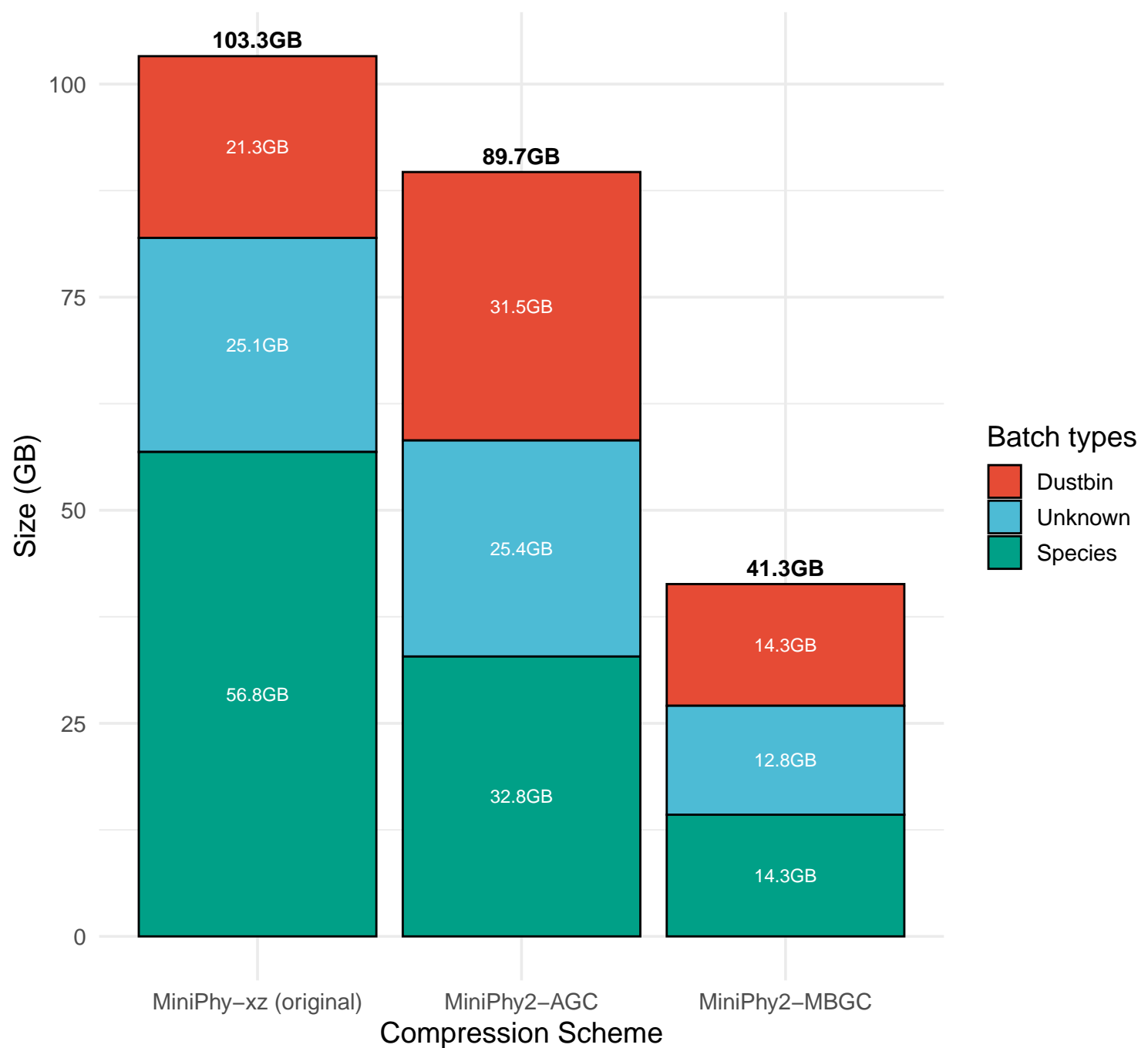


# Compression results – ATB



PARAMETERS: AGC with `-a -b 500 -s 1500`; MBGC with `-m 3`. All with 25 threads. Batches of Species clusters have 4000 genomes, of dustbin and unknown have 1000 genomes.