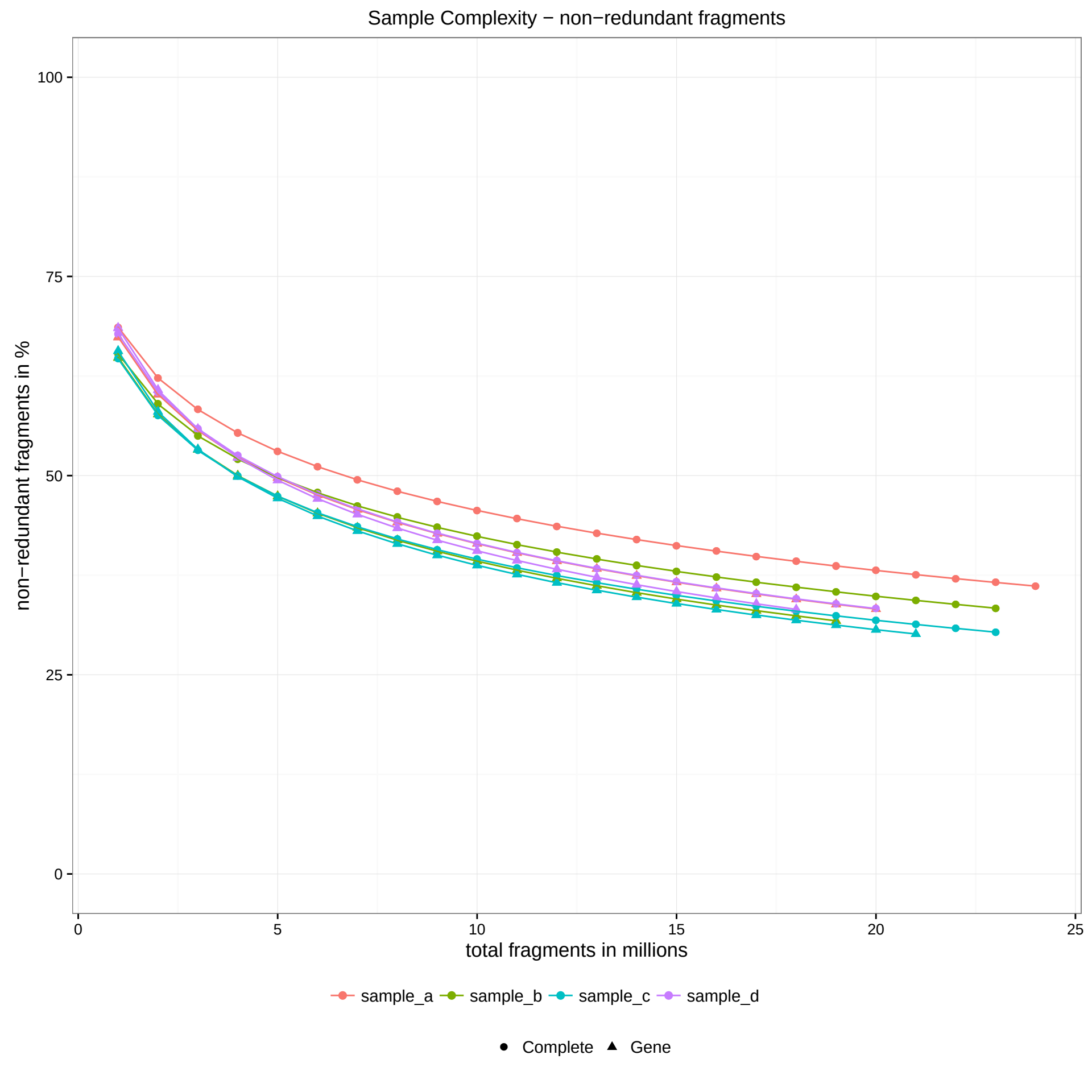

Sample Complexity − non−redundant fragments
100
75
50
25
0
non−redundant fragments in %
0
5
10
15
20
25
total fragments in millions
sample_a
sample_b
sample_c
sample_d
Complete
Gene