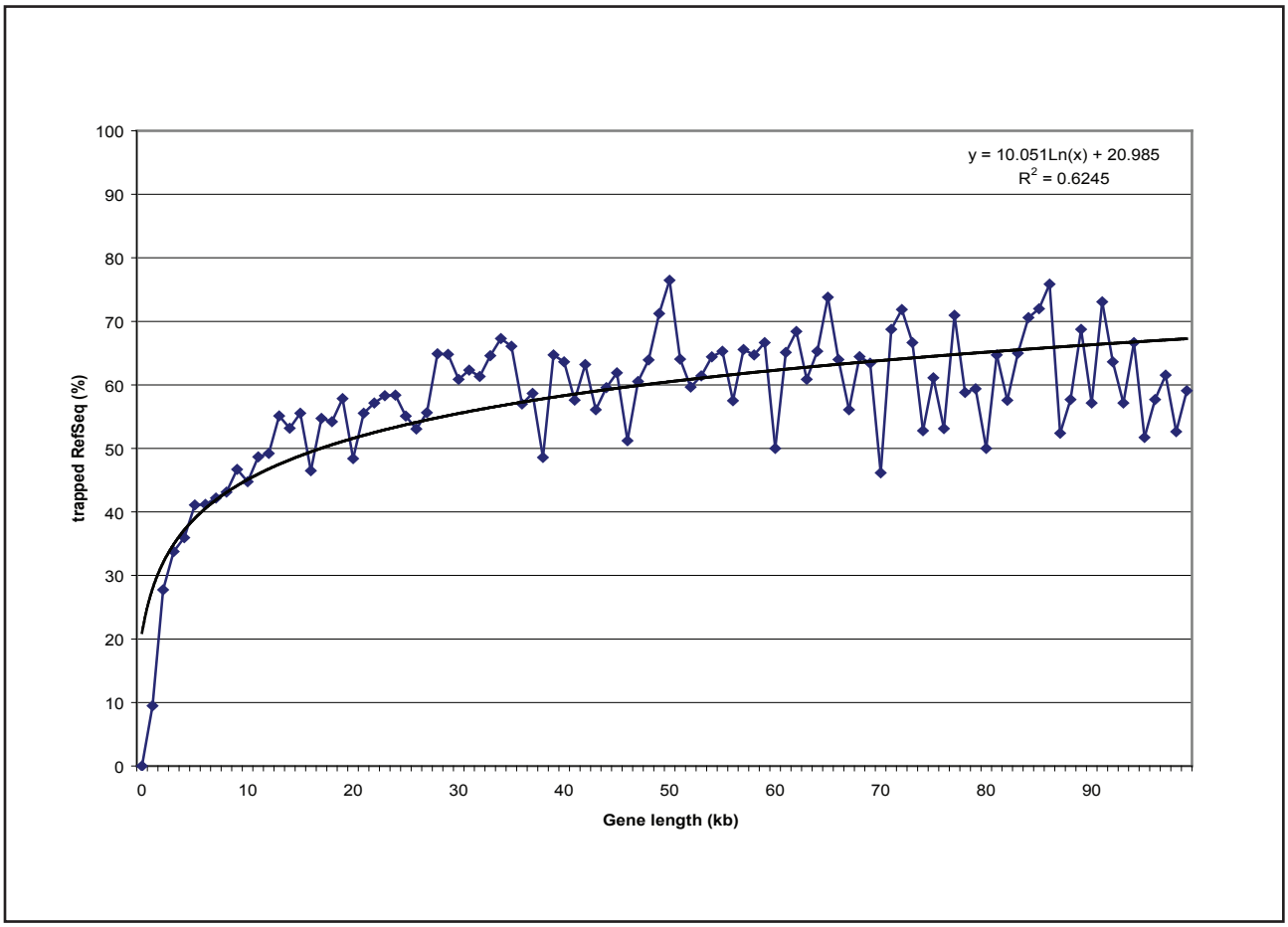
y = 10.051Ln(x) + 20.985
R2 = 0.6245
trapped RefSeq (%)
Gene length (kb)
0
10
20
30
40
50
60
70
80
90
100