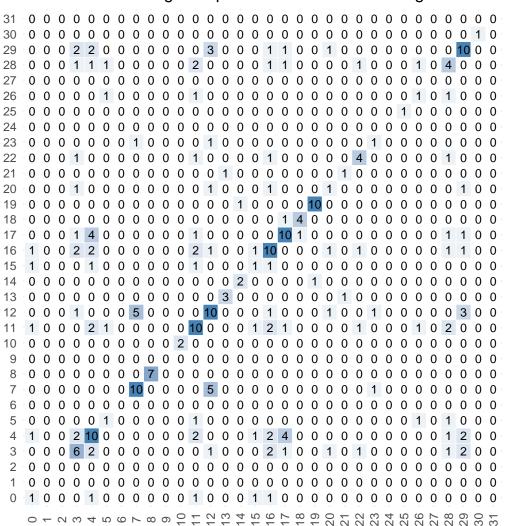
Number of similar genes per cluster and considering confounders



res 10.0 7.5 5.0 2.5 0.0