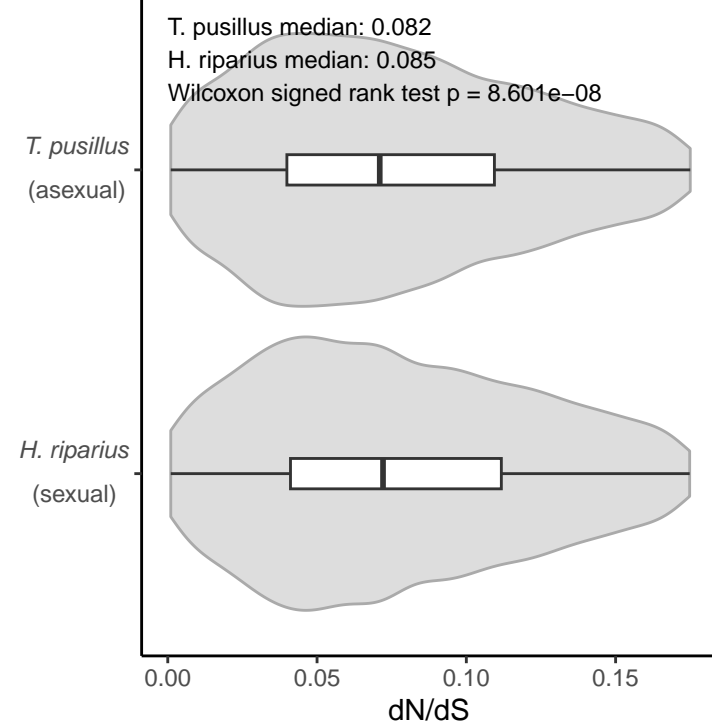
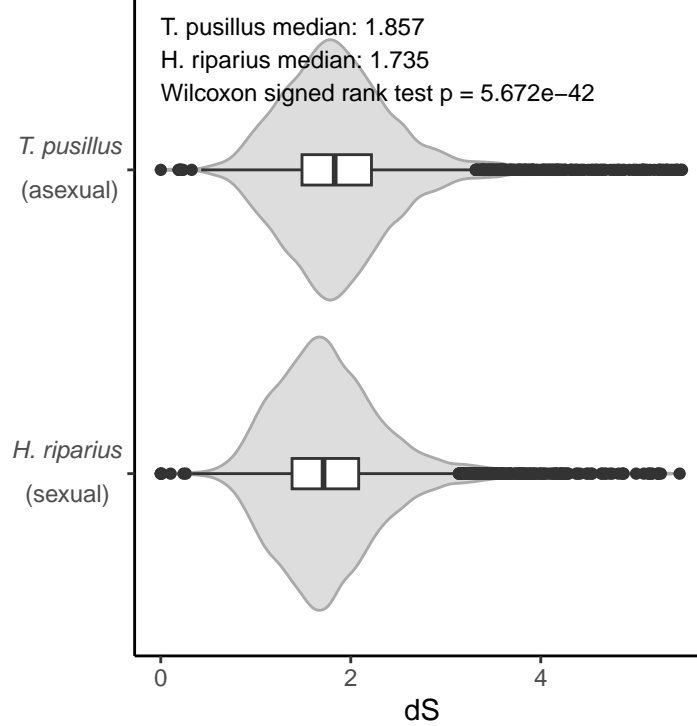
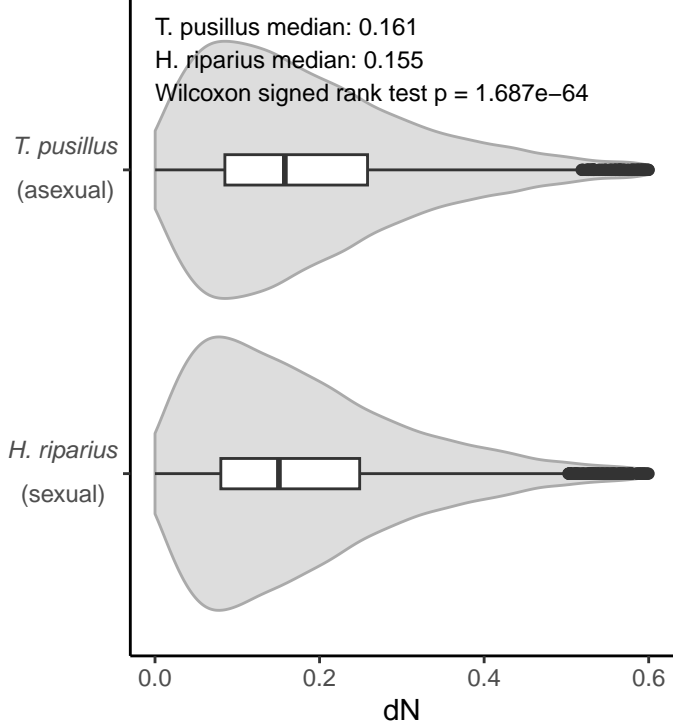
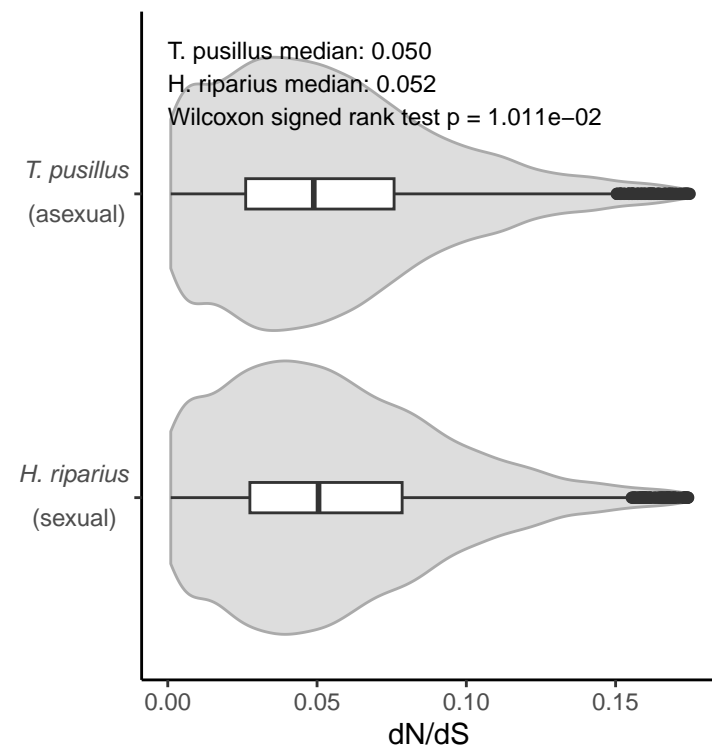
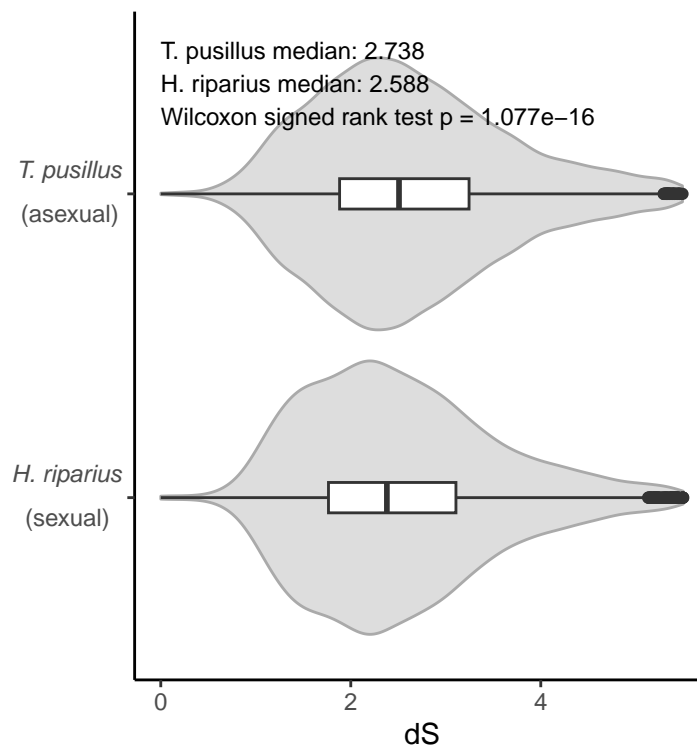
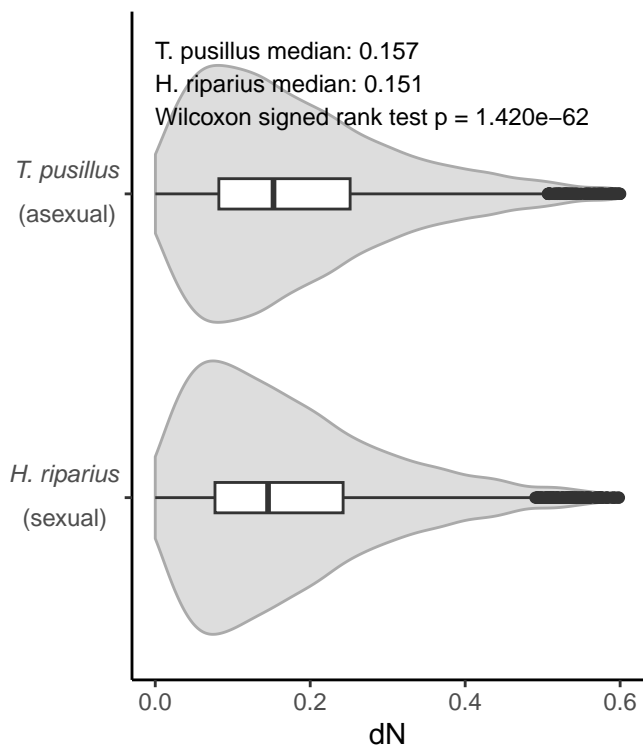


Model with equal codon frequencies  
(CodonFreq=0)



Codon frequencies estimated from  
average nucleotide frequencies  
(CodonFreq=1)



Codon frequencies estimated from average  
nucleotide frequencies at the each codon position  
(CodonFreq=2)

