Cell Cluster Confusion Matrix, ARI:0.80 Precision: 0.81 Recall: 0.87 1038 4 -845 3 predicted label 366 132 [382] [1049] 810 0 -Ö 3 true label Mutation Cluster Confusion Matrix, ARI:0.87 Precision: 0.90 Recall: 0.90 9 400 1 1 258 predicted label 3 37 6 115 1 40 361 1 -243 0 -Ó 3 true label