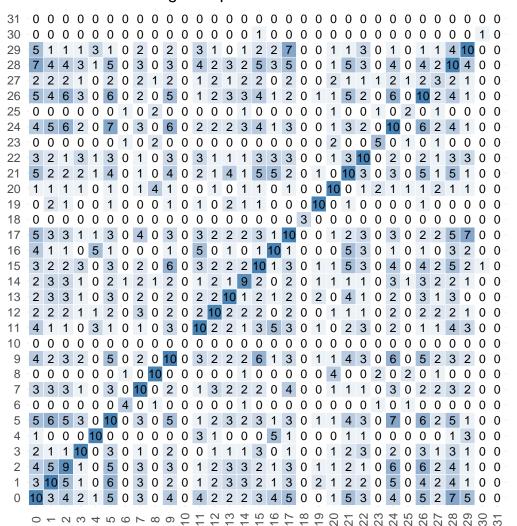
## Number of similar genes per cluster and without confounders



res

10.0 7.5

5.0

2.5

0.0