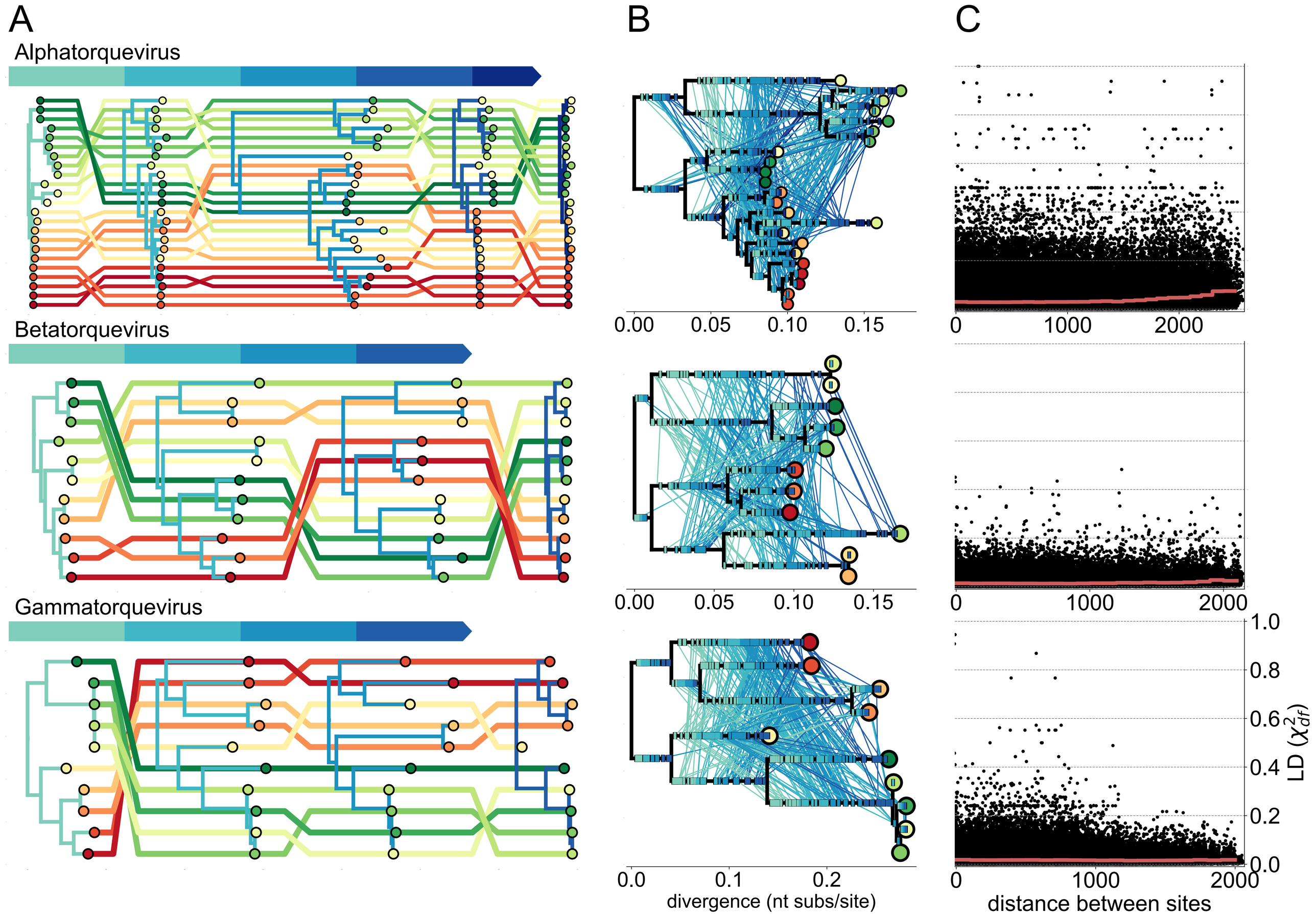
A
Alphatorquevirus
Betatorquevirus
Gammatorquevirus
B
0.00
0.05
0.10
0.15
0.0
0.1
0.2
divergence (nt subs/site)
C
0
1000
2000
1.0
0.8
0.6
0.4
0.2
0.0
LD (χ^2_{df})
distance between sites