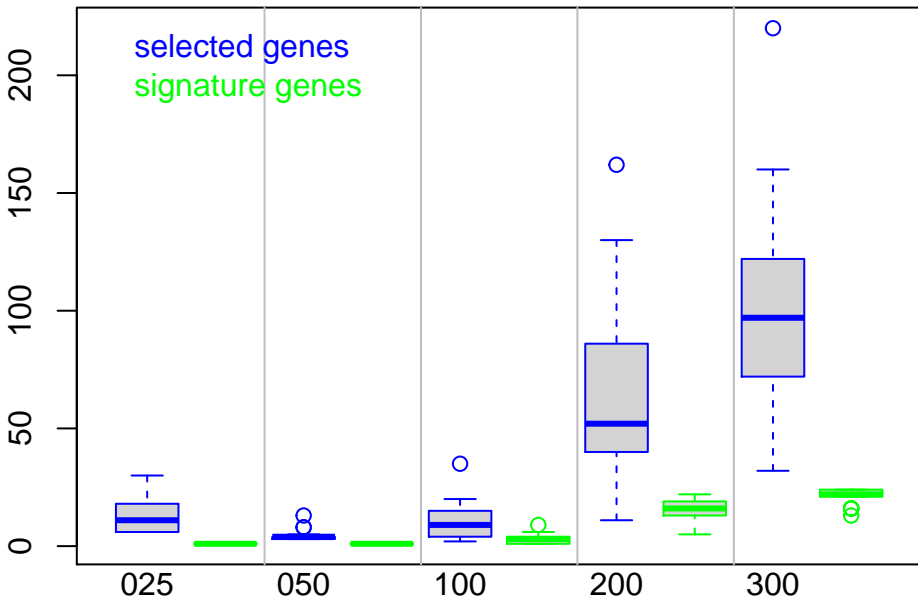**Sim = Sim_01**

one se lambda – selected gene counts   **min lambda – selected gene counts**

selected genes
signature genes