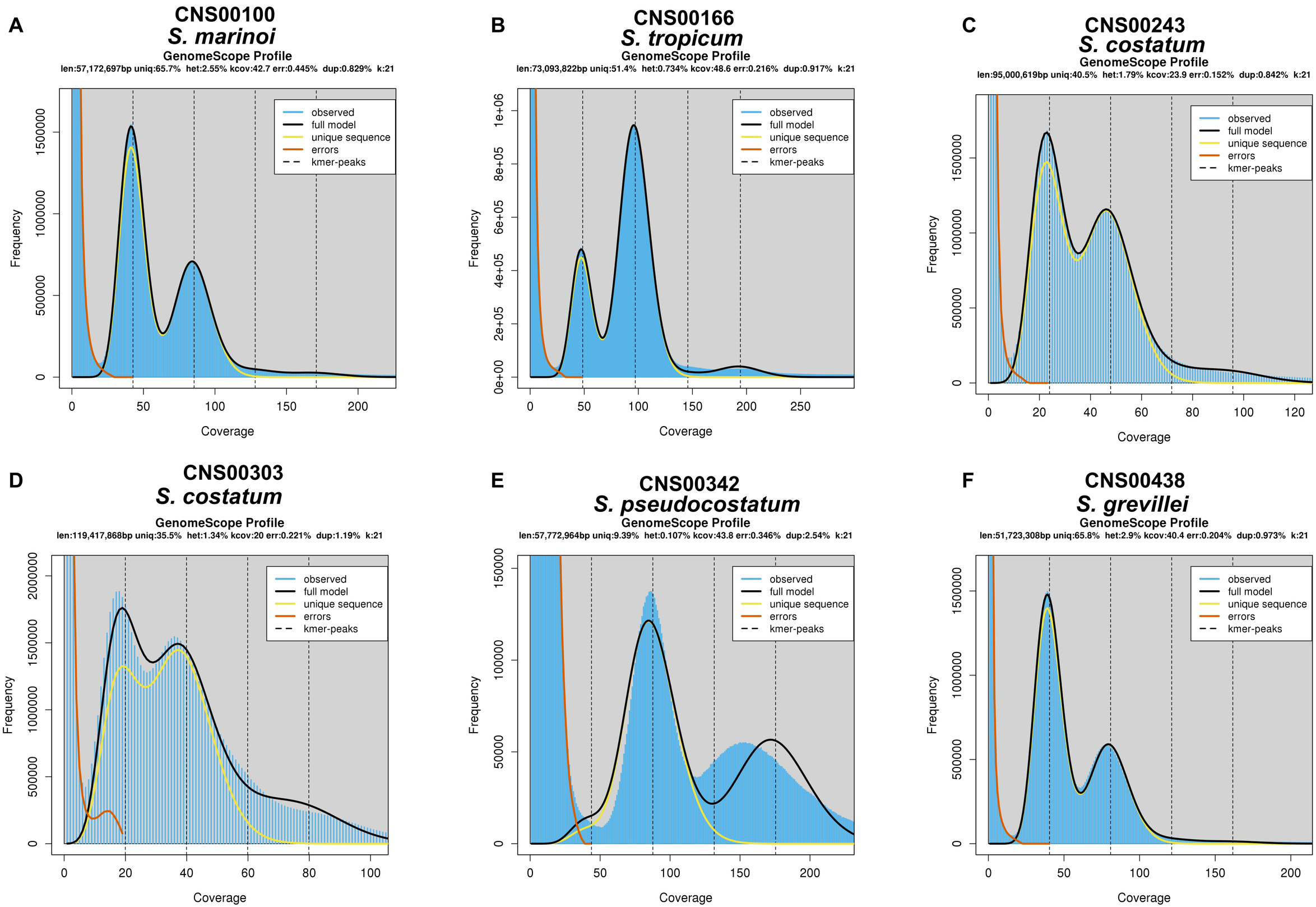
A
CNS00100
S. marinoi
GenomeScope Profile
len:57,172,697bp uniq:65.7% het:2.55% kcov:42.7 err:0.445% dup:0.829% k:21
observed
full model
unique sequence
errors
kmer-peaks
Frequency
Coverage
B
CNS00166
S. tropicum
GenomeScope Profile
len:73,093,822bp uniq:51.4% het:0.734% kcov:48.6 err:0.216% dup:0.917% k:21
observed
full model
unique sequence
errors
kmer-peaks
Frequency
Coverage
C
CNS00243
S. costatum
GenomeScope Profile
len:95,000,619bp uniq:40.5% het:1.79% kcov:23.9 err:0.152% dup:0.842% k:21
observed
full model
unique sequence
errors
kmer-peaks
Frequency
Coverage
D
CNS00303
S. costatum
GenomeScope Profile
len:119,417,868bp uniq:35.5% het:1.34% kcov:20 err:0.221% dup:1.19% k:21
observed
full model
unique sequence
errors
kmer-peaks
Frequency
Coverage
E
CNS00342
S. pseudocostatum
GenomeScope Profile
len:57,772,964bp uniq:9.39% het:0.107% kcov:43.8 err:0.346% dup:2.54% k:21
observed
full model
unique sequence
errors
kmer-peaks
Frequency
Coverage
F
CNS00438
S. grevillei
GenomeScope Profile
len:51,723,308bp uniq:65.8% het:2.9% kcov:40.4 err:0.204% dup:0.973% k:21
observed
full model
unique sequence
errors
kmer-peaks
Frequency
Coverage