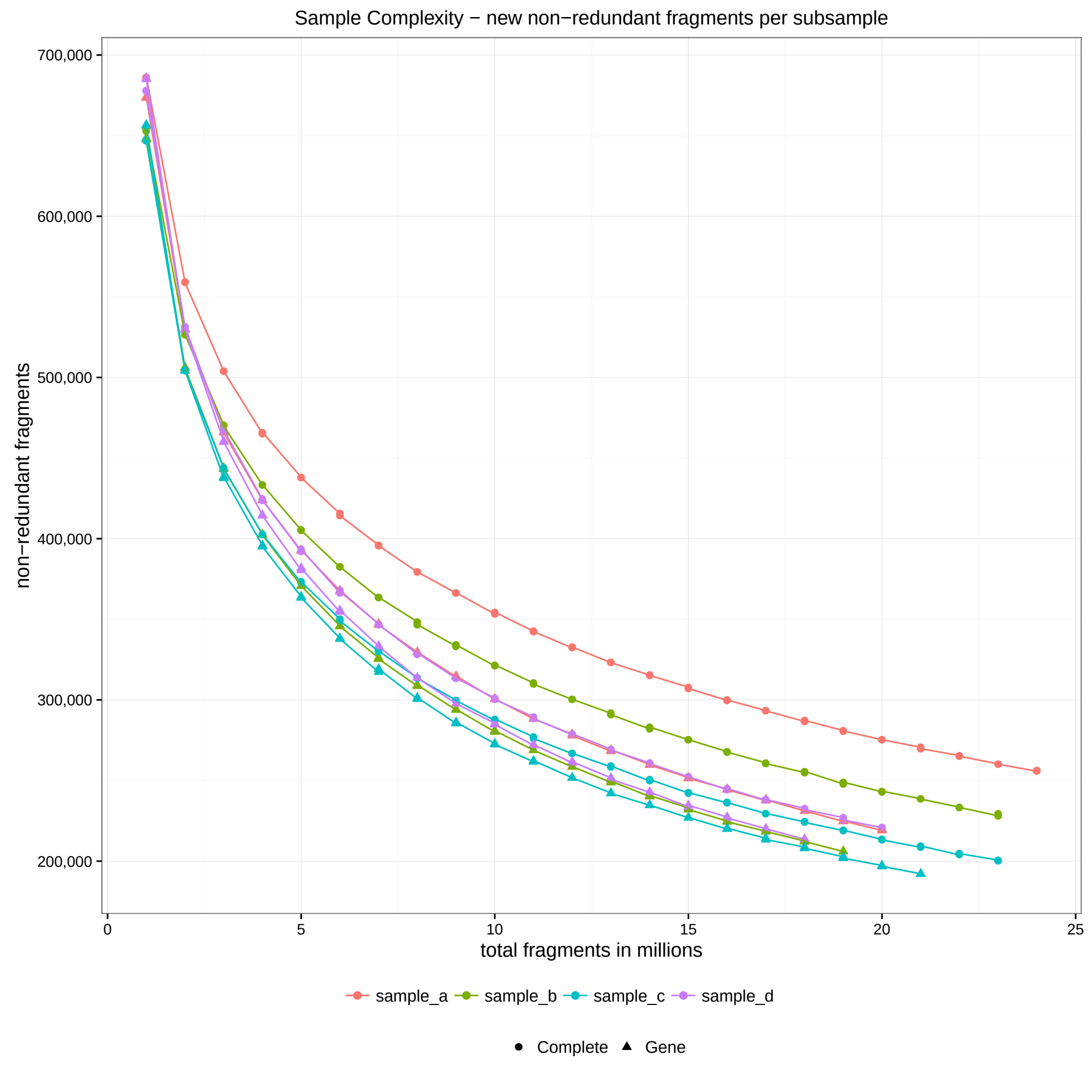

Sample Complexity − new non−redundant fragments per subsample
non−redundant fragments
700,000
600,000
500,000
400,000
300,000
200,000
0
5
10
15
20
25
total fragments in millions
sample_a
sample_b
sample_c
sample_d
Complete
Gene