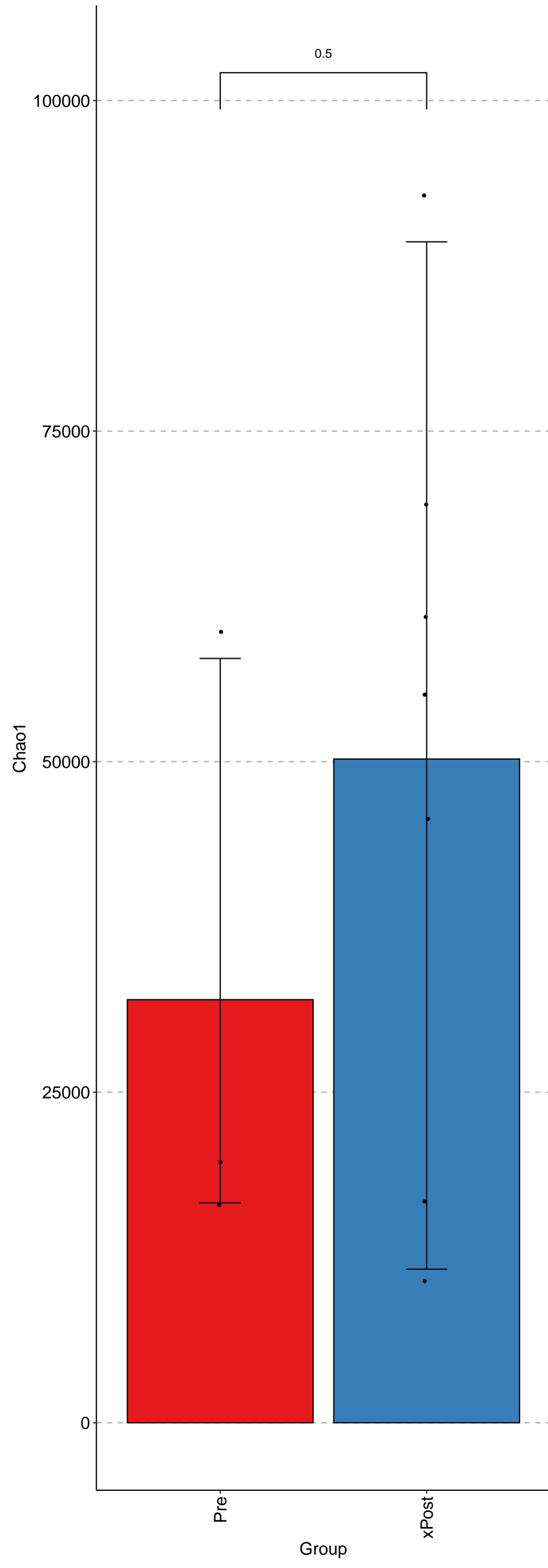
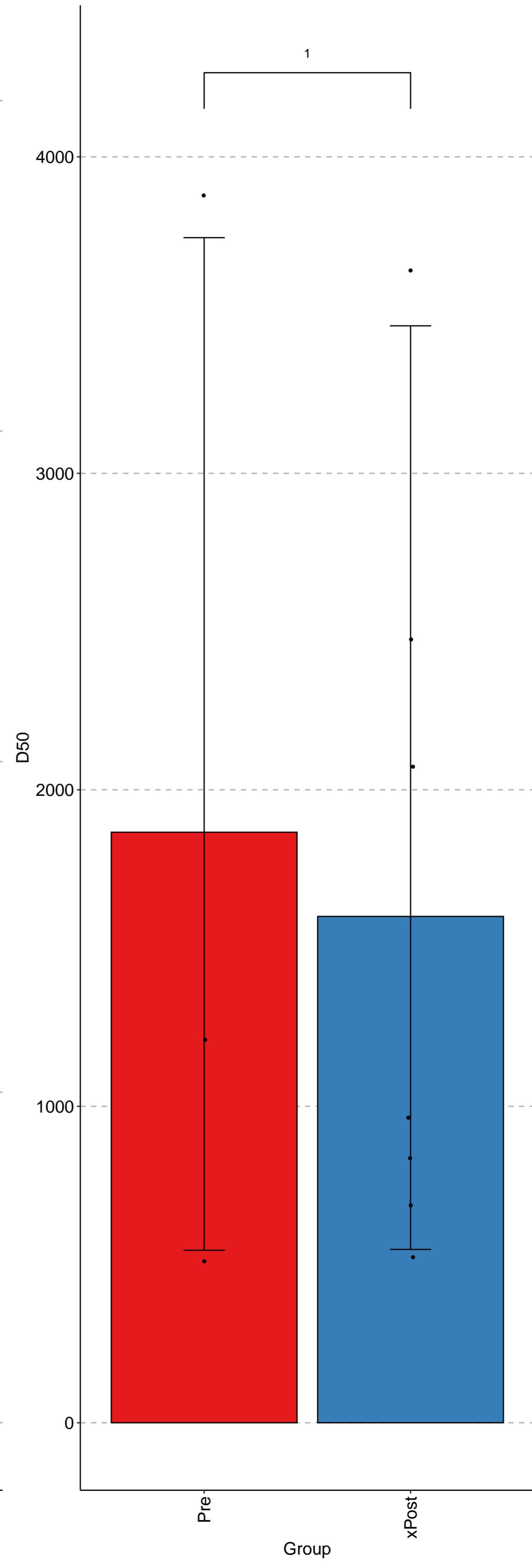


**Chao1**  
Sample diversity estimation using Chao1



**D50 diversity index**  
Number of clonotypes occupying the 50% of repertoires



**Inverse Simpson index**  
Sample diversity estimation using the inverse Simpson index

