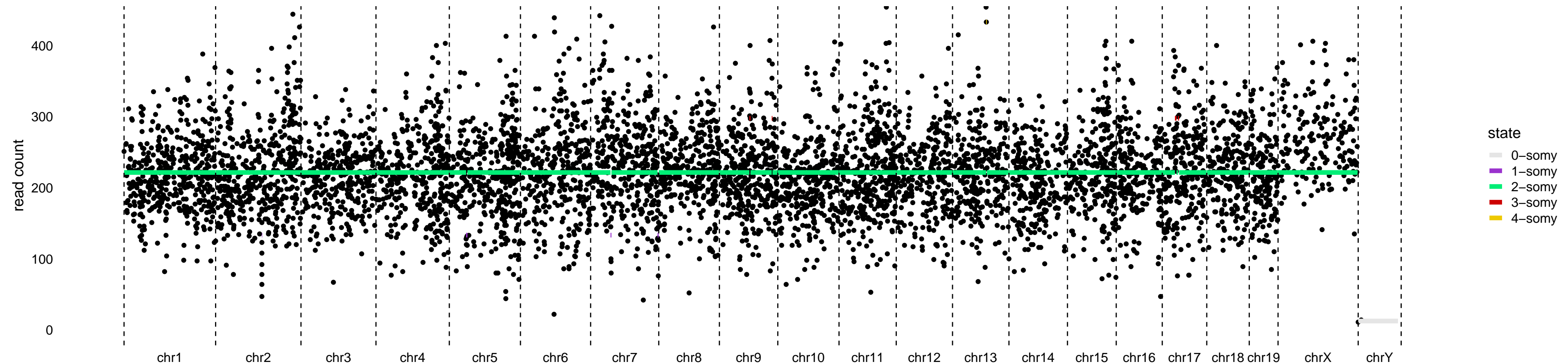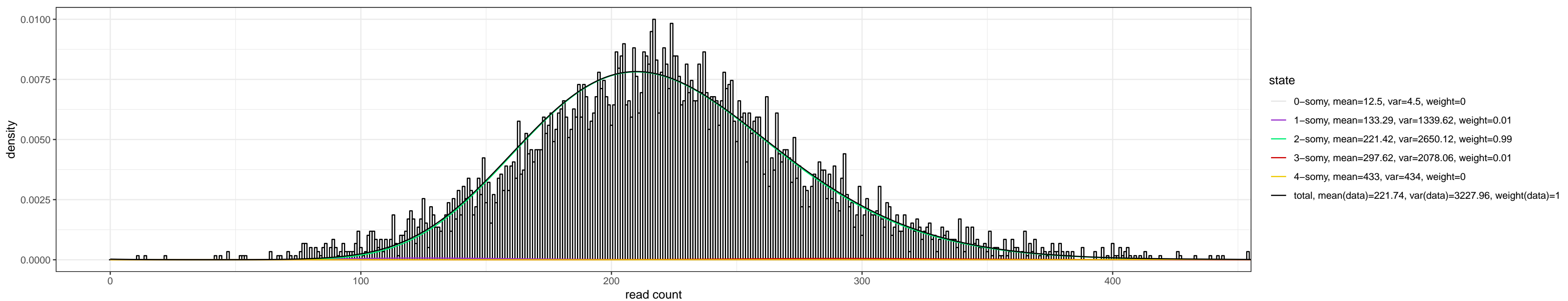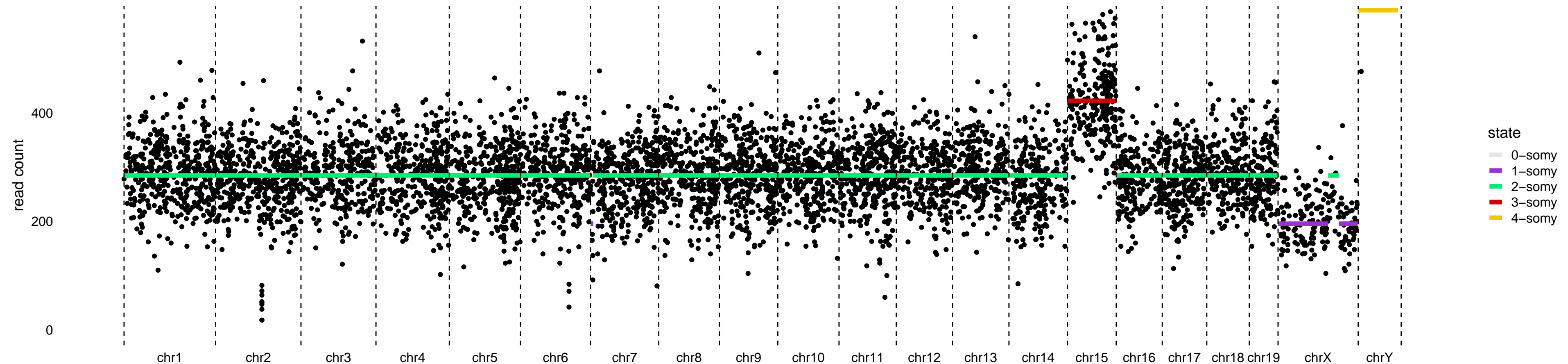# DNA_140619_C1-sc-1996983_sorted_dedup.bam

reads = 1.31M, complexity = 197.65M, spikiness = 0.25, entropy = 8.65, bhattacharyya = 0.51, num.segments = 38, loglik = NA, sos = 18396183

DNA_140619_C1-sc-1996983_sorted_dedup.bam

reads = 1.31M, complexity = 197.65M, spikiness = 0.25, entropy = 8.65, bhattacharyya = 0.51, num.segments = 38, loglik = NA, sos = 18396183

state

0-somy, mean=12.5, var=4.5, weight=0
1-somy, mean=133.29, var=1339.62, weight=0.01
2-somy, mean=221.42, var=2650.12, weight=0.99
3-somy, mean=297.62, var=2078.06, weight=0.01
4-somy, mean=433, var=434, weight=0
total, mean(data)=221.74, var(data)=3227.96, weight(data)=1
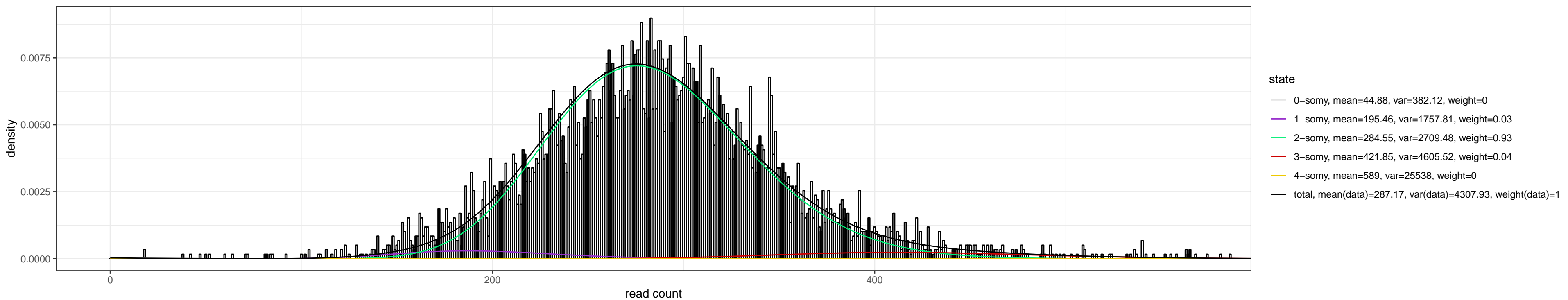
DNA_140619_C2-sc-1996984_sorted_dedup.bam

reads = 1.69M, complexity = 176.02M, spikiness = 0.2, entropy = 8.66, bhattacharyya = 0.45, num.segments = 28, loglik = NA, sos = 18899768
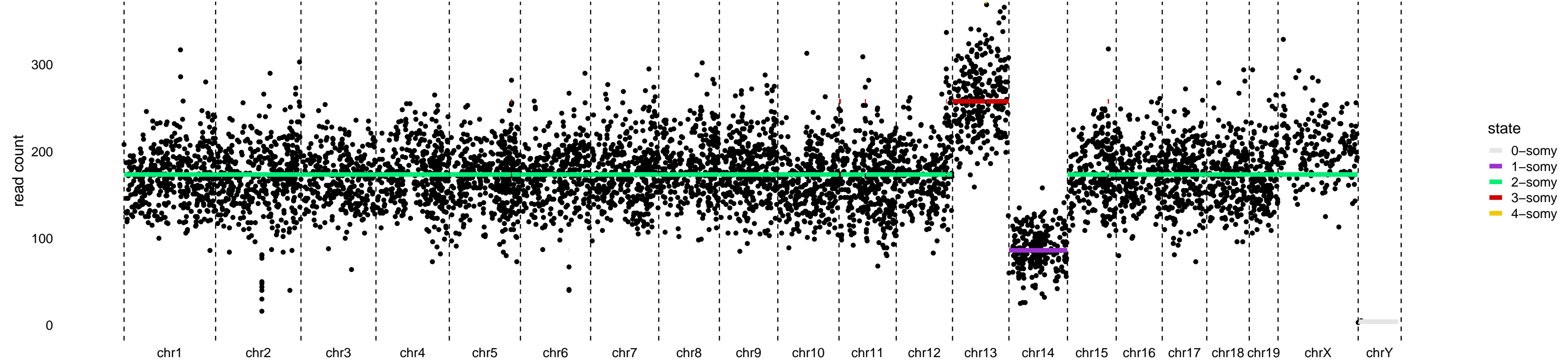

DNA_140619_C2-sc-1996984_sorted_dedup.bam

reads = 1.69M, complexity = 176.02M, spikiness = 0.2, entropy = 8.66, bhattacharyya = 0.45, num.segments = 28, loglik = NA, sos = 18899768
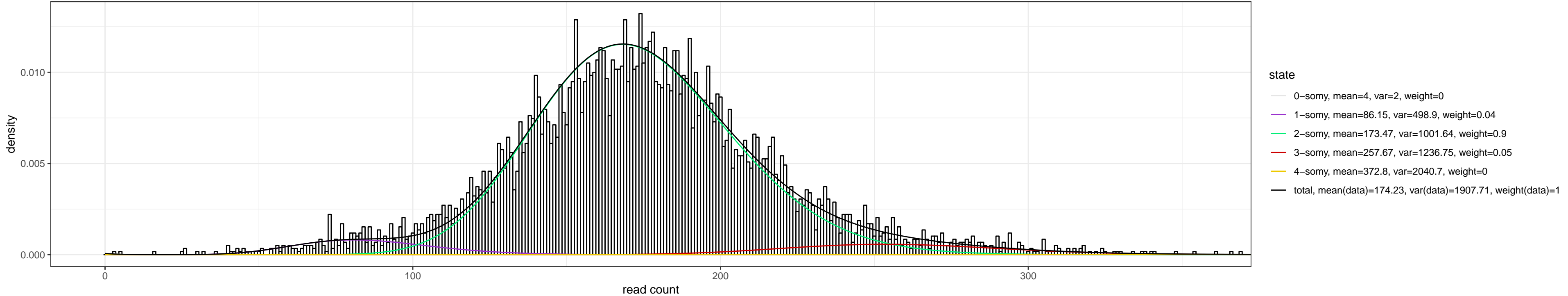

state

- 0-somy, mean=44.88, var=382.12, weight=0
- 1-somy, mean=195.46, var=1757.81, weight=0.03
- 2-somy, mean=284.55, var=2709.48, weight=0.93
- 3-somy, mean=421.85, var=4605.52, weight=0.04
- 4-somy, mean=589, var=25538, weight=0
- total, mean(data)=287.17, var(data)=4307.93, weight(data)=1

DNA_140619_C3−sc−1996985_sorted_dedup.bam

reads = 1.03M, complexity = 191.69M, spikiness = 0.19, entropy = 8.65, bhattacharyya = 1.32, num.segments = 36, loglik = NA, sos = 6833390
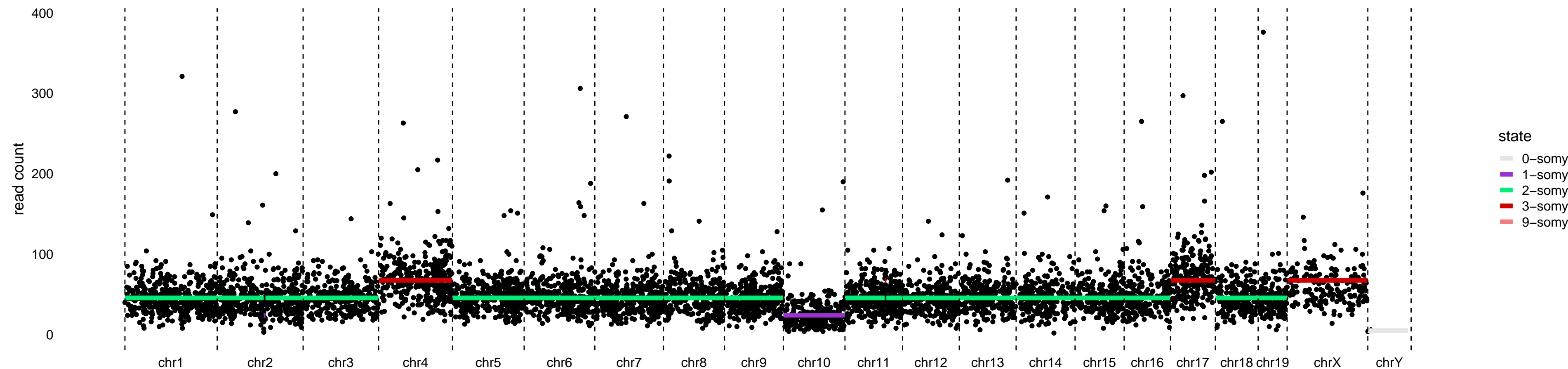

DNA_140619_C3−sc−1996985_sorted_dedup.bam

state

- 0−somy, mean=4, var=2, weight=0
- 1−somy, mean=86.15, var=498.9, weight=0.04
- 2−somy, mean=173.47, var=1001.64, weight=0.9
- 3−somy, mean=257.67, var=1236.75, weight=0.05
- 4−somy, mean=372.8, var=2040.7, weight=0
- total, mean(data)=174.23, var(data)=1907.71, weight(data)=1
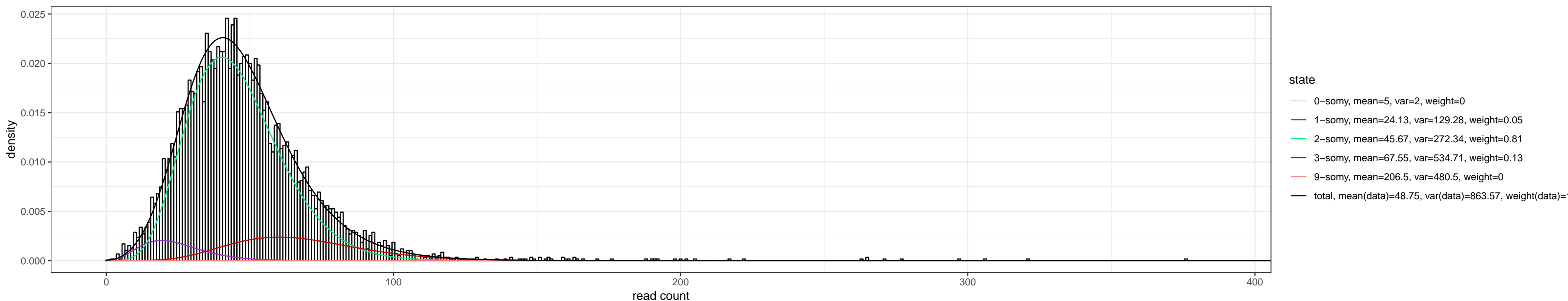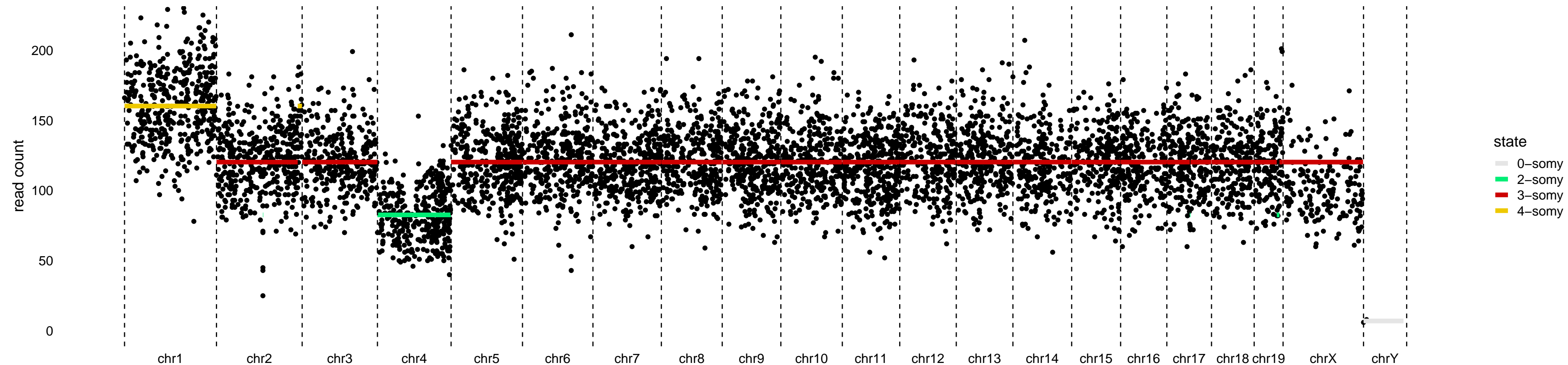
DNA_140619_C4-sc-1996986_sorted_dedup.bam

reads = 0.29M, complexity = 23.64M, spikiness = 0.44, entropy = 8.56, bhattacharyya = 0.31, num.segments = 27, loglik = NA, sos = 4503097

DNA_140619_C4-sc-1996986_sorted_dedup.bam

reads = 0.29M, complexity = 23.64M, spikiness = 0.44, entropy = 8.56, bhattacharyya = 0.31, num.segments = 27, loglik = NA, sos = 4503097

state

- 0-somy, mean=5, var=2, weight=0
- 1-somy, mean=24.13, var=129.28, weight=0.05
- 2-somy, mean=45.67, var=272.34, weight=0.81
- 3-somy, mean=67.55, var=534.71, weight=0.13
- 9-somy, mean=206.5, var=480.5, weight=0
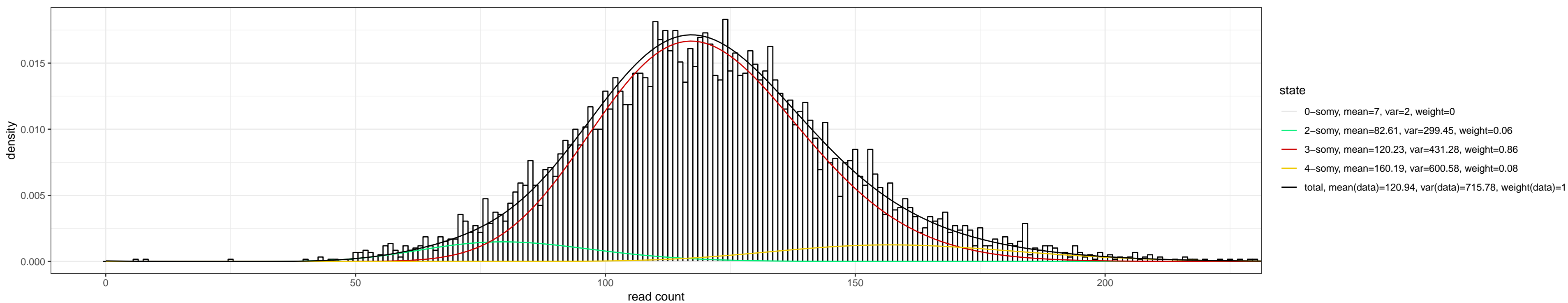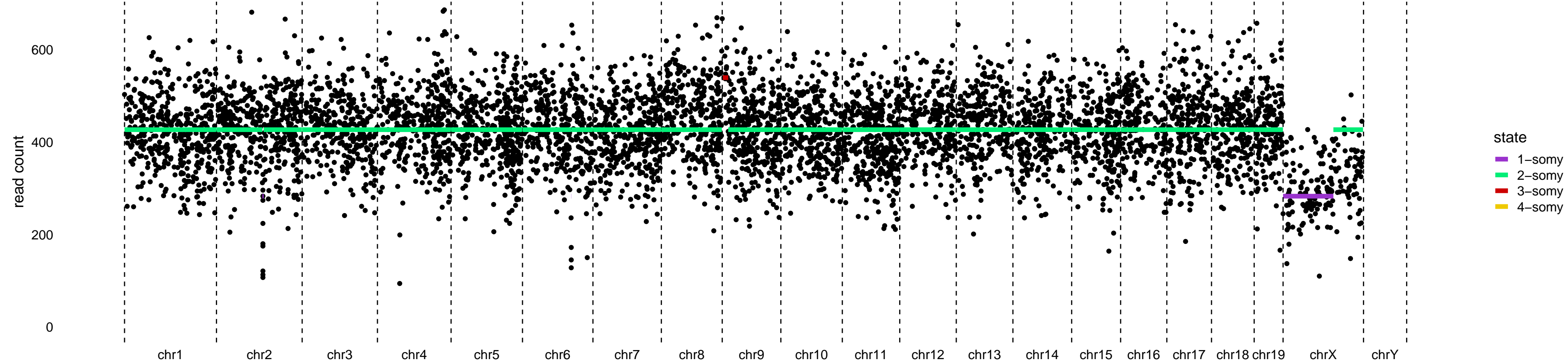- total, mean(data)=48.75, var(data)=863.57, weight(data)=1

DNA_140619_C5-sc-1996987_sorted_dedup.bam

reads = 0.71M, complexity = 165.05M, spikiness = 0.19, entropy = 8.66, bhattacharyya = 0.49, num.segments = 28, loglik = NA, sos = 2945559
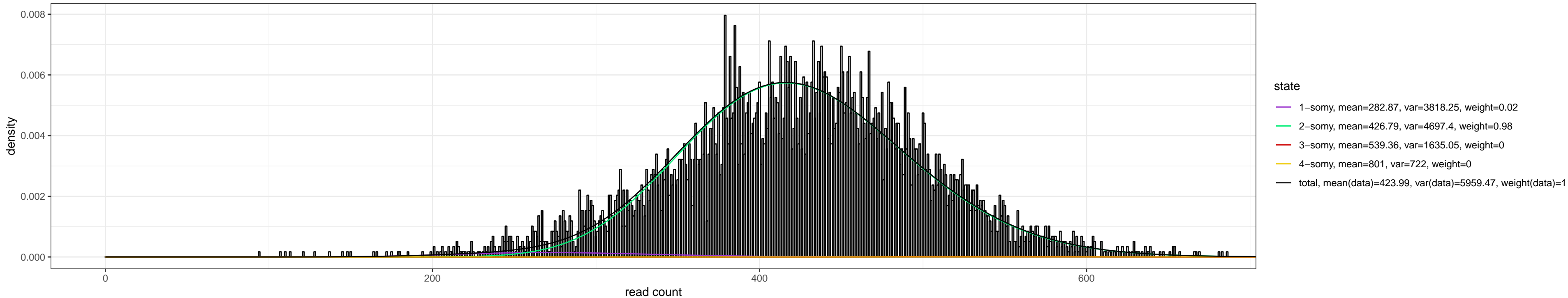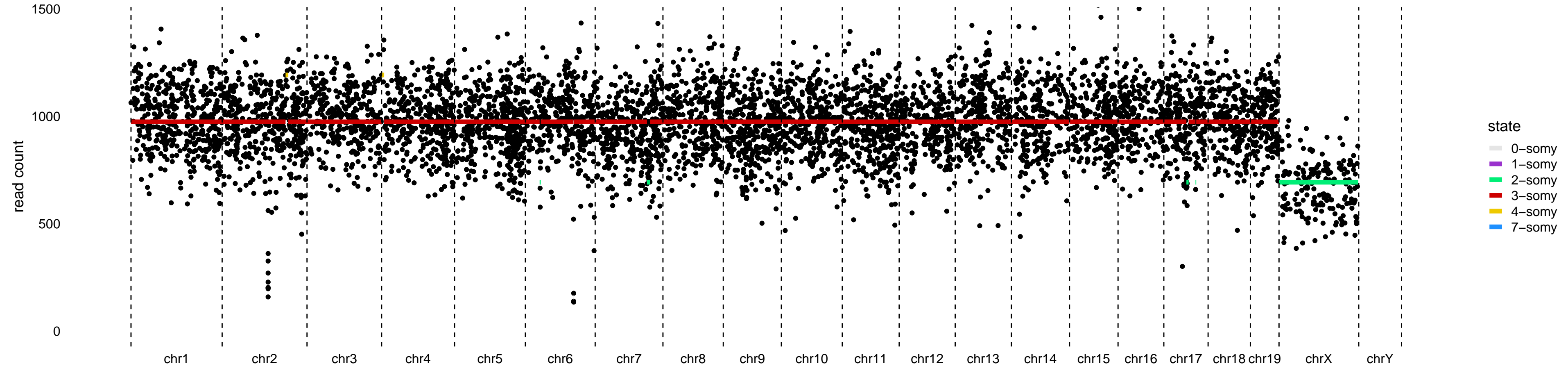
DNA_140619_C7-sc-1996989_sorted_dedup.bam

reads = 2.5M, complexity = 207.41M, spikiness = 0.18, entropy = 8.67, bhattacharyya = 0.62, num.segments = 27, loglik = NA, sos = 32004230

DNA_140619_C7-sc-1996989_sorted_dedup.bam

reads = 2.5M, complexity = 207.41M, spikiness = 0.18, entropy = 8.67, bhattacharyya = 0.62, num.segments = 27, loglik = NA, sos = 32004230

DNA_140619_C8–sc–1996990_sorted_dedup.bam

reads = 5.68M, complexity = 238.5M, spikiness = 0.15, entropy = 8.67, bhattacharyya = 2.91, num.segments = 36, loglik = NA, sos = 118192376