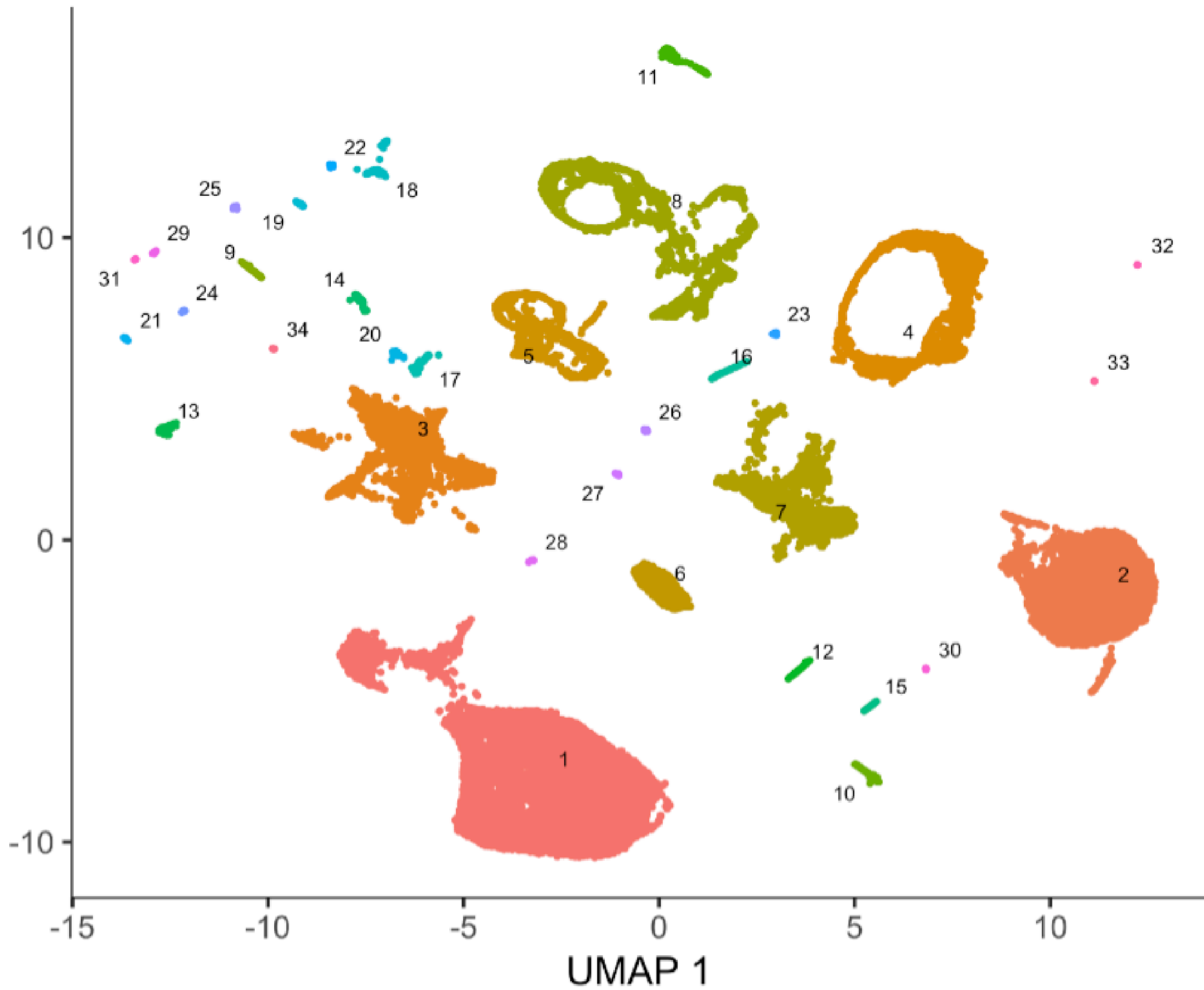
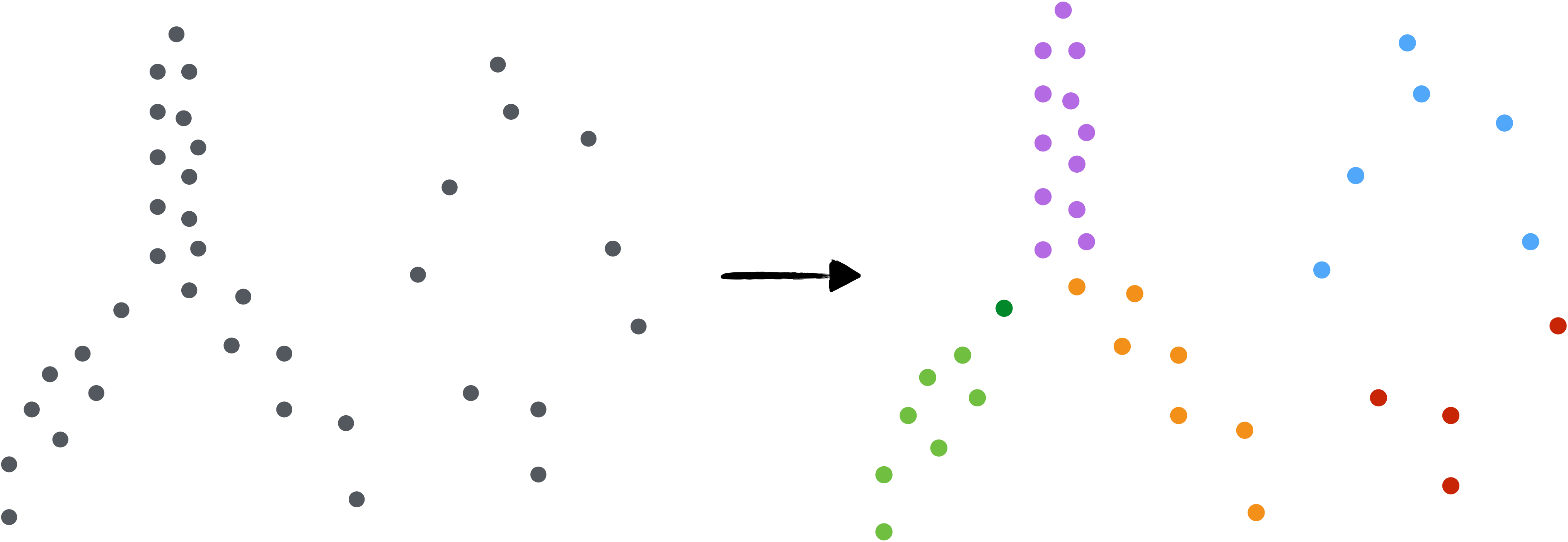




UMAP 2



Group cells by partitions



```
cds <- cluster_cells(cds)
plot_cells(cds, color_cells_by="partition",
group_cells_by="partition")
```

**Pre-process**



```
graph TD; A[Pre-process] --> B[Reduce dimension]; B --> C[Cluster];
```

**Reduce  
dimension**

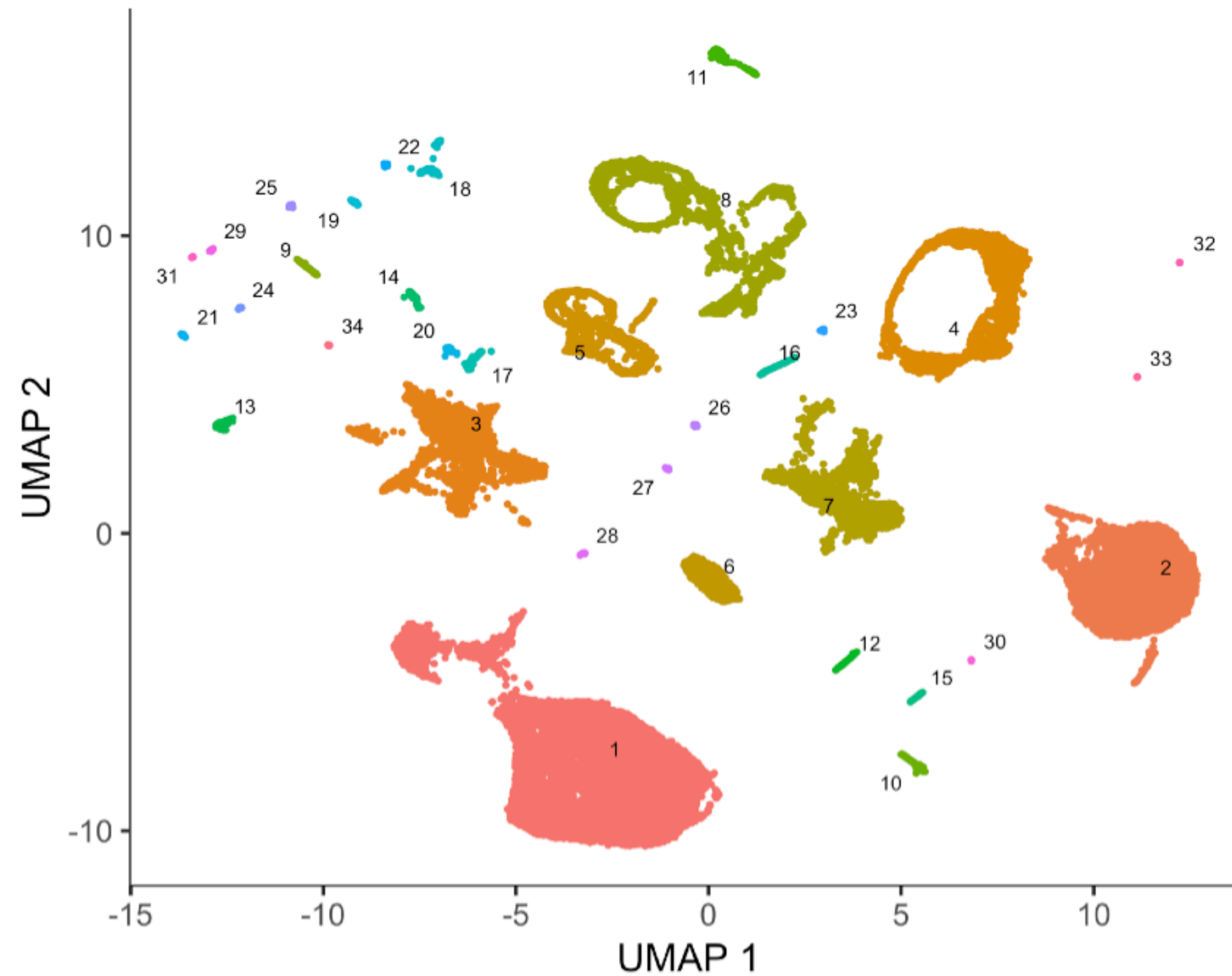
**Cluster**

Pre-process

Reduce  
dimension

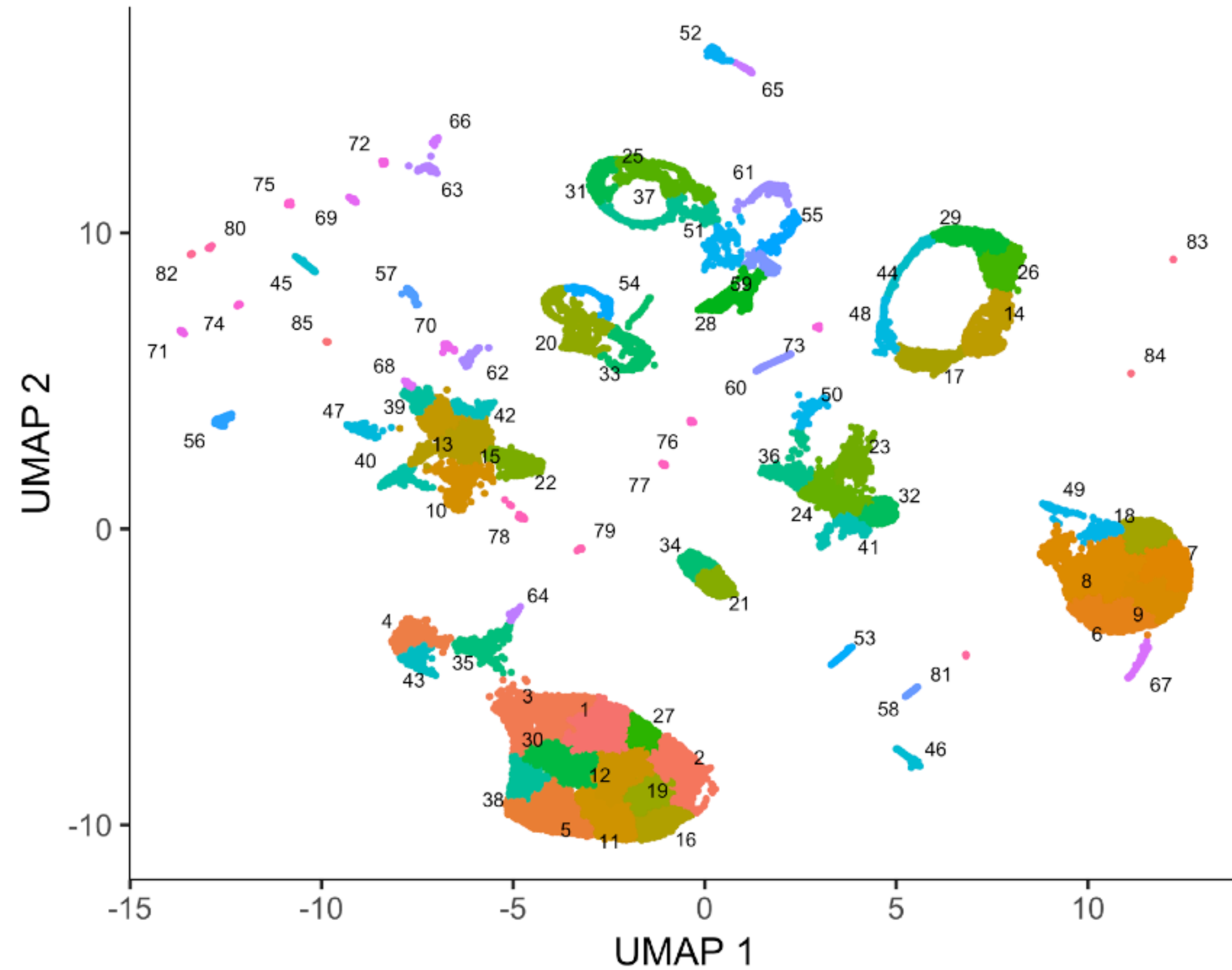
Cluster

# Group cells by partitions



```
cds <- cluster_cells(cds)
plot_cells(cds, color_cells_by="partition",
group_cells_by="partition")
```

# Group cells by clusters



```
cds <- cluster_cells(cds)
plot_cells(cds)
```