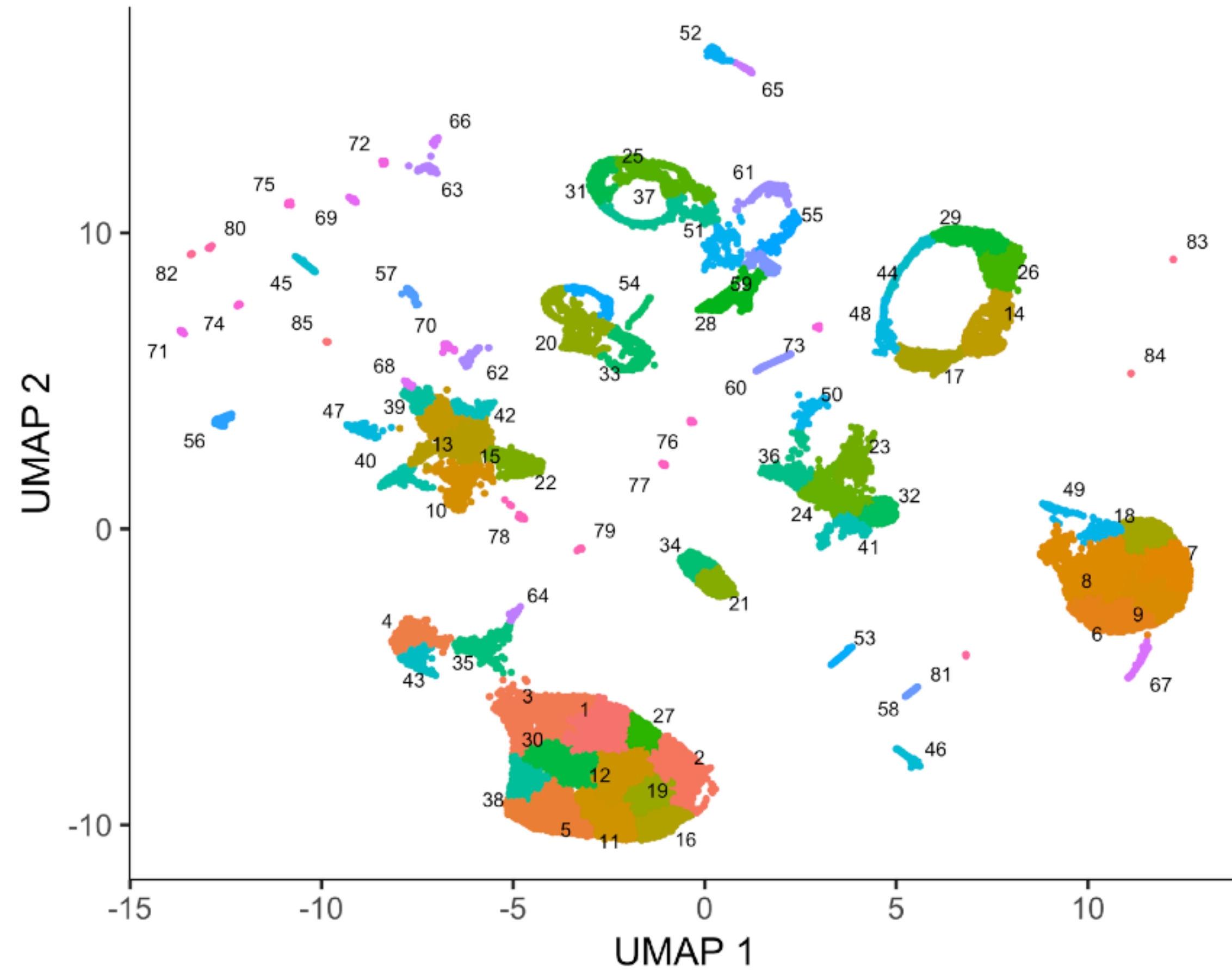


Group cells by clusters



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

Compare clusters

