation Cluster Confusion Matrix, ARI:0.46 Precision: 0.45 Recall: 0.74 8 -[203] 7 predicted label [30] [1] [55] 3 -0 -i true label