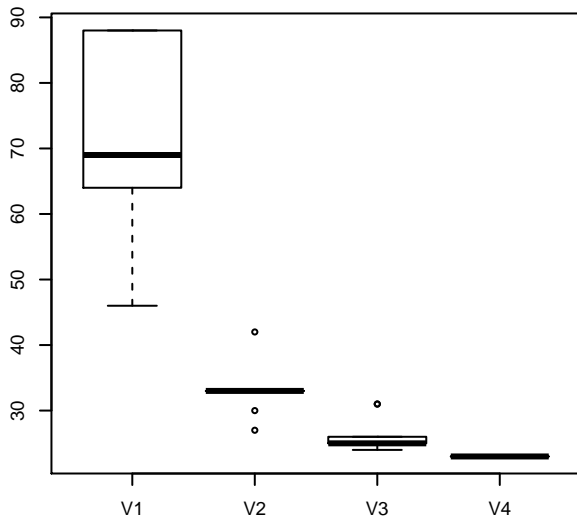
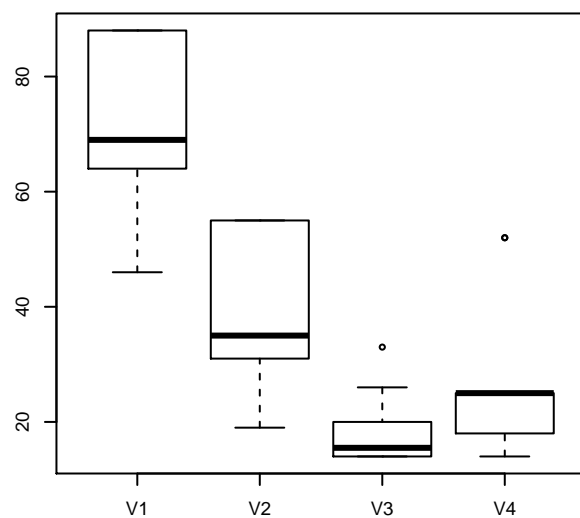


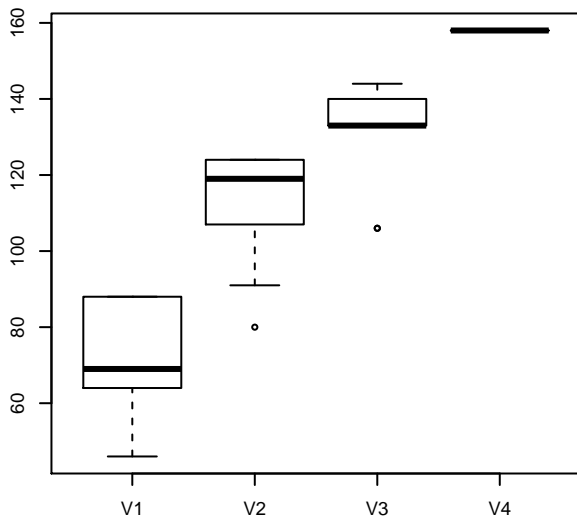
**number\_of\_conserved\_genes.Rtab**



**number\_of\_new\_genes.Rtab**



**number\_of\_genes\_in\_pan\_genome.Rtab**



**number\_of\_unique\_genes.Rtab**

