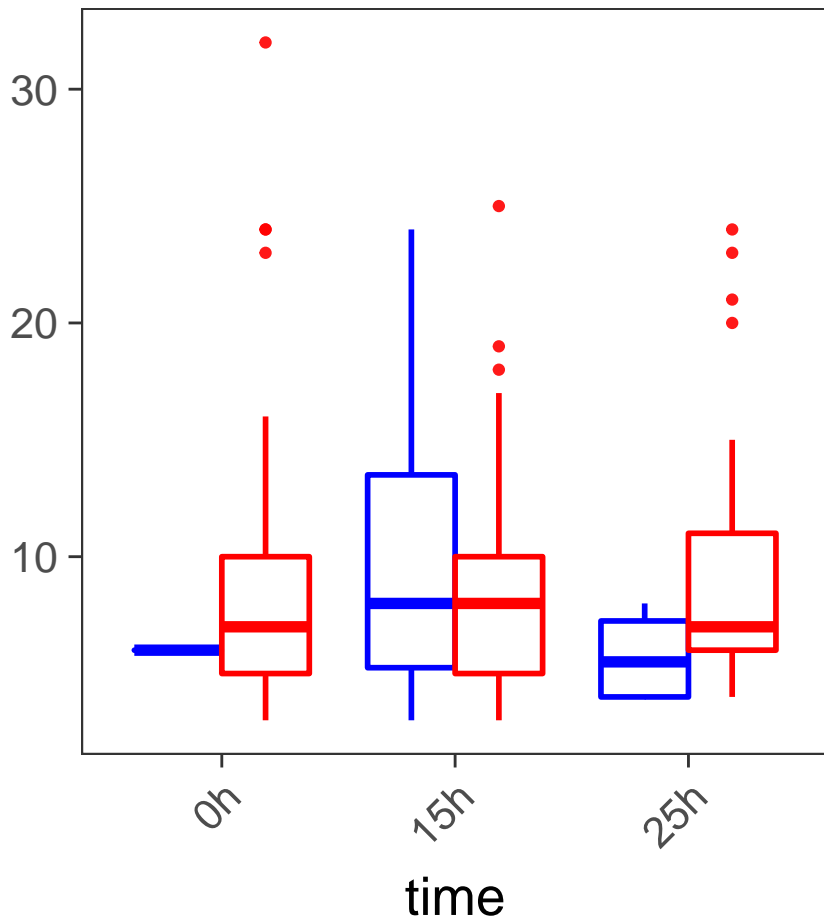


Number of genes per cluster



group

downregulation

upregulation