Cell Cluster Confusion Matrix, ARI:0.64 Precision: 0.75 Recall: 0.69 9 -8 predicted label **-**true label Mutation Cluster Confusion Matrix, ARI:0.36 Precision: 0.47 Recall: 0.44 9 -[101] [224] 8 -[121] predicted label [121] [77][22] [182] [7] **-**0 -true label