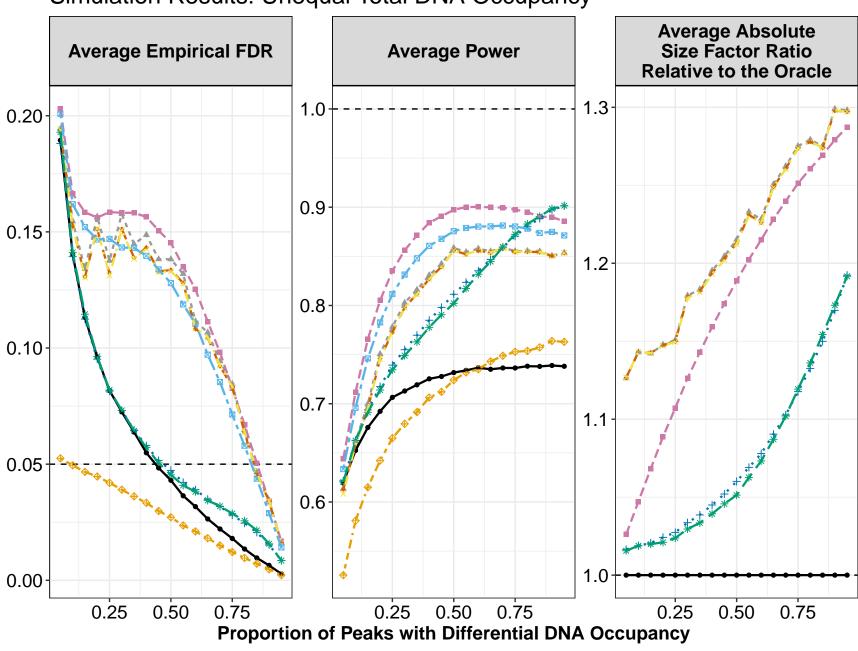
Simulation Results: Unequal Total DNA Occupancy



Technique

- → Oracle
- Library Size (Background Bins)
- -- Library Size (Reads in Peaks)
- → RLE (Background Bins)
- ·+ RLE (Reads in Peaks)
- TMM (Background Bins)
- * TMM (Reads in Peaks)
 - Loess Adjusted Fit (Reads in Peaks)
- MAnorm2 (Common Peaks)