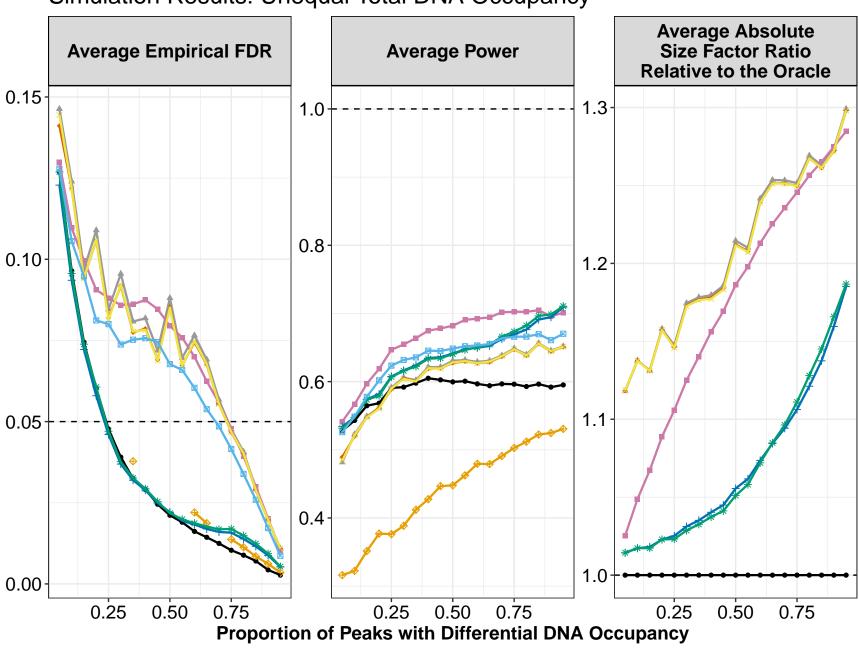
Simulation Results: Unequal Total DNA Occupancy



## **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)