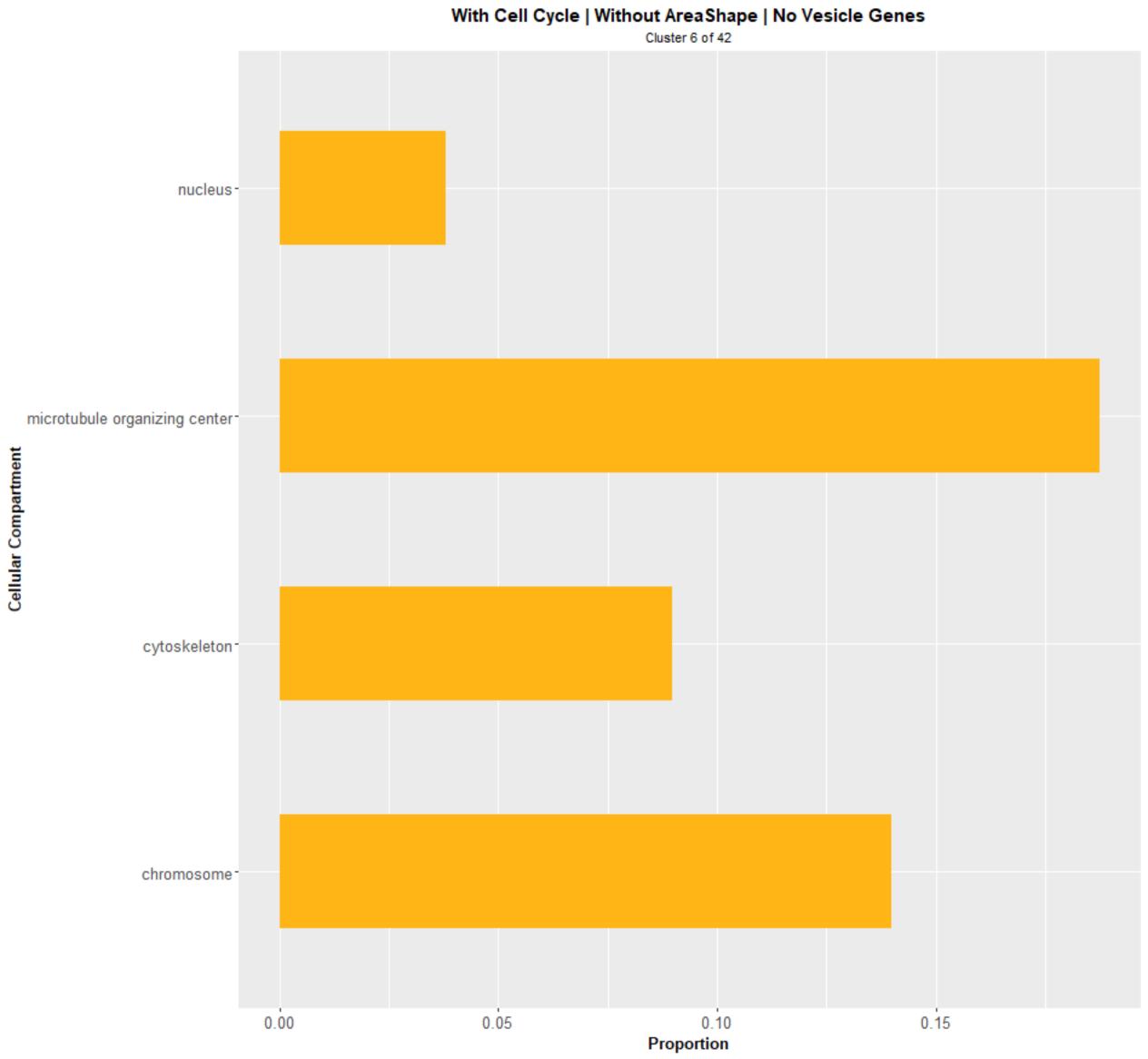
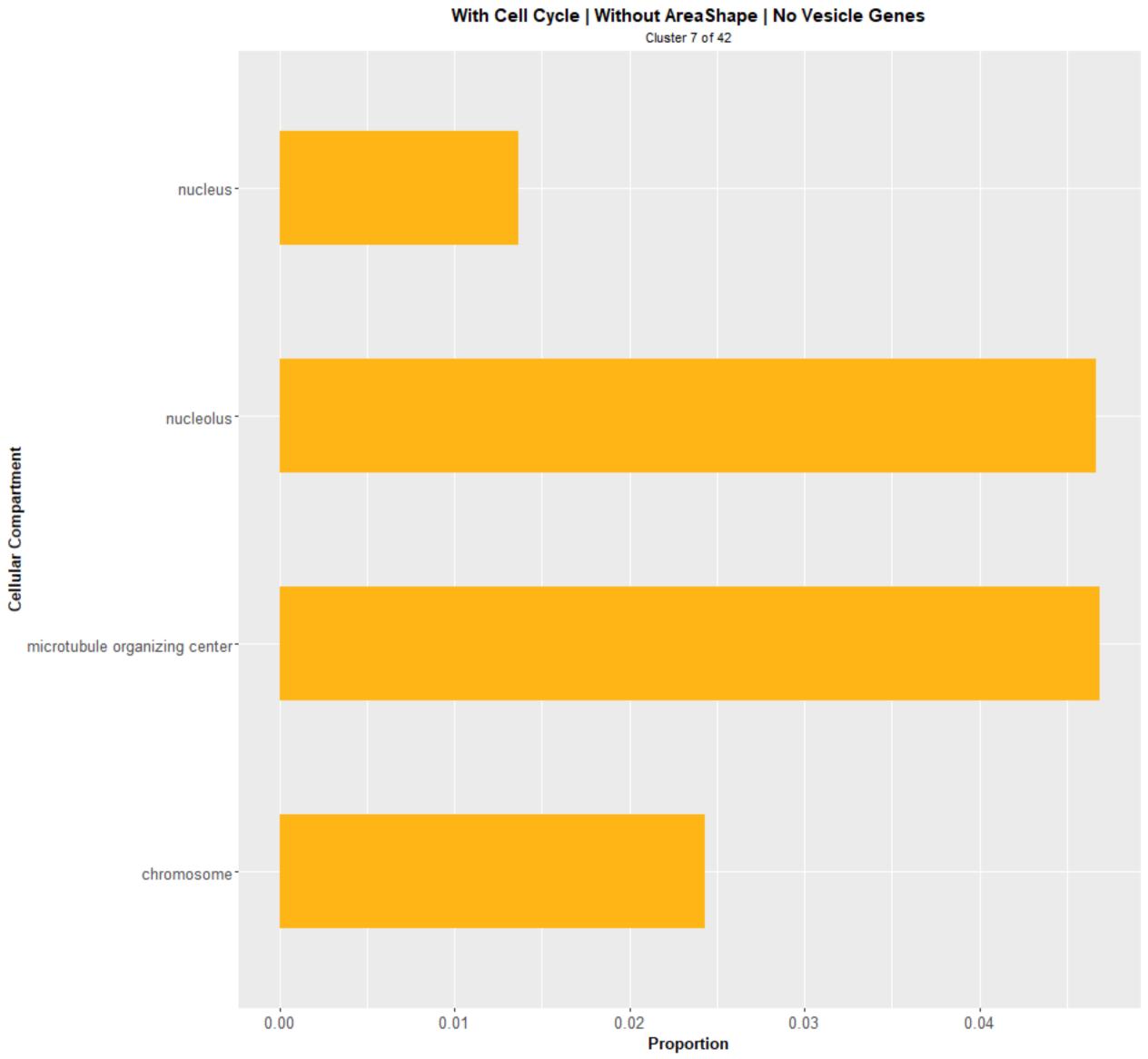
With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 1 of 42 nucleus-Cellular Compartment nucleolus-0.01 0.02 0.03 0.00 Proportion

With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 1 of 42 rRNA processing RNA catabolic process Biological Process ribosome assembly ribosomal large subunit_ biogenesis 0.00 0.02 0.04 0.08 Proportion

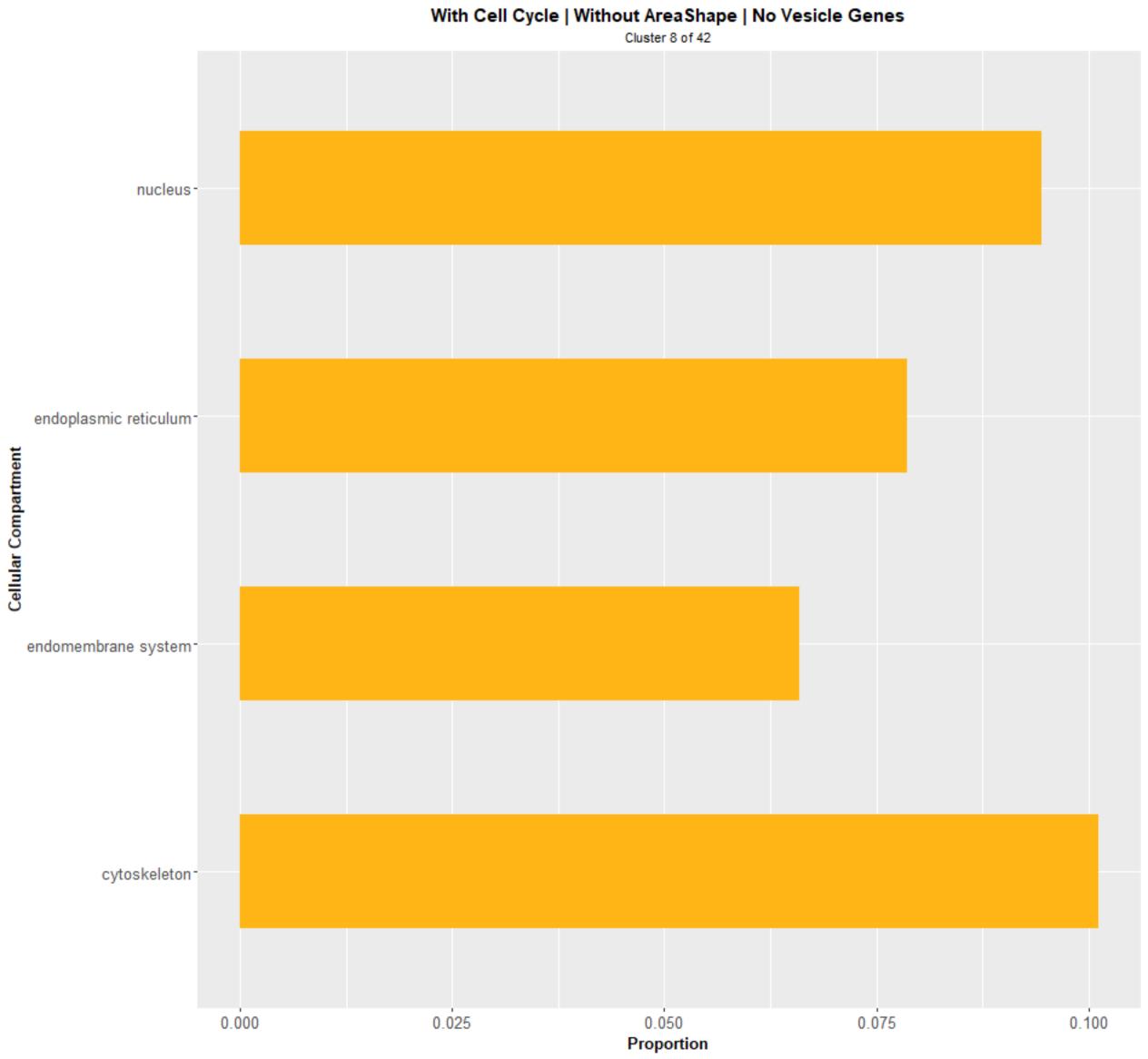


With Cell Cycle | Without AreaShape | No Vesicle Genes

Cluster 6 of 42 telomere organization regulation of organelle organization regulation of DNA metabolic_ regulation of cell cycle proteolysis involved in cellular protein catabolicprocess protein modification by small protein conjugation or removal protein complex biogenesis **Biological Process** organelle fission mitotic cell cycle meiotic cell cycle -DNA replication -DNA repair -DNA recombination cytoskeleton organization chromosome segregation chromatin organization cellular response to DNA_ damage stimulus 0.10 0.05 0.15 0.20 0.00 Proportion



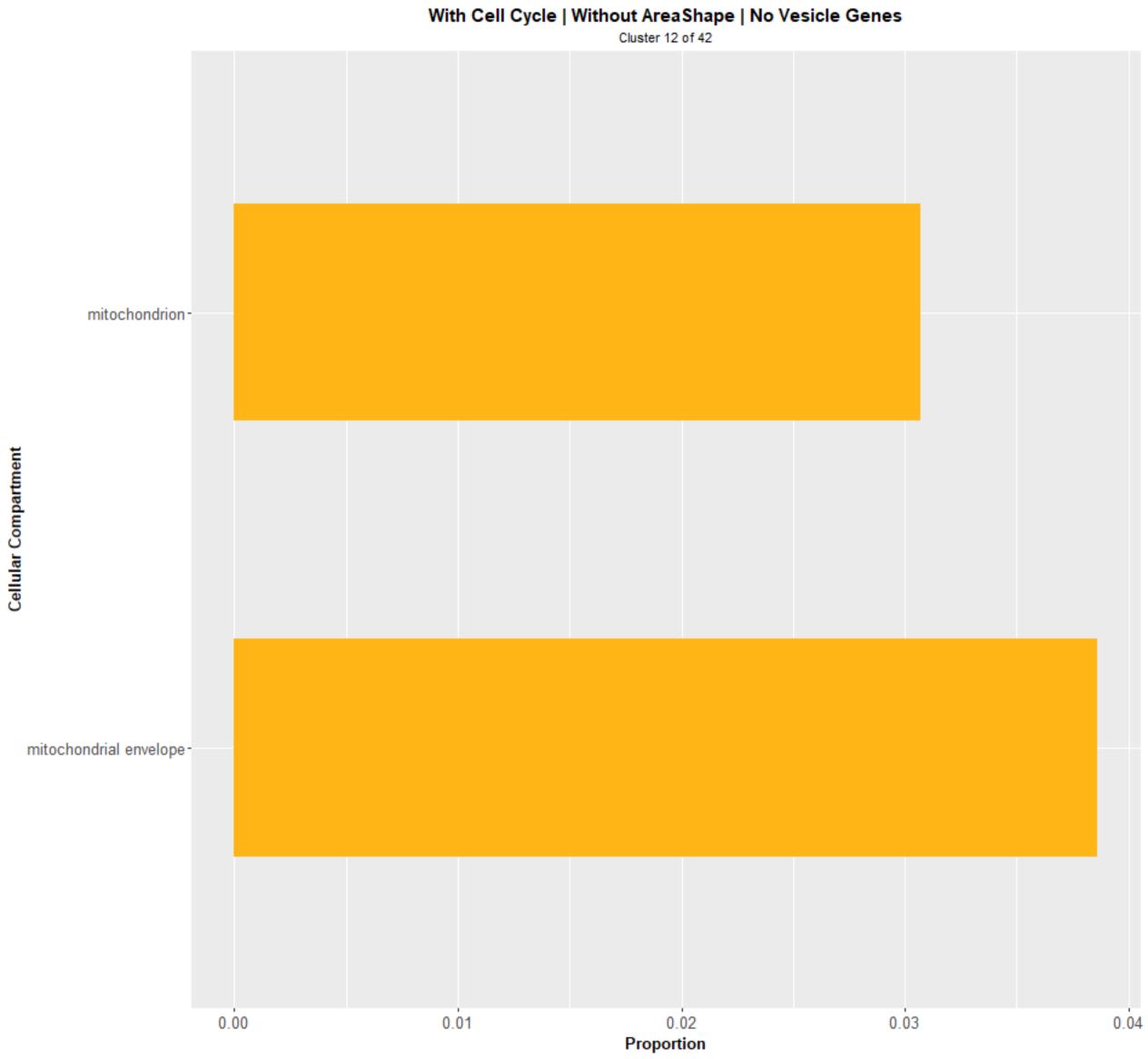
With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 7 of 42 transcription from RNA polymerase III promoter snoRNA processing rRNA processing ribosomal subunit export from nucleus **Biological Process** organelle fission nuclear transport mitotic cell cyclecytoskeleton organization chromosome segregation chromatin organization 0.000 0.025 0.050 0.075 0.100 0.125 Proportion



With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 8 of 42 translational initiation transcription from RNA polymerase II promoter snoRNA processing rRNA processing RNA splicing ribosomal subunit export from nucleus proteolysis involved in cellular protein catabolicprocess protein targeting protein lipidation protein complex biogenesis **Biological Process** protein acylation peptidyl-amino acid_ modification nucleobase-containing compound. nuclear transport mRNA processing lipid metabolic process DNA repair DNA-templated transcription, termination DNA-templated transcription, initiation DNA-templated transcription, _ elongation cytoskeleton organization cytokinesis cellular response to DNA damage stimulus 0.0 0.2 0.4 0.1 0.3 Proportion

With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 11 of 42 Cellular Compartment 0.000 0.005 0.015 0.020 0.010 Proportion

With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 11 of 42 rRNA processing ribosomal large subunit_ biogenesis Biological Process nuclear transport DNA-templated transcription, _ termination cytoskeleton organization 0.050 0.075 0.000 0.025 0.100 Proportion



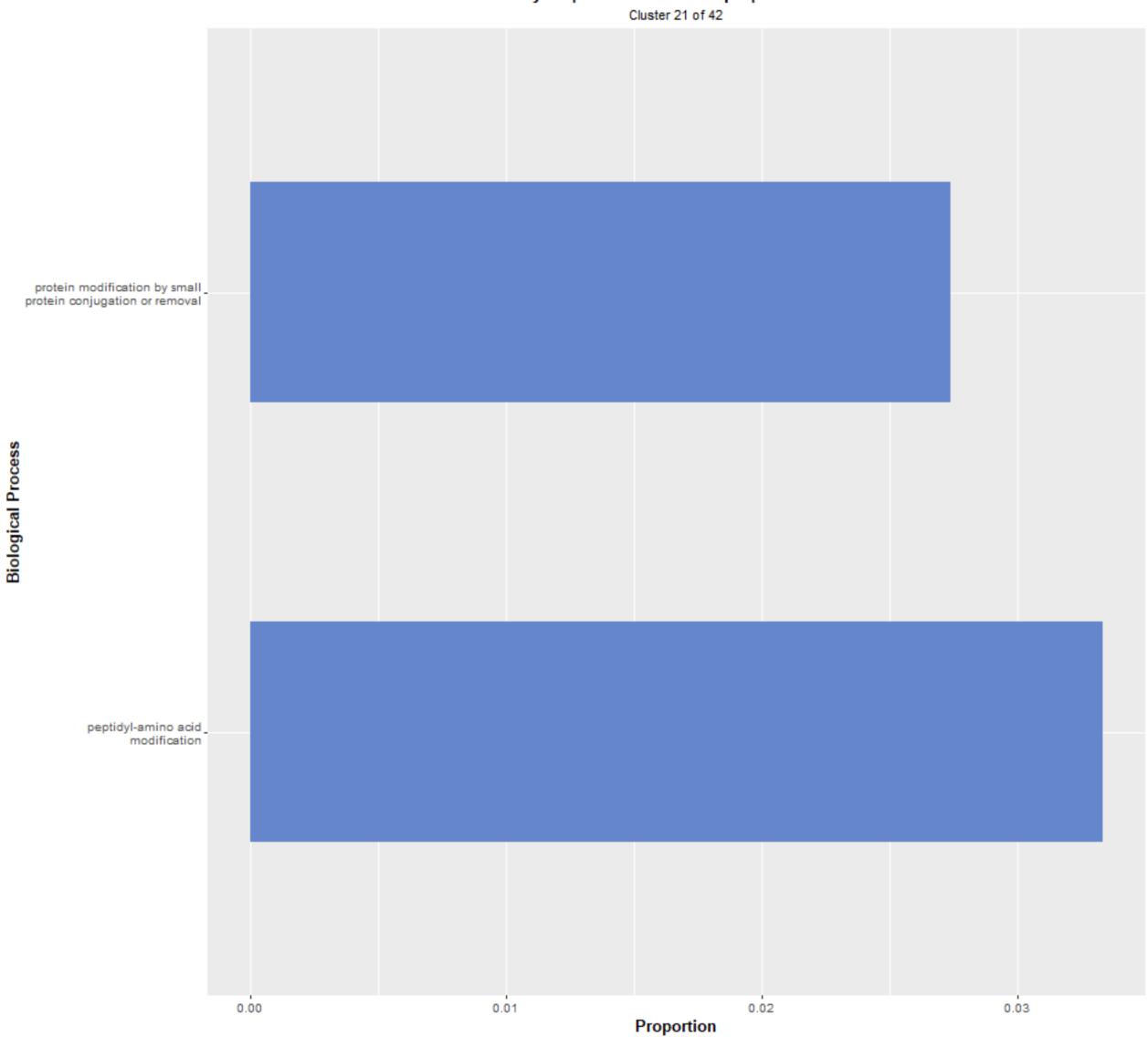
With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 12 of 42 nucleobase-containing small molecule metabolic process mitochondrion organization mitochondrial translation -**Biological Process** generation of precursor_ metabolites and energy cofactor metabolic process carbohydrate metabolic process 0.050 0.000 0.025 0.075 Proportion

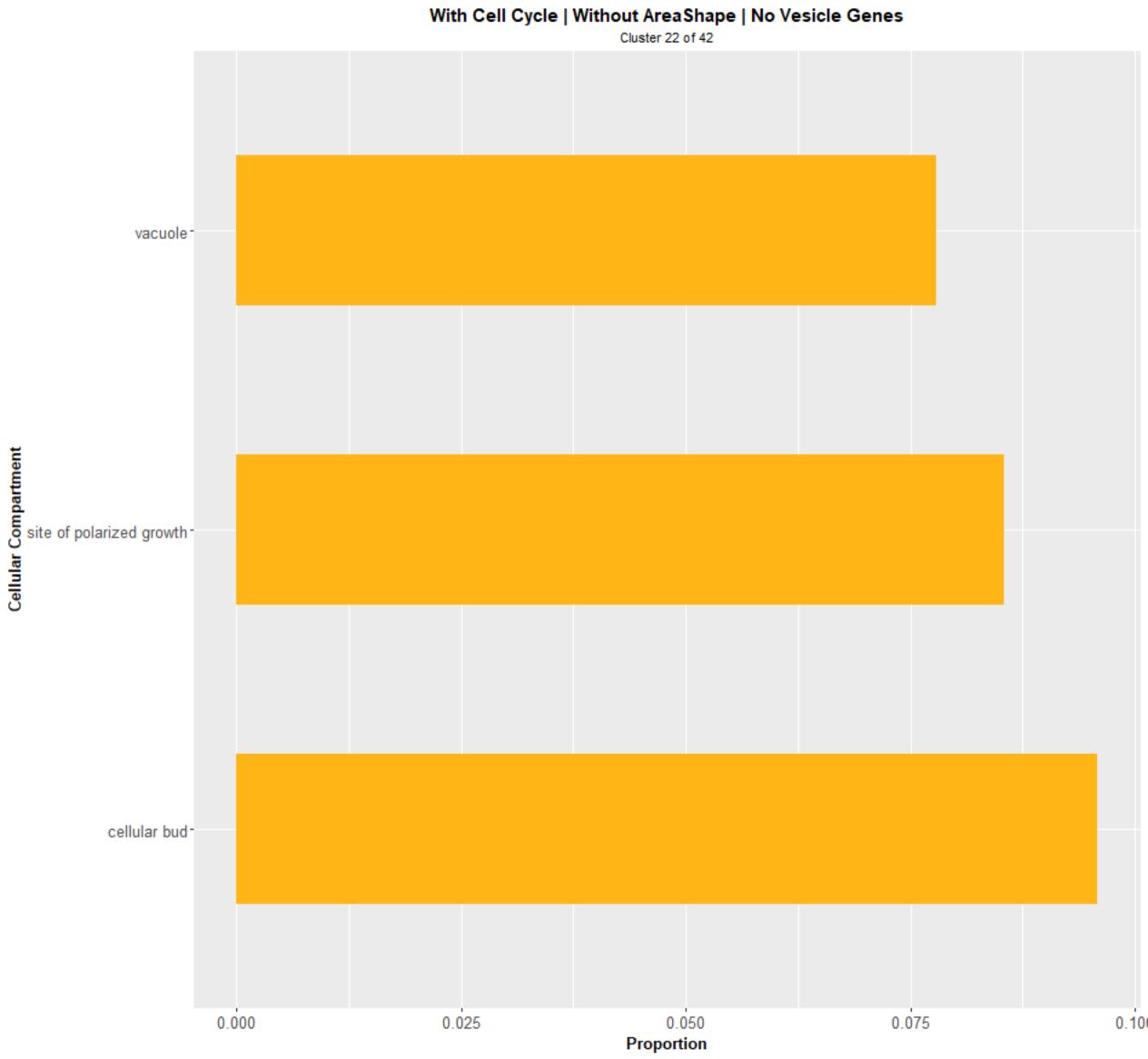
With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 14 of 42 Cellular Compartment 0.000 0.005 0.010 0.015 Proportion

With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 14 of 42 tRNA aminoacylation for protein translation transcription from RNA polymerase II promoter RNA splicing Biological Process mRNA processing -DNA-templated transcription, initiation chromosome segregation -0.06 0.00 0.03 0.09 0.12 Proportion

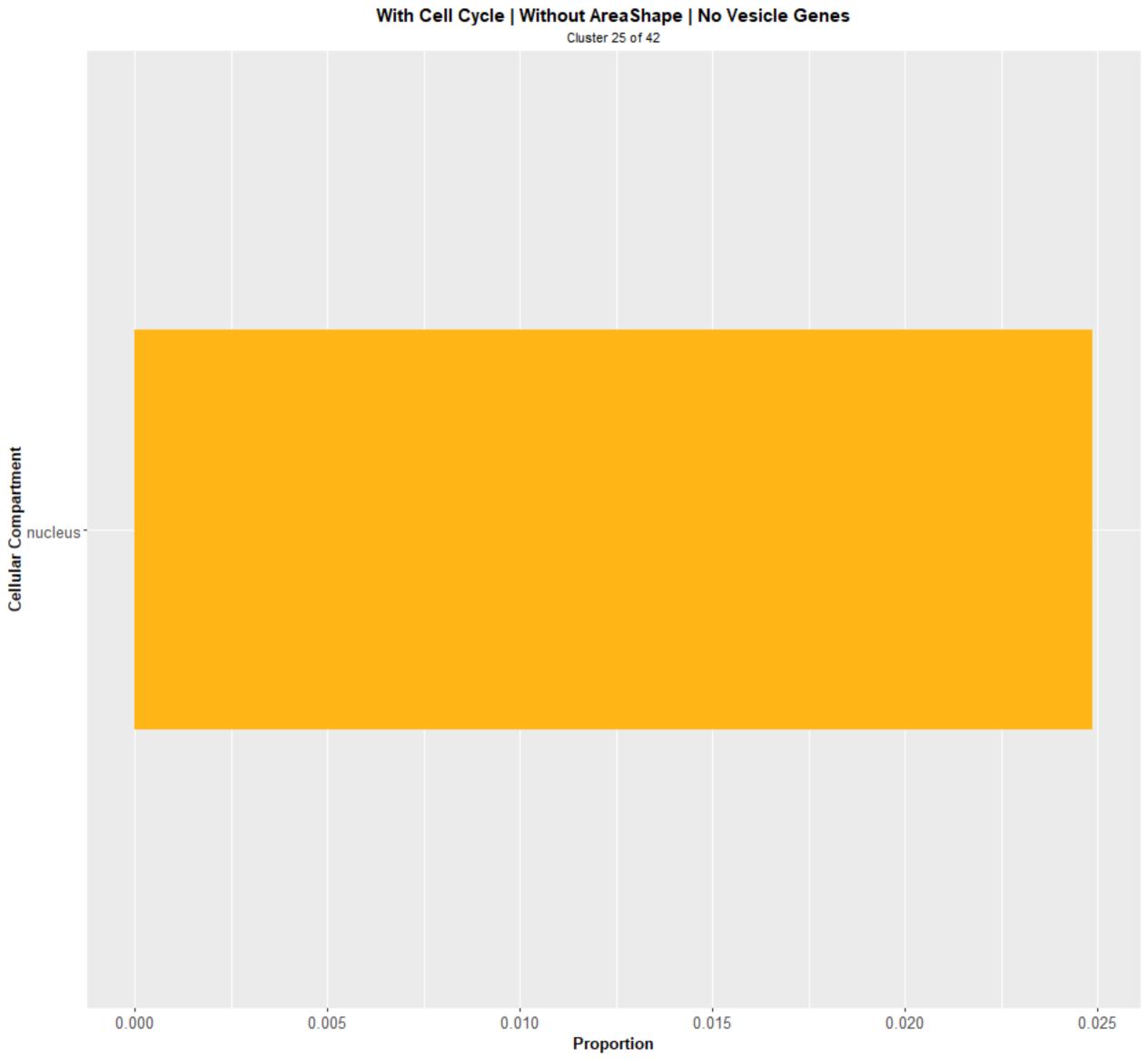
With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 16 of 42 Biological Process Lessonse to starvation -0.02 0.05 0.00 0.01 0.03 0.04 Proportion

With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 19 of 42 Cellular Compartment 0.00 0.01 0.02 0.03 0.04 Proportion



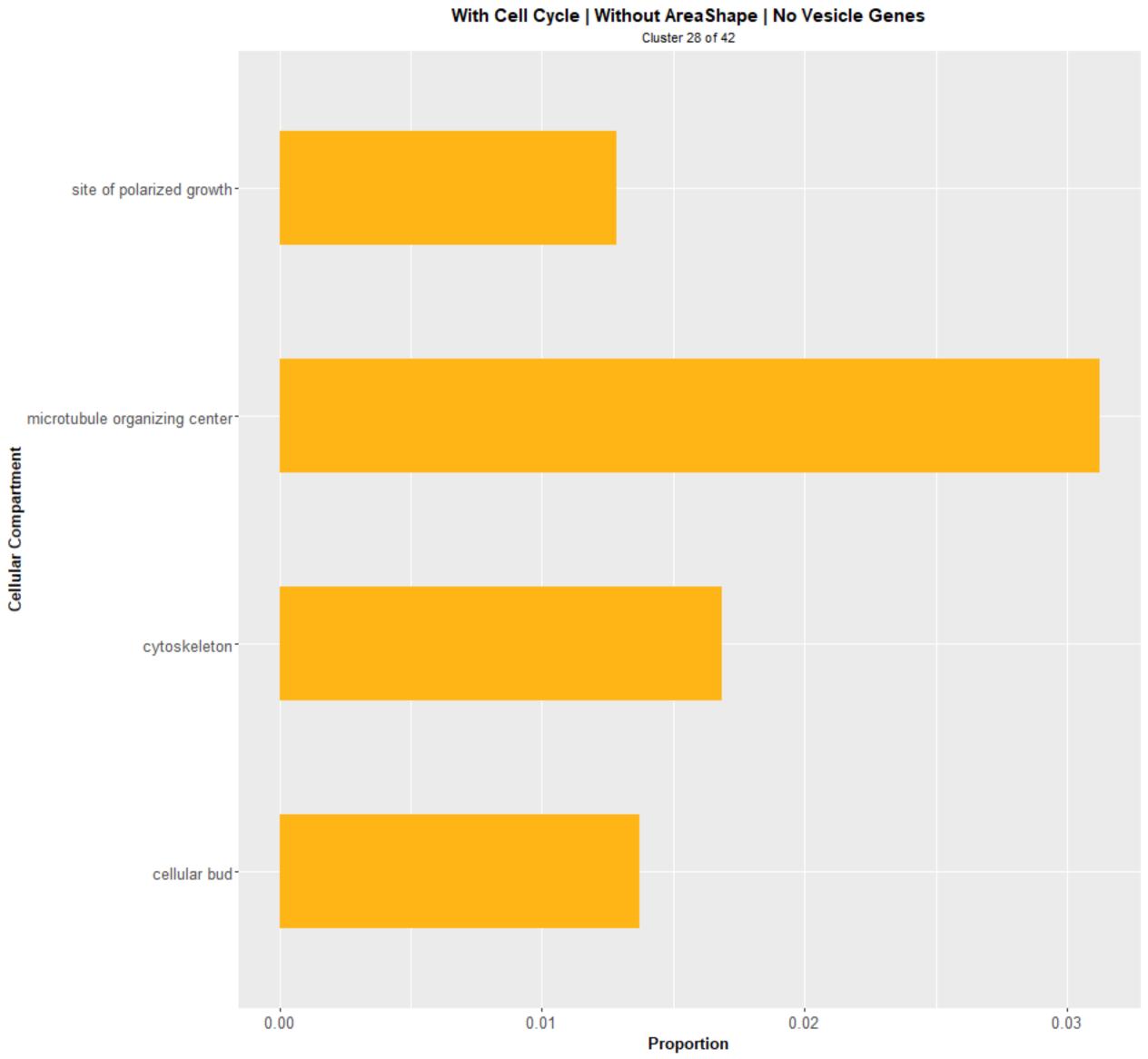


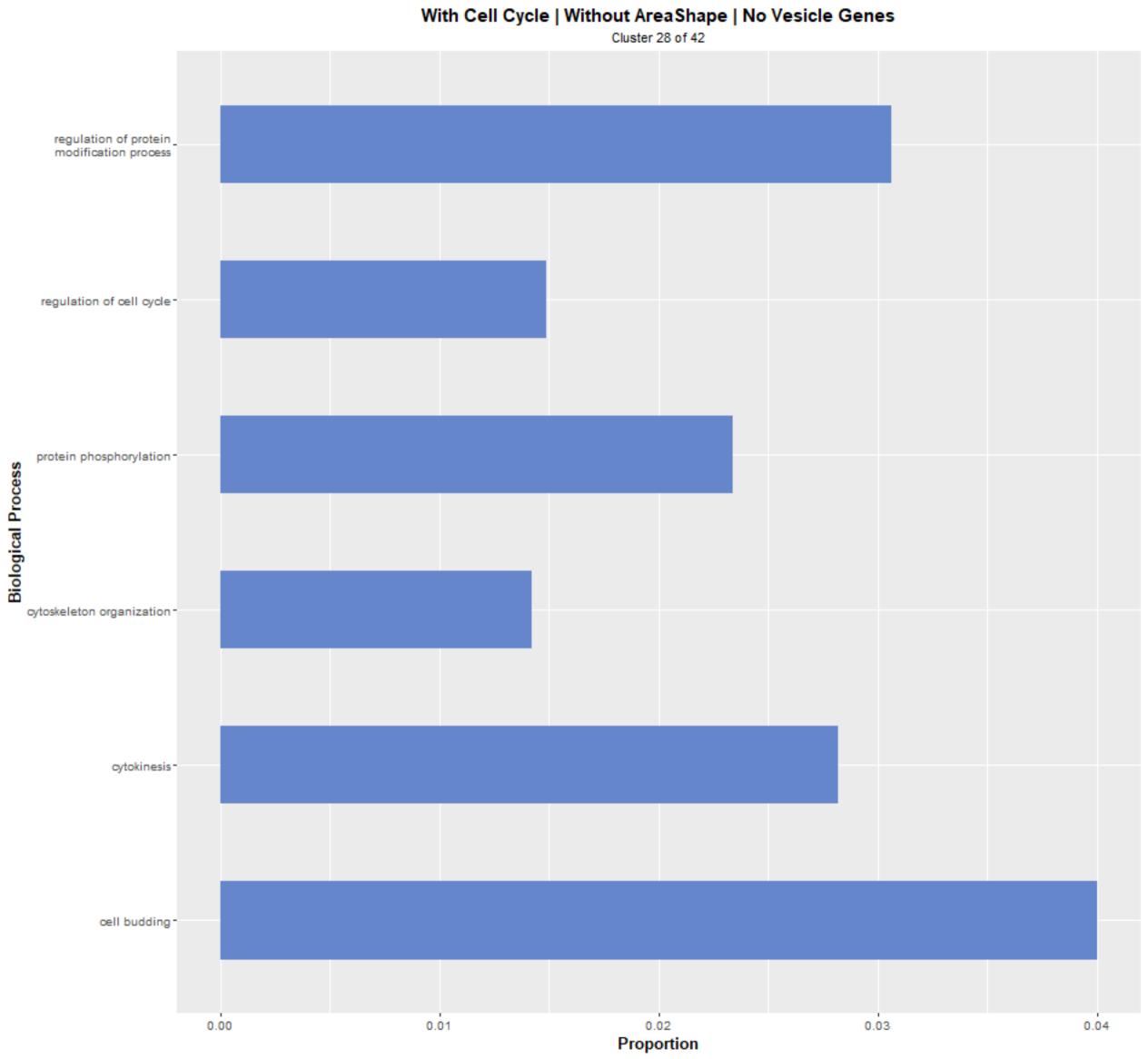
With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 22 of 42 regulation of transport Biological Process chromatin organization -0.00 0.04 0.08 0.12 Proportion



With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 25 of 42 ribosomal small subunit biogenesis Biological Process protein maturation -0.050 0.000 0.025 0.075 Proportion

With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 26 of 42 organelle fission -Biological Process cytoplasmic translation -0.04 0.08 0.02 0.00 0.06 Proportion





With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 31 of 42 translational elongation transcription from RNA polymerase II promoter transcription from RNA polymerase I promoter Biological Process response to heatprotein acylation endosomal transport cytoplasmic translation chromatin organization -0.10 0.00 0.05 0.15 Proportion

0.02

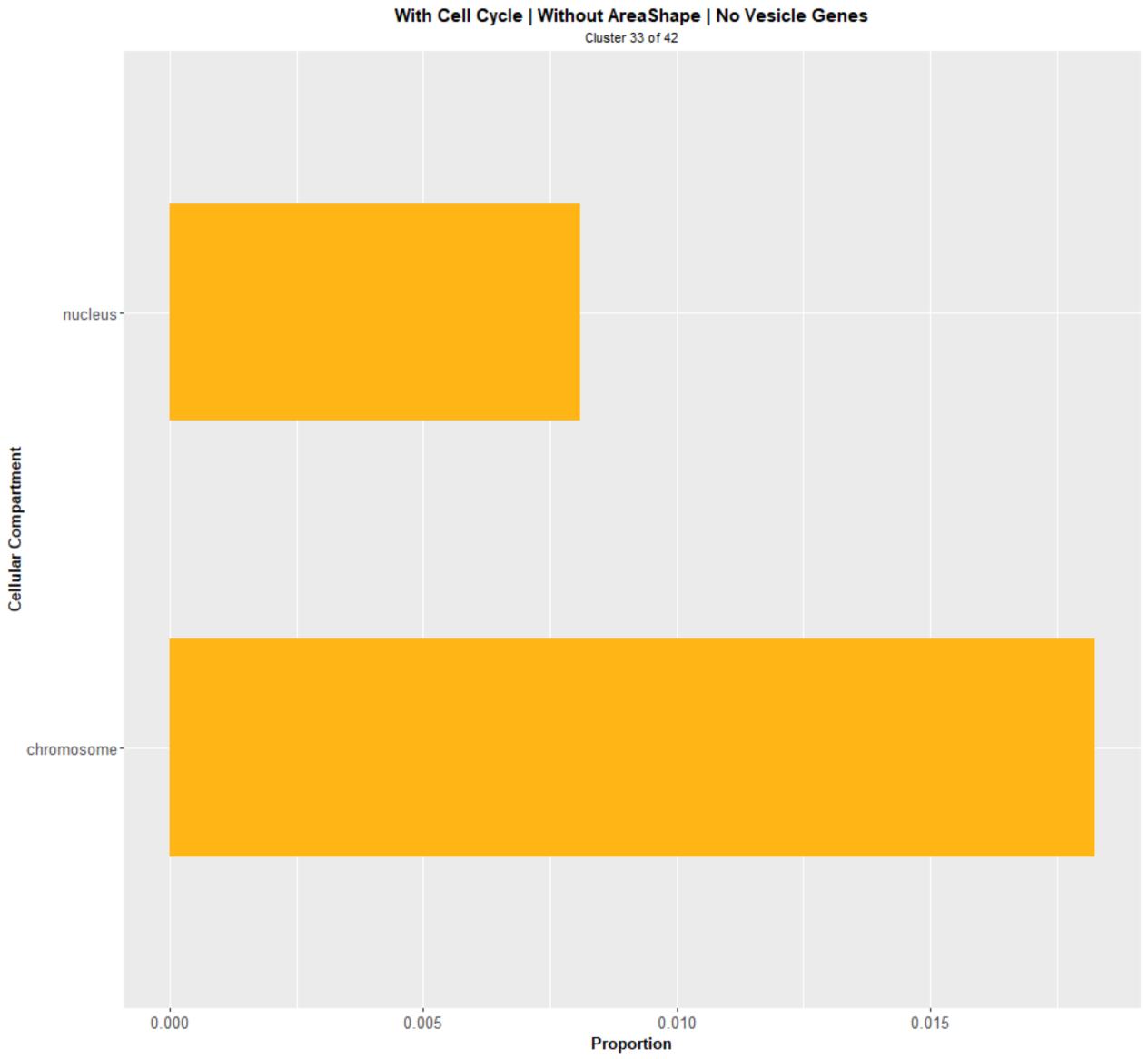
Proportion

0.03

0.04

0.01

0.00



With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 33 of 42 transposition telomere organization -RNA catabolic process regulation of DNA metabolic_ process organelle fission -Biological Process mitotic cell cycle -DNA replication -DNA repair DNA recombination chromosome segregation cellular response to DNA damage stimulus 0.00 0.05 0.10 0.15 Proportion

With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 40 of 42 Biological Process cellular amino acid metabolic_ process 0.02 Proportion 0.00 0.01 0.03 0.04

With Cell Cycle | Without AreaShape | No Vesicle Genes Cluster 41 of 42 Cellular Compartment 0.01 0.02 0.00 0.03 Proportion