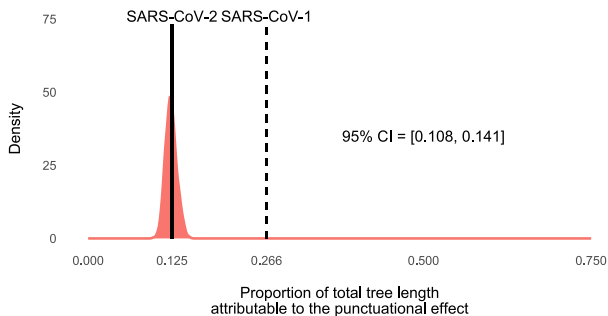
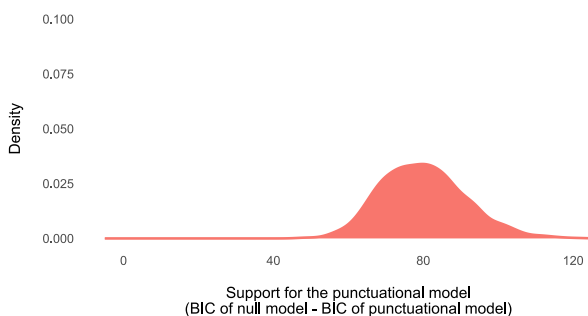


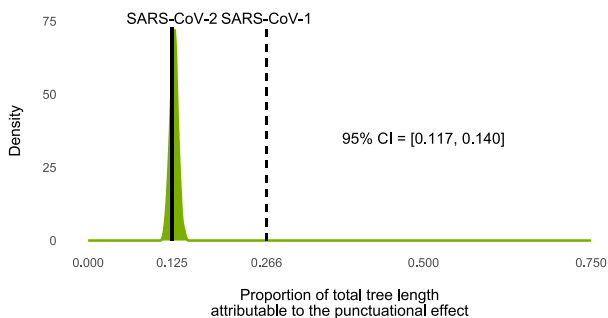
(1A) Subsampling 1,000 genomes 1,000 times
Simple random sampling



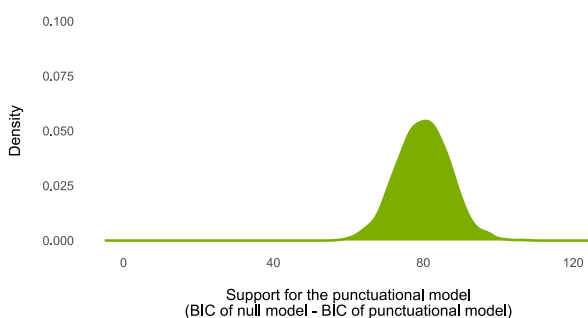
(1B) Subsampling 1,000 genomes 1,000 times
Simple random sampling



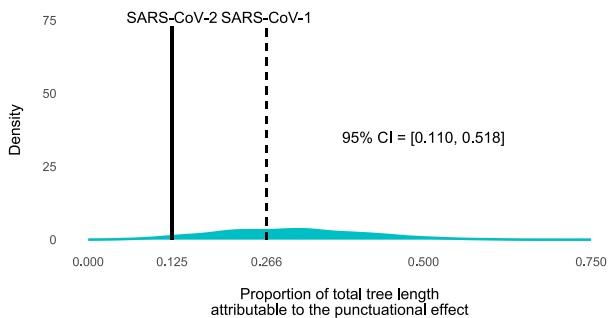
(2A) Subsampling 1,000 genomes 1,000 times
Using SARS-CoV-1 by-continent sampling proportion



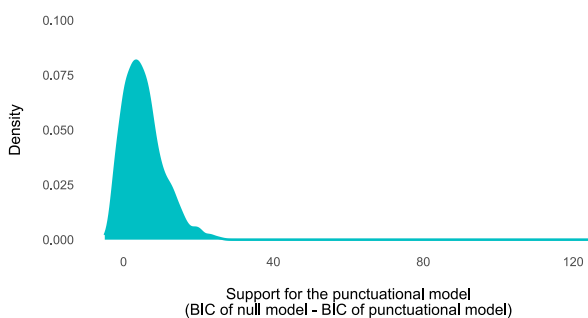
(2B) Subsampling 1,000 genomes 1,000 times
Using SARS-CoV-1 by-continent sampling proportion



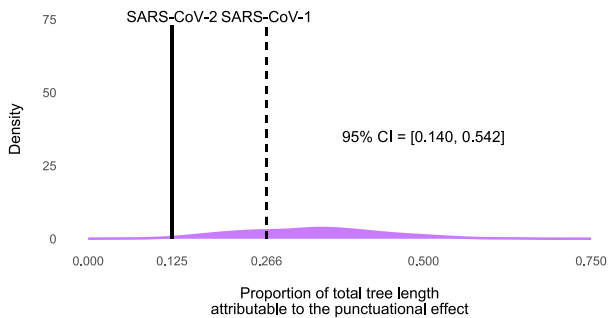
(3A) Subsampling 42 genomes 1,000 times
Simple random sampling



(3B) Subsampling 42 genomes 1,000 times
Simple random sampling



(4A) Subsampling 42 genomes 1,000 times
Using SARS-CoV-1 by-continent sampling proportion



(4B) Subsampling 42 genomes 1,000 times
Using SARS-CoV-1 by-continent sampling proportion

