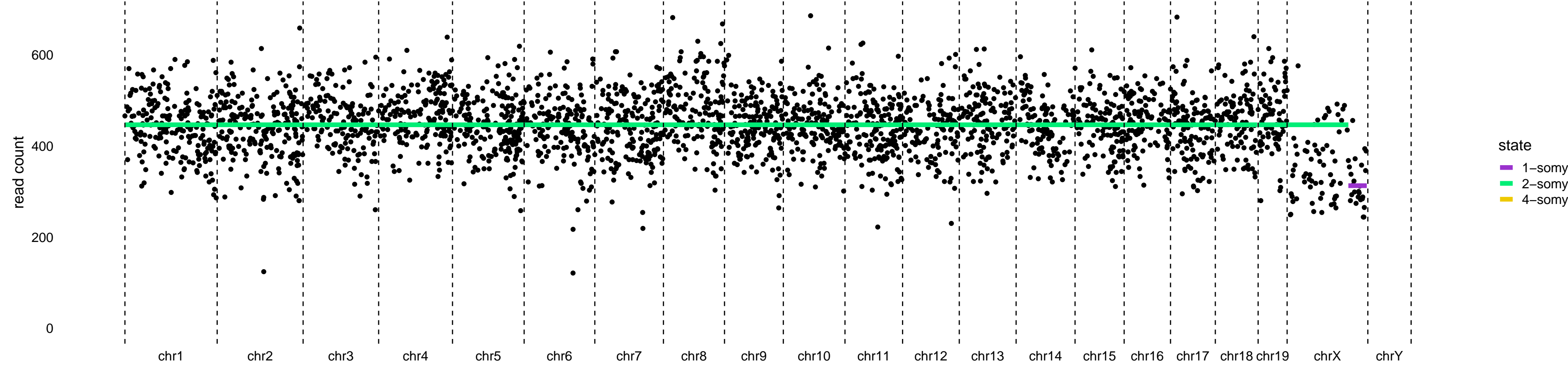# DNA_140619_B1-sc-1996973_sorted_dedup.bam

reads = 1.29M, complexity = 212.45M, spikiness = 0.15, entropy = 7.96, bhattacharyya = 0.85, num.segments = 22, loglik = NA, sos = 12264388

state
- 1-somy
- 2-somy
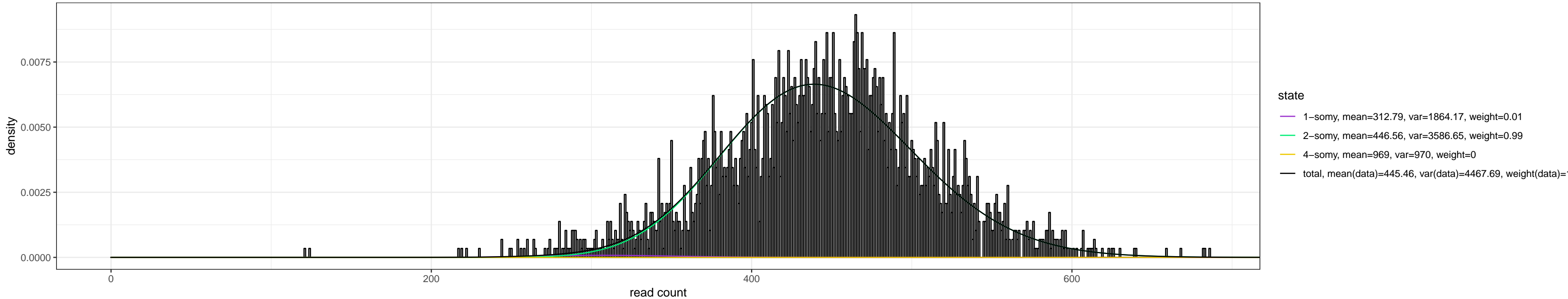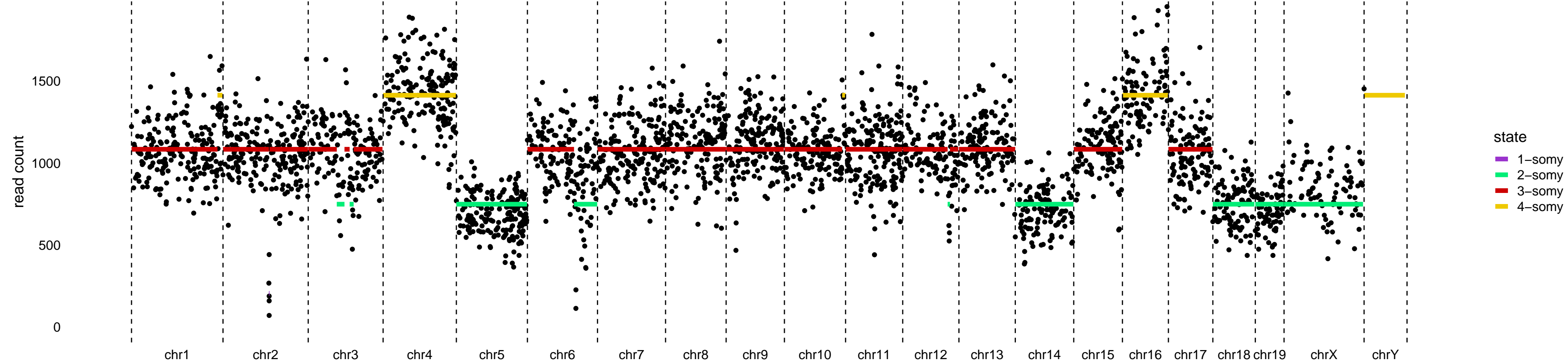- 4-somy

DNA_140619_B1-sc-1996973_sorted_dedup.bam

reads = 1.29M, complexity = 212.45M, spikiness = 0.15, entropy = 7.96, bhattacharyya = 0.85, num.segments = 22, loglik = NA, sos = 12264388
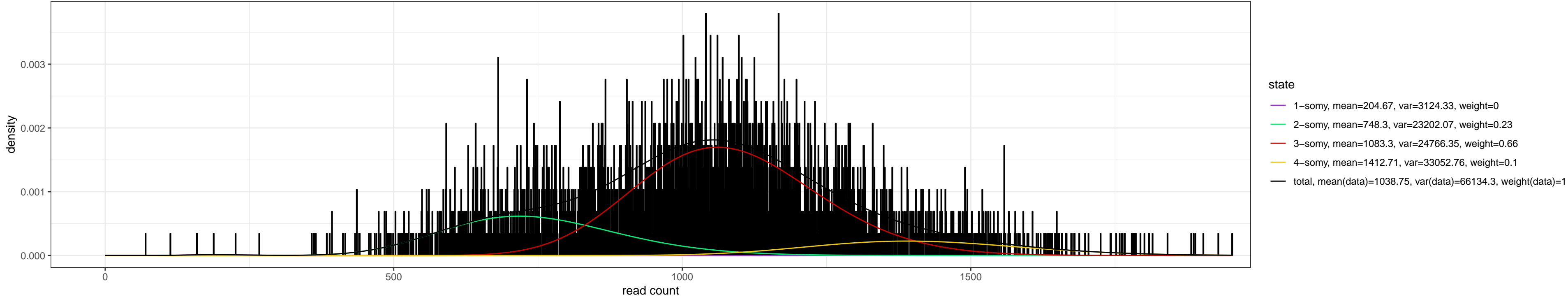

state
- 1-somy, mean=312.79, var=1864.17, weight=0.01
- 2-somy, mean=446.56, var=3586.65, weight=0.99
- 4-somy, mean=969, var=970, weight=0
- total, mean(data)=445.46, var(data)=4467.69, weight(data)=1

DNA_140619_B2–sc–1996974_sorted_dedup.bam

reads = 3.01M, complexity = 258.02M, spikiness = 0.15, entropy = 7.94, bhattacharyya = 3.85, num.segments = 32, loglik = NA, sos = 86944167

DNA_140619_B2–sc–1996974_sorted_dedup.bam

reads = 3.01M, complexity = 258.02M, spikiness = 0.15, entropy = 7.94, bhattacharyya = 3.85, num.segments = 32, loglik = NA, sos = 86944167
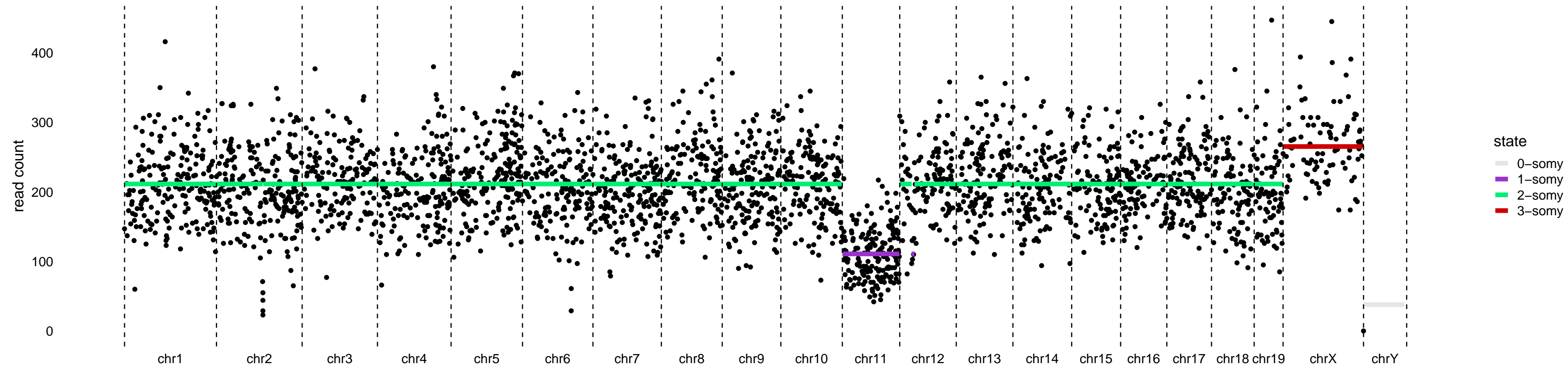

state

1–somy, mean=204.67, var=3124.33, weight=0

2–somy, mean=748.3, var=23202.07, weight=0.23

3–somy, mean=1083.3, var=24766.35, weight=0.66

4–somy, mean=1412.71, var=33052.76, weight=0.1

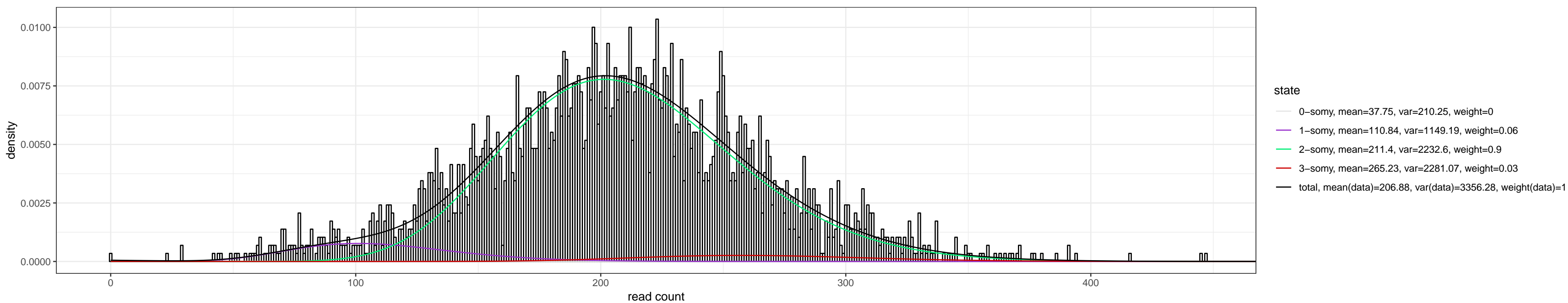total, mean(data)=1038.75, var(data)=66134.3, weight(data)=1

DNA_140619_B3-sc-1996975_sorted_dedup.bam

reads = 0.6M, complexity = 46.01M, spikiness = 0.26, entropy = 7.93, bhattacharyya = 0.78, num.segments = 25, loglik = NA, sos = 7488140
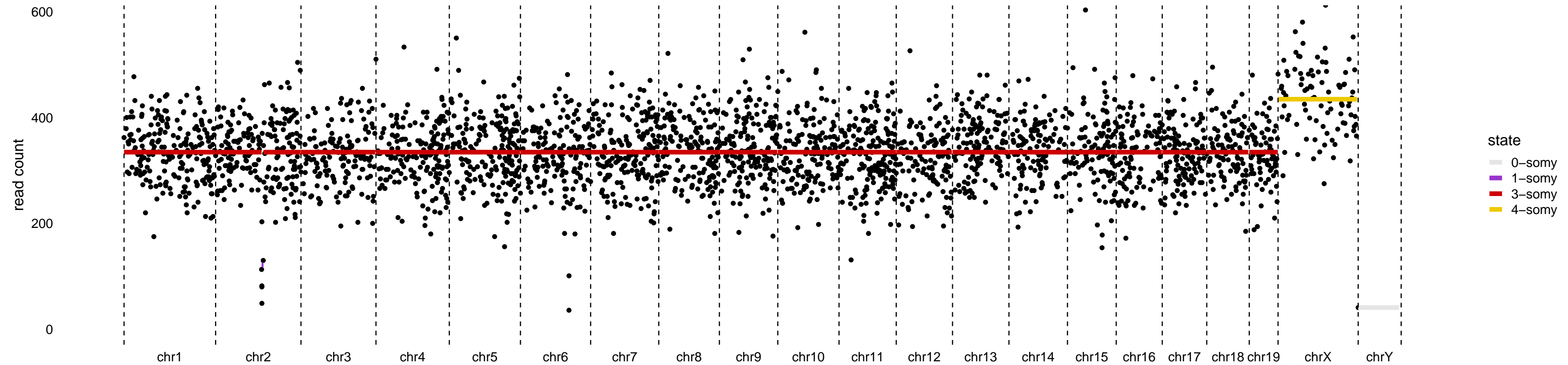

DNA_140619_B3-sc-1996975_sorted_dedup.bam

reads = 0.6M, complexity = 46.01M, spikiness = 0.26, entropy = 7.93, bhattacharyya = 0.78, num.segments = 25, loglik = NA, sos = 7488140
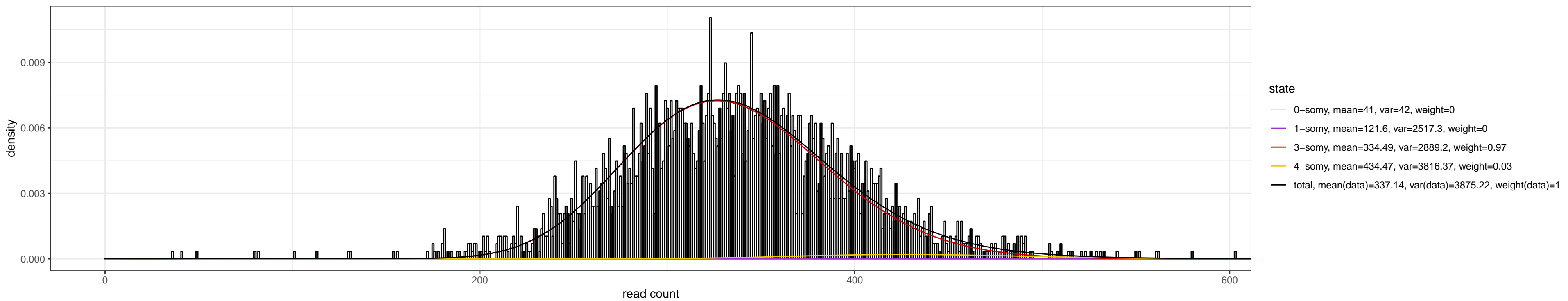
DNA_140619_B4-sc-1996976_sorted_dedup.bam

reads = 0.98M, complexity = 175.03M, spikiness = 0.18, entropy = 7.95, bhattacharyya = 1.85, num.segments = 23, loglik = NA, sos = 10025072
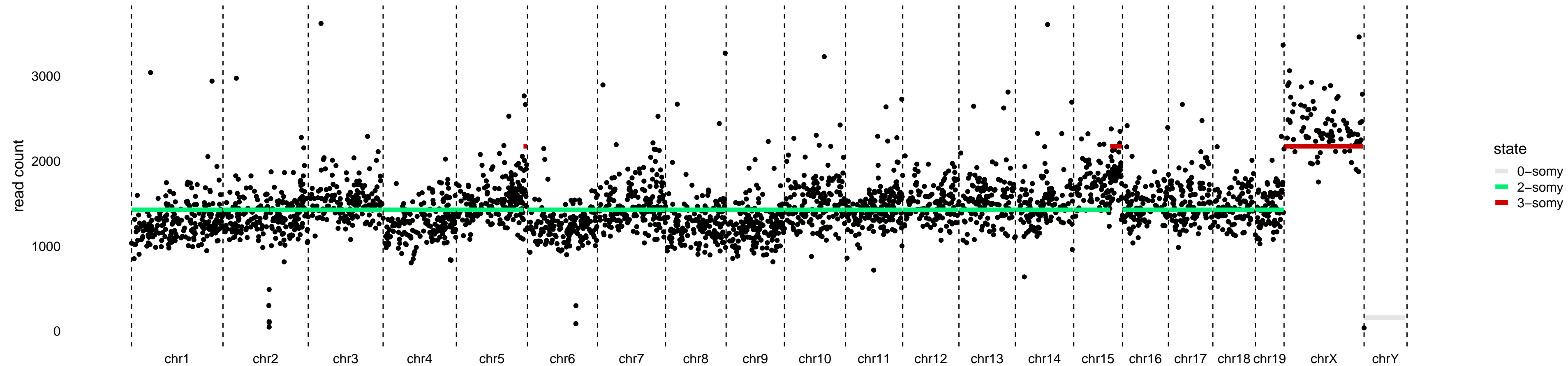

DNA_140619_B4-sc-1996976_sorted_dedup.bam

reads = 0.98M, complexity = 175.03M, spikiness = 0.18, entropy = 7.95, bhattacharyya = 1.85, num.segments = 23, loglik = NA, sos = 10025072
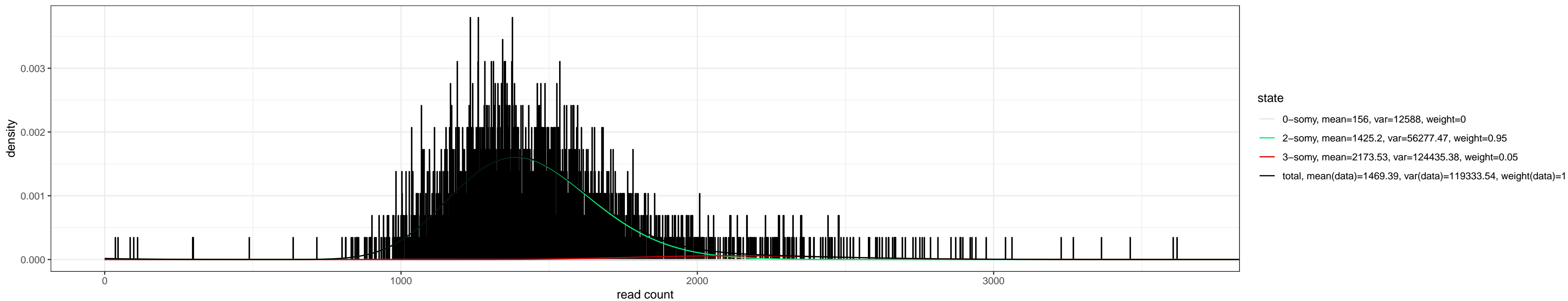
DNA_140619_B5−sc−1996977_sorted_dedup.bam

reads = 4.26M, complexity = 95.81M, spikiness = 0.16, entropy = 7.95, bhattacharyya = 0.82, num.segments = 27, loglik = NA, sos = 255648560
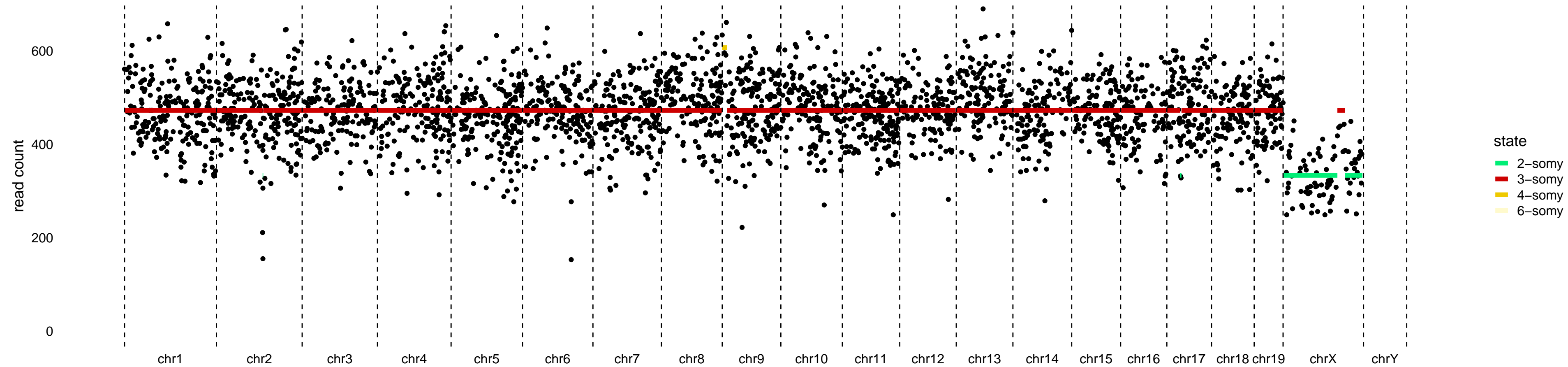

DNA_140619_B5−sc−1996977_sorted_dedup.bam

state

0−somy, mean=156, var=12588, weight=0

2−somy, mean=1425.2, var=56277.47, weight=0.95

3−somy, mean=2173.53, var=124435.38, weight=0.05

total, mean(data)=1469.39, var(data)=119333.54, weight(data)=1

DNA_140619_B7−sc−1996979_sorted_dedup.bam

reads = 1.36M, complexity = 189.87M, spikiness = 0.14, entropy = 7.96, bhattacharyya = 0.83, num.segments = 28, loglik = NA, sos = 11492682
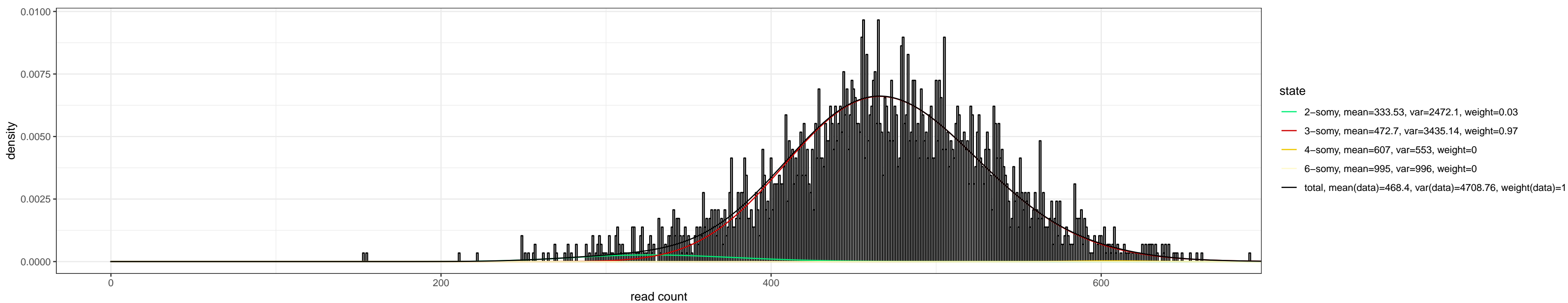

DNA_140619_B7−sc−1996979_sorted_dedup.bam

reads = 1.36M, complexity = 189.87M, spikiness = 0.14, entropy = 7.96, bhattacharyya = 0.83, num.segments = 28, loglik = NA, sos = 11492682
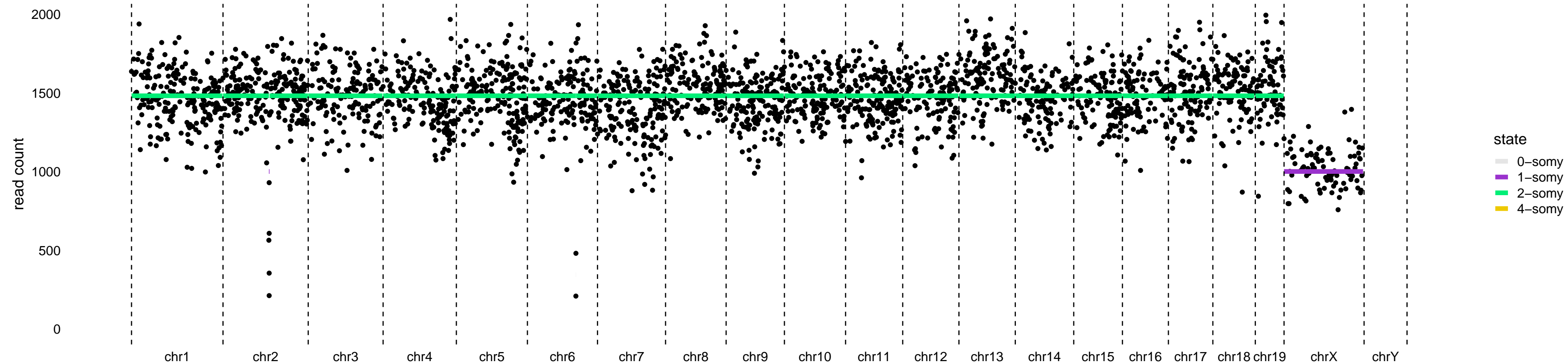
DNA_140619_B8−sc−1996980_sorted_dedup.bam

reads = 4.25M, complexity = 266.81M, spikiness = 0.11, entropy = 7.96, bhattacharyya = 1.26, num.segments = 25, loglik = NA, sos = 77904535

DNA_140619_B8−sc−1996980_sorted_dedup.bam

reads = 4.25M, complexity = 266.81M, spikiness = 0.11, entropy = 7.96, bhattacharyya = 1.26, num.segments = 25, loglik = NA, sos = 77904535

state

- 0−somy, mean=345, var=36992, weight=0
- 1−somy, mean=1001.13, var=22446.91, weight=0.03
- 2−somy, mean=1482.33, var=23055.49, weight=0.97
- 4−somy, mean=3271, var=3272, weight=0
- total, mean(data)=1465.93, var(data)=36263.97, weight(data)=1