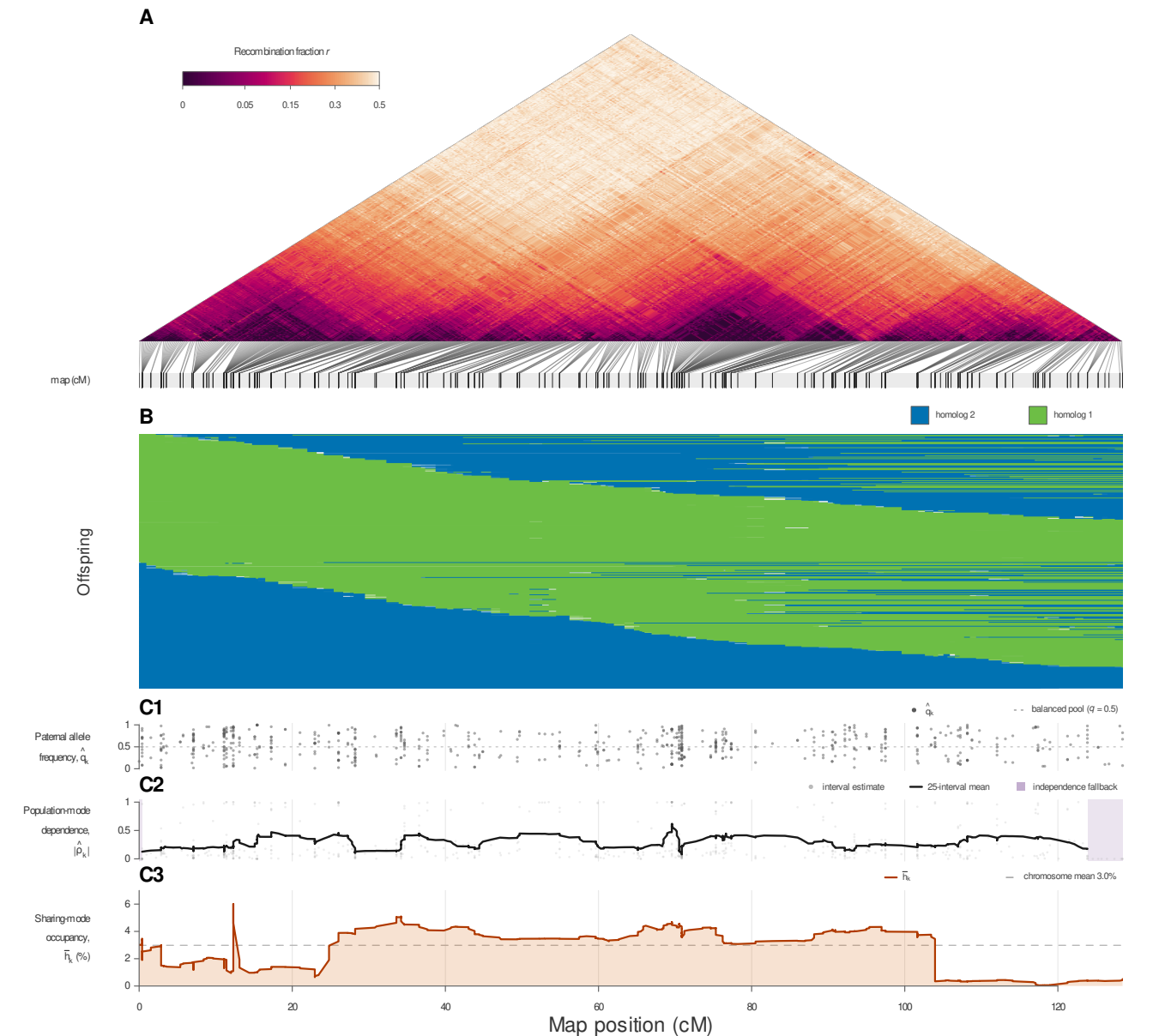

A
Recombination fraction r
0
0.05
0.15
0.3
0.5
map (cM)
B
homolog 2
homolog 1
Offspring
C1
Paternal allele frequency, q̂k
balanced pool (q = 0.5)
C2
Population-mode dependence, |p̂k|
interval estimate
25-interval mean
independence fallback
C3
Sharing-mode occupancy, π̄k (%)
chromosome mean 3.0%
Map position (cM)