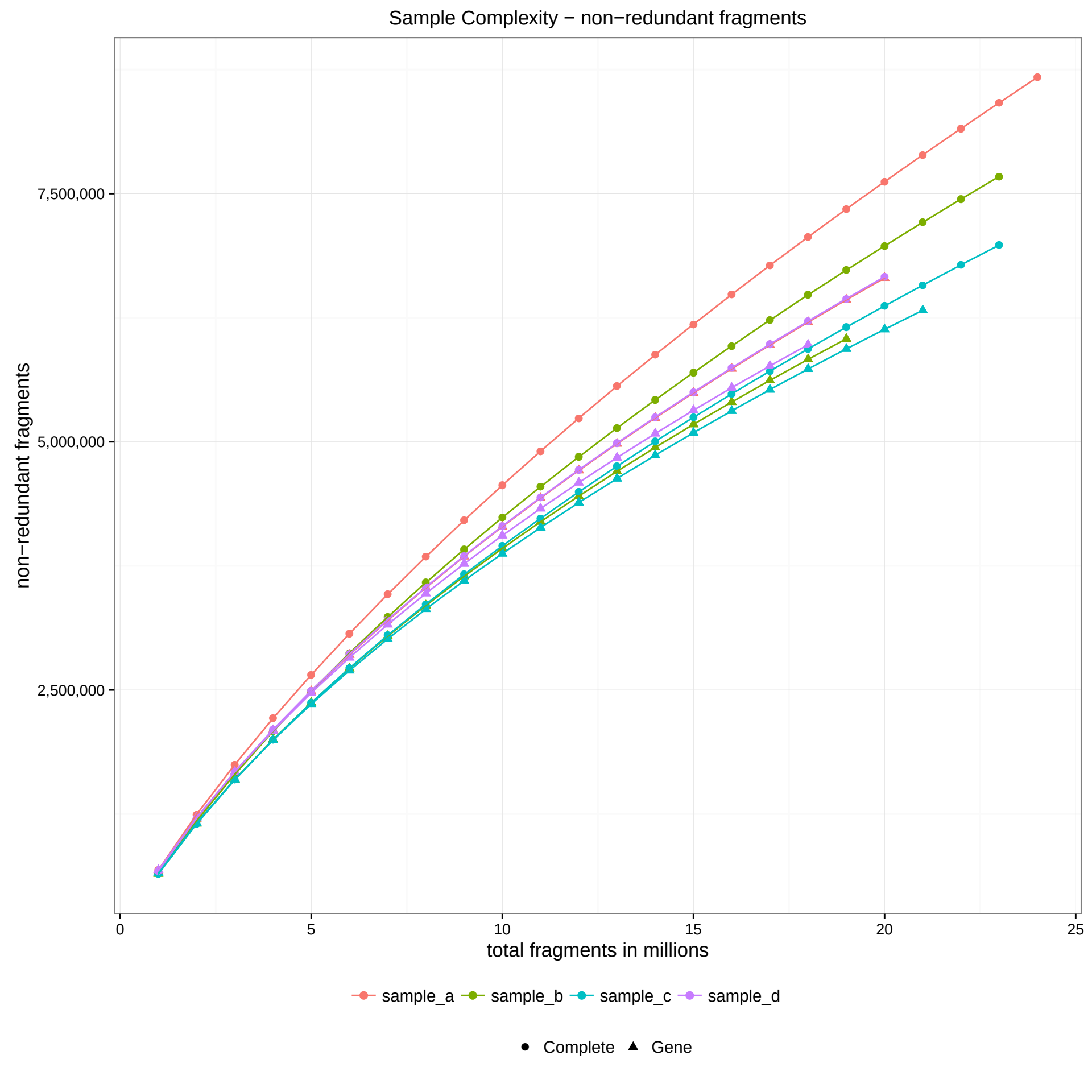

Sample Complexity − non−redundant fragments
non−redundant fragments
7,500,000
5,000,000
2,500,000
0
5
10
15
20
25
total fragments in millions
sample_a
sample_b
sample_c
sample_d
Complete
Gene