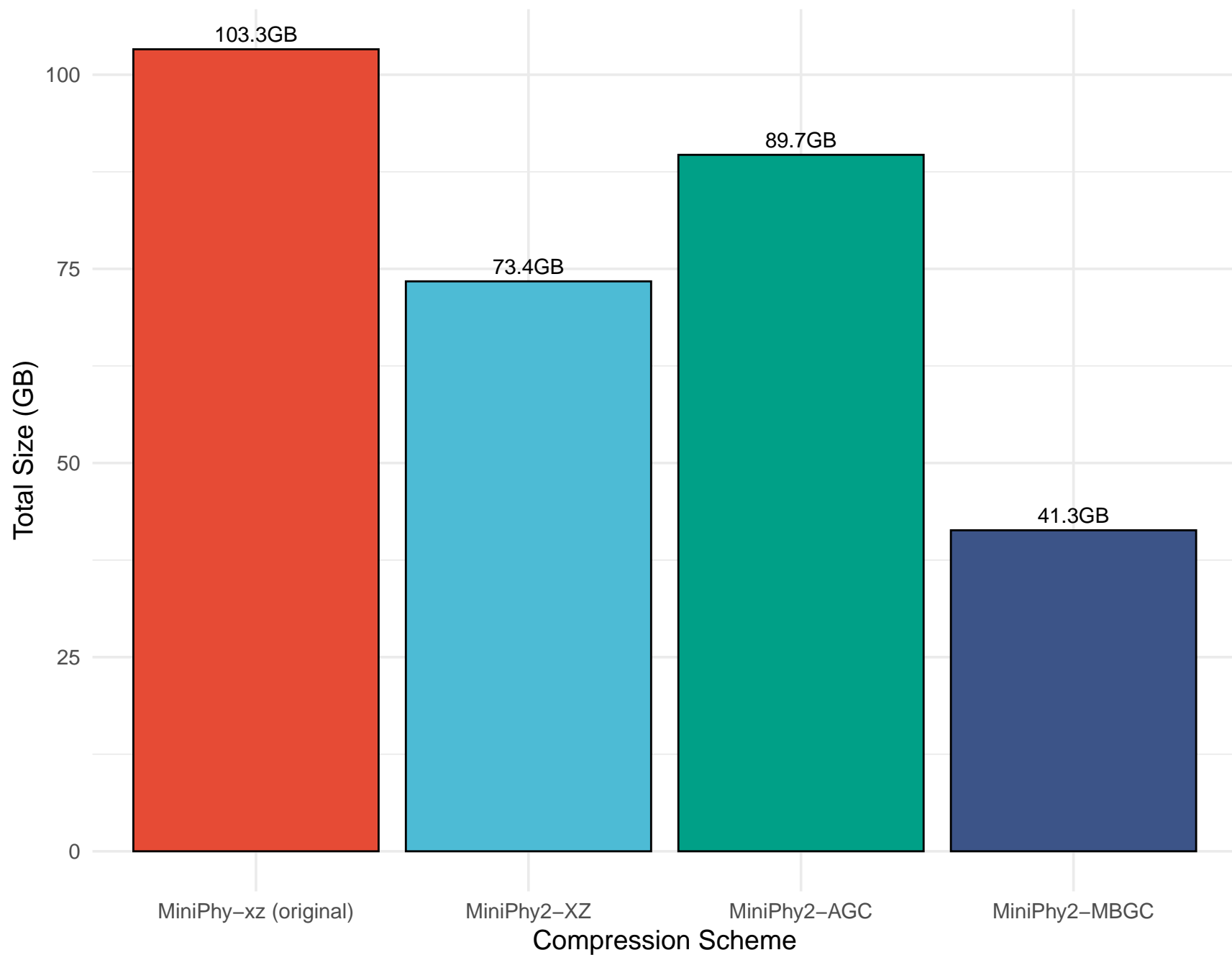


## Compression results – ATB



PARAMETERS: AGC with `-a -b 500 -s 1500`; the AGC reference genome is the first genome in each batch. MBGC with `-m 3`. All with 25 threads. Batches of Species clusters have 4000 genomes. Batches of dustbin and unknown have 1000 genomes for AGC and MBGC, 4000 genomes for XZ.