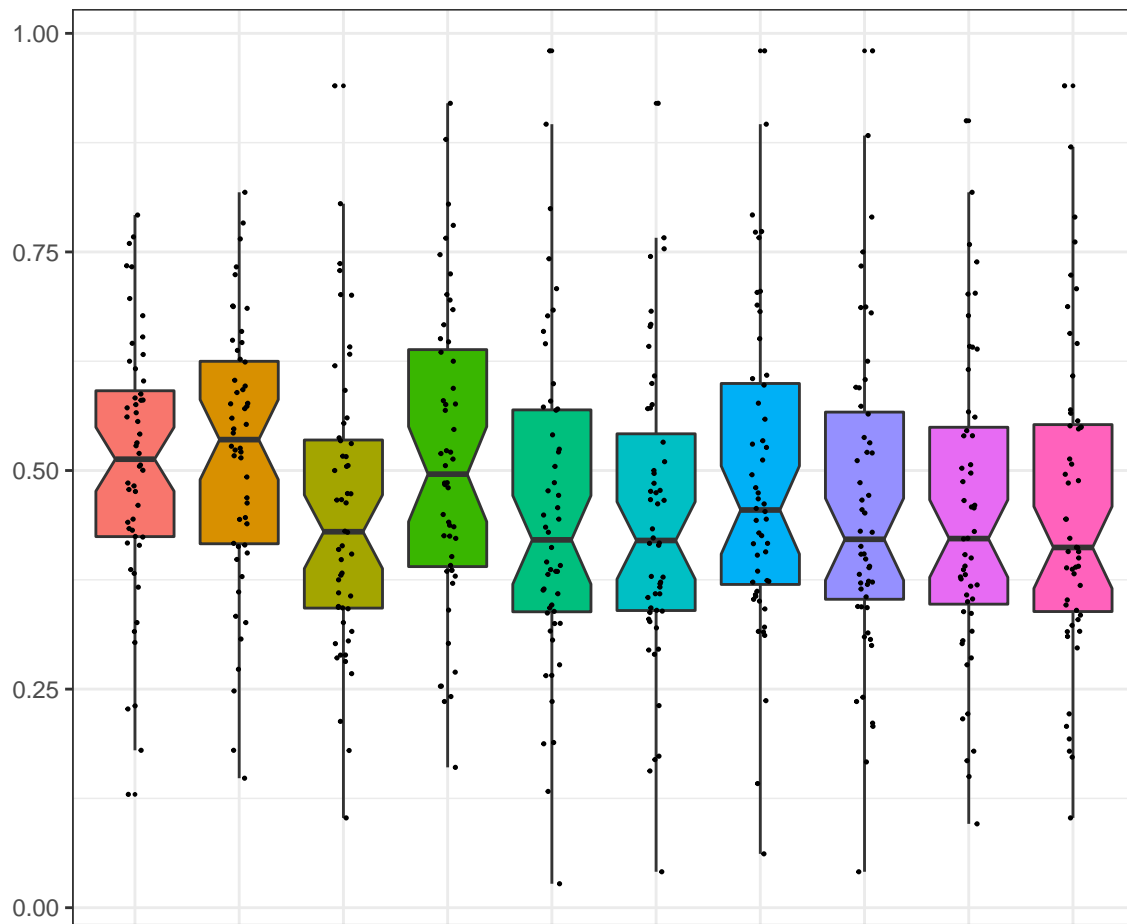


dnabackbone  
#(domains)=52



factor(variable)

- instances\_individuals\_change\_ratio
- aa\_ref\_overlap\_individuals\_change\_ratio
- jsd\_100way\_instances\_major\_ratio
- jsd\_100way\_aa\_not\_used\_ratio
- jsd\_mul\_aa\_ref\_SE
- jsd\_SE\_diff\_ratio
- SE\_jsd\_diff\_ratio
- jsd\_SE\_sum
- jsds\_ratio
- jsds\_subtraction