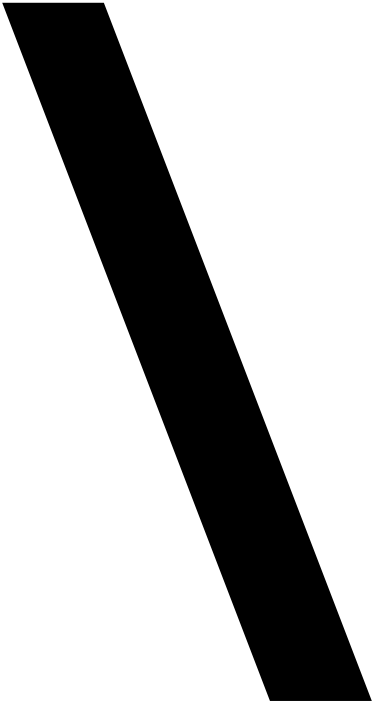
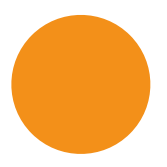
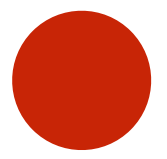
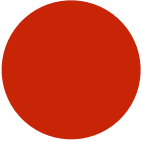
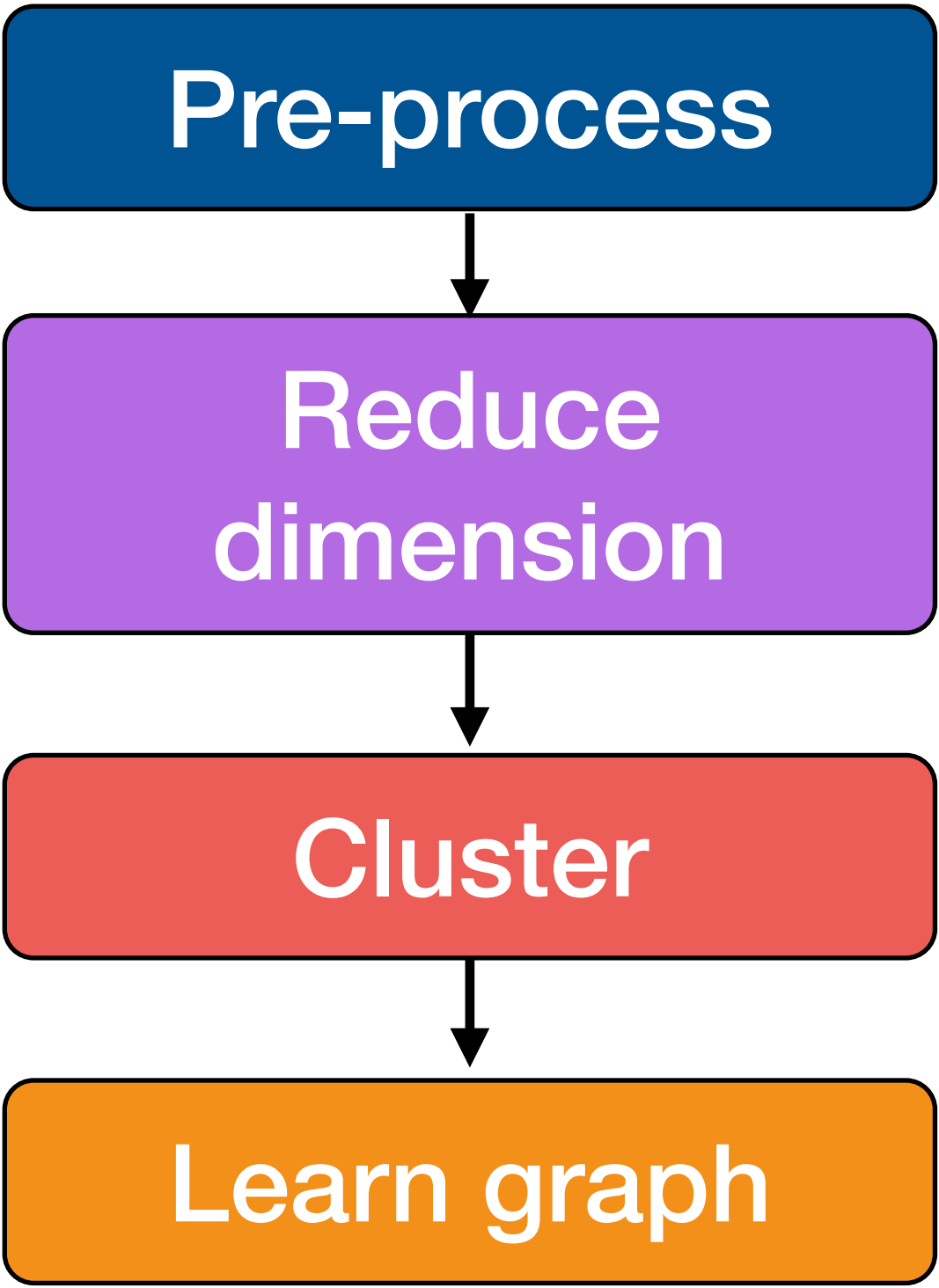


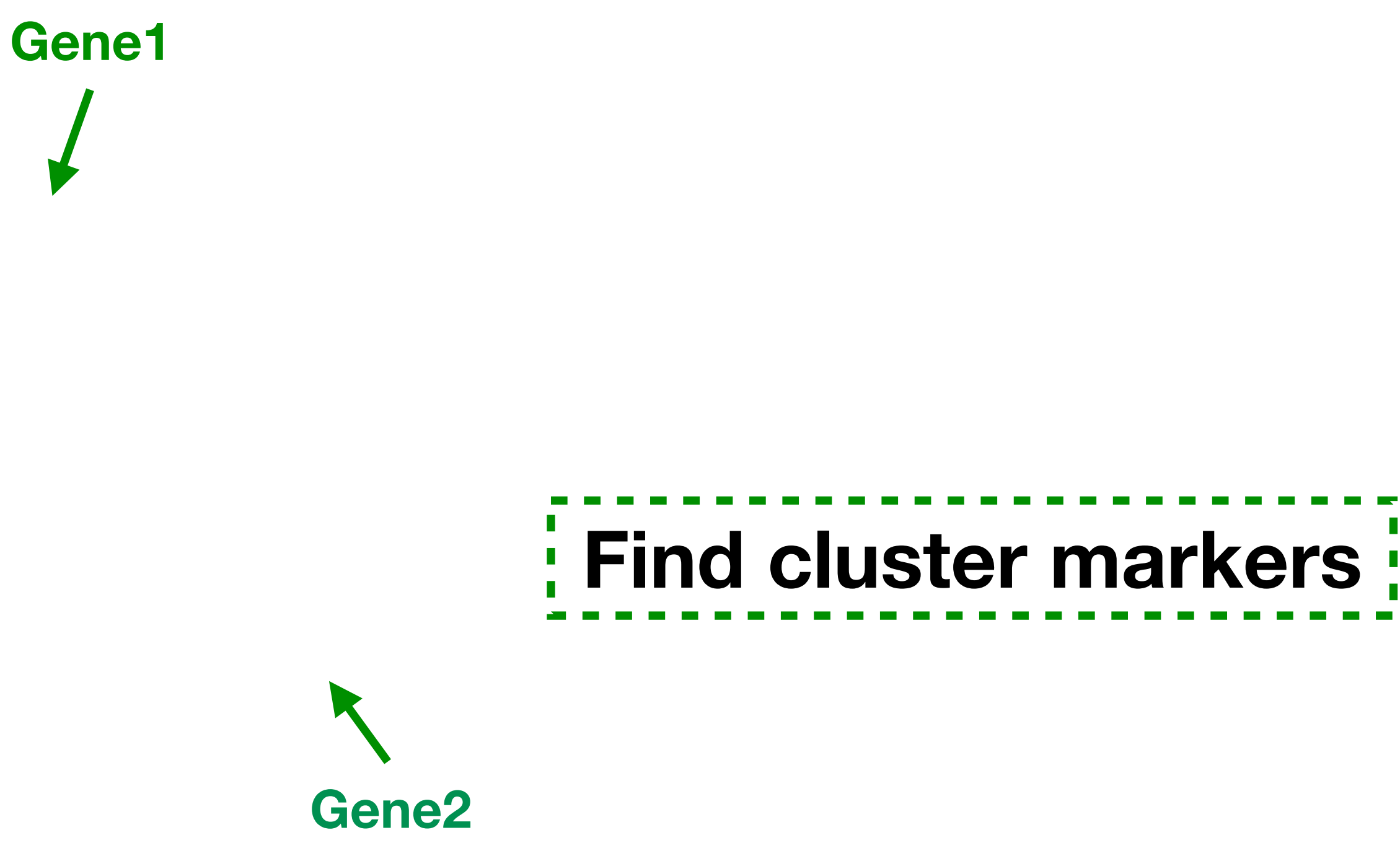







Pre-process
Reduce dimension
Cluster
Learn graph

Compare expression between clusters

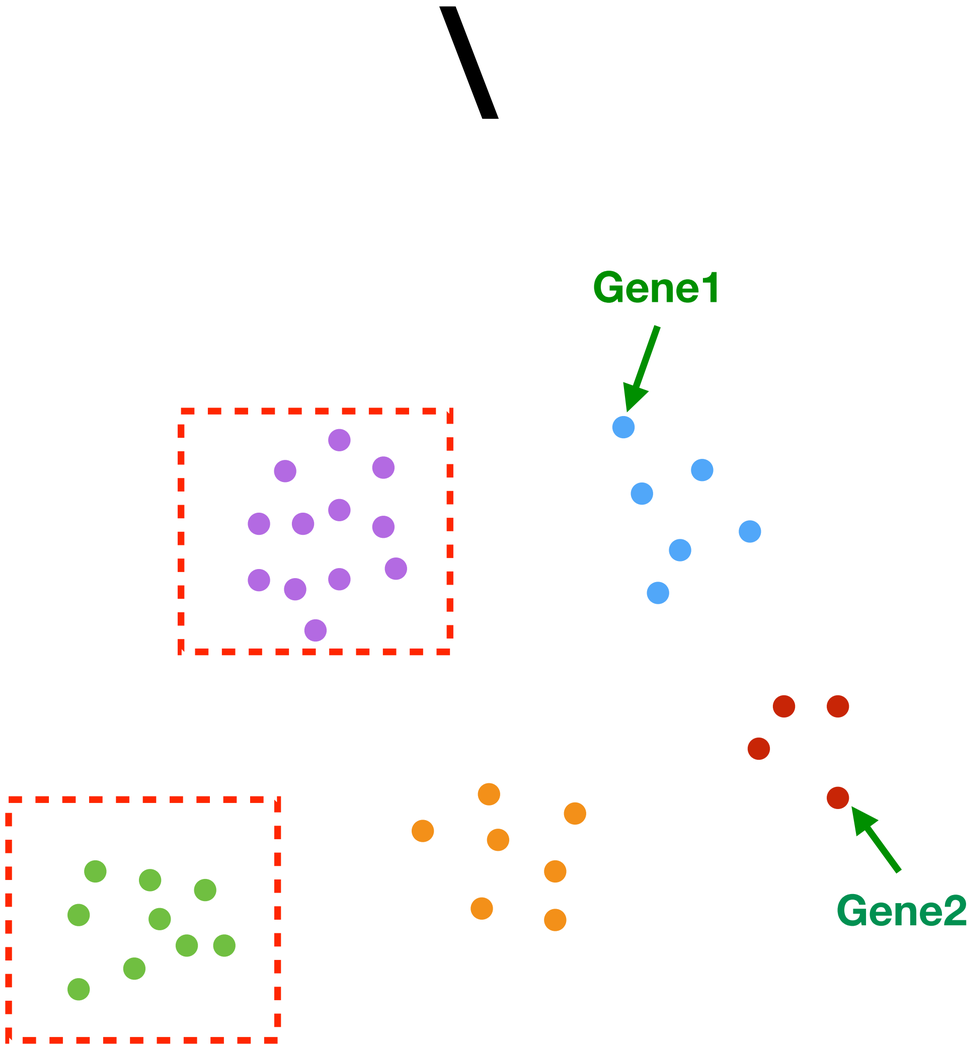

Gene1
Find cluster markers
Gene2

Pre-process

Reduce dimension

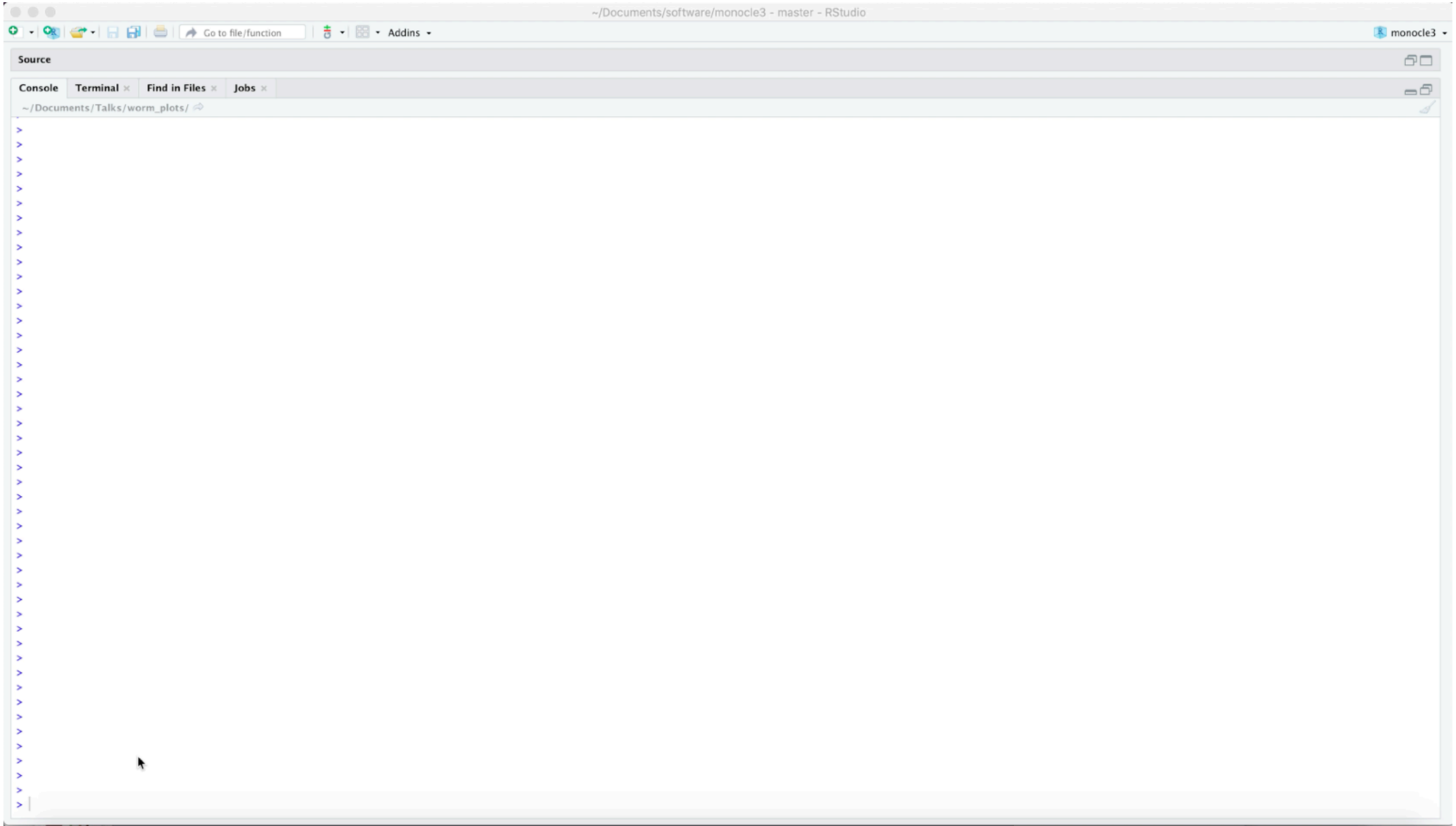
Cluster

Learn graph



Compare expression between clusters

Find cluster markers

Compare selected clusters

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cds_subset <- choose_cells(cds)
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