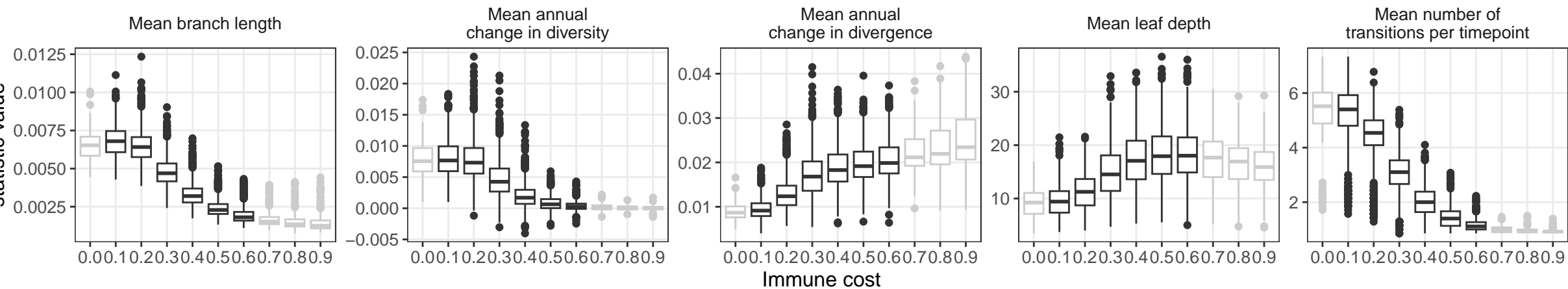
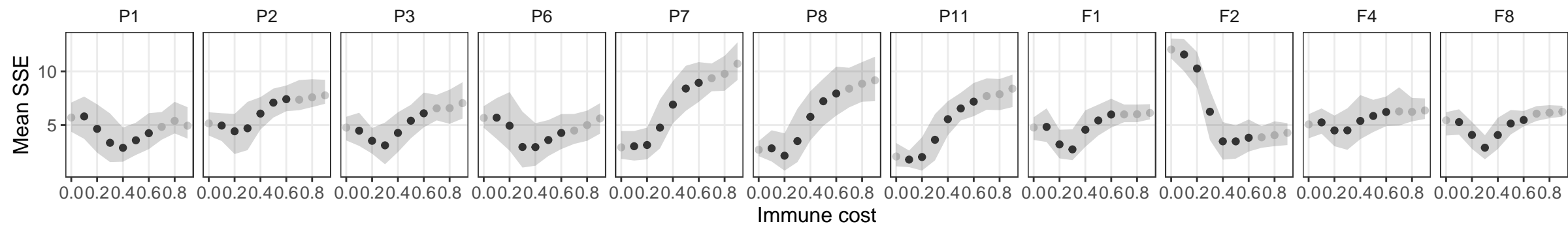


**A**Phylogenetic summary  
statistic value**B**

Number of replicates • 20 • 100