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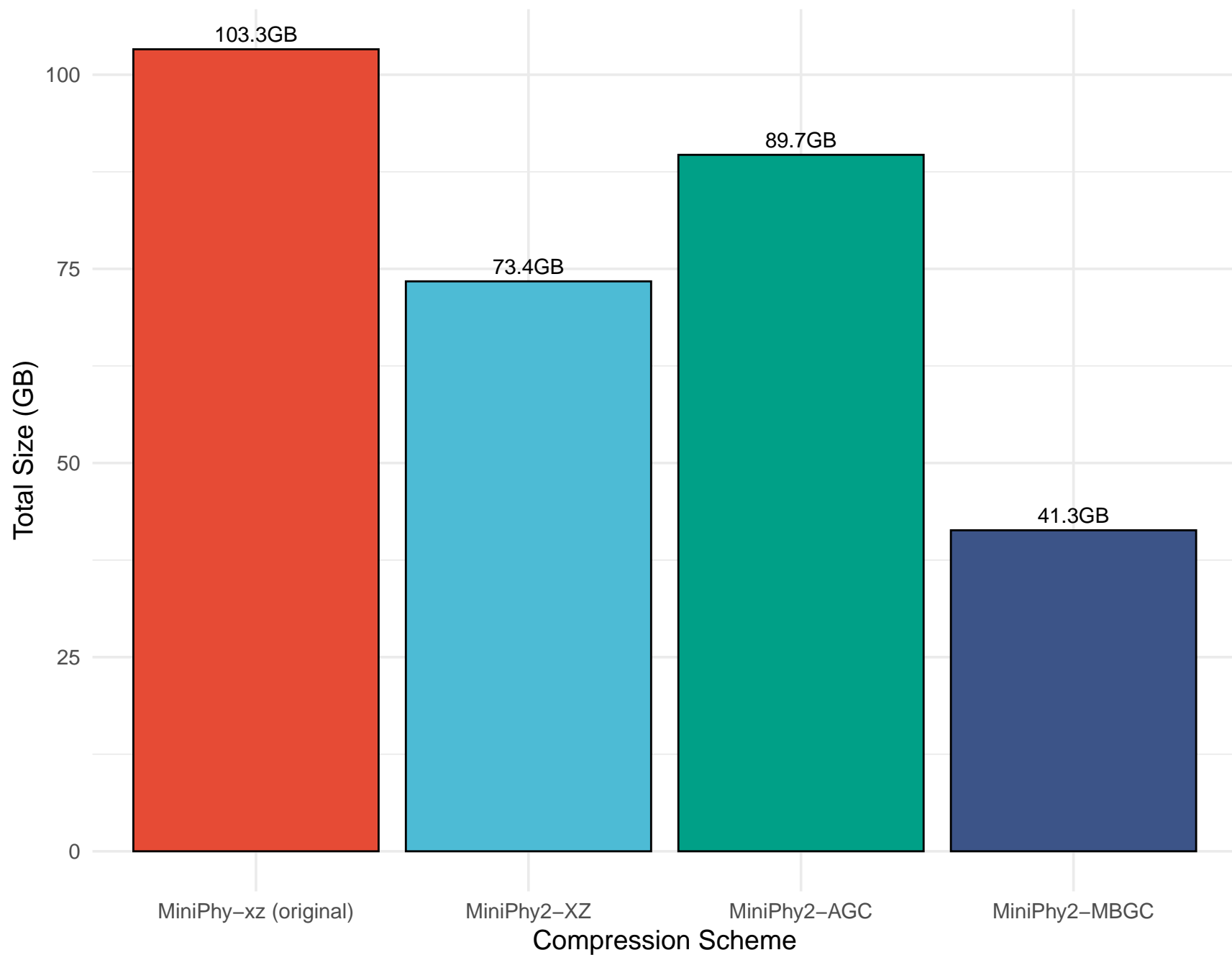
PARAMETERS: AGC with –a –b 500 –s 1500; the AGC reference genome is the first genome in each batch.
MBGC with –m 3. All with 25 threads. Batches of Species clusters have 4000 genomes. Batches of
dustbin and unknown have 1000 genomes for AGC and MBGC, 4000 genomes for XZ.