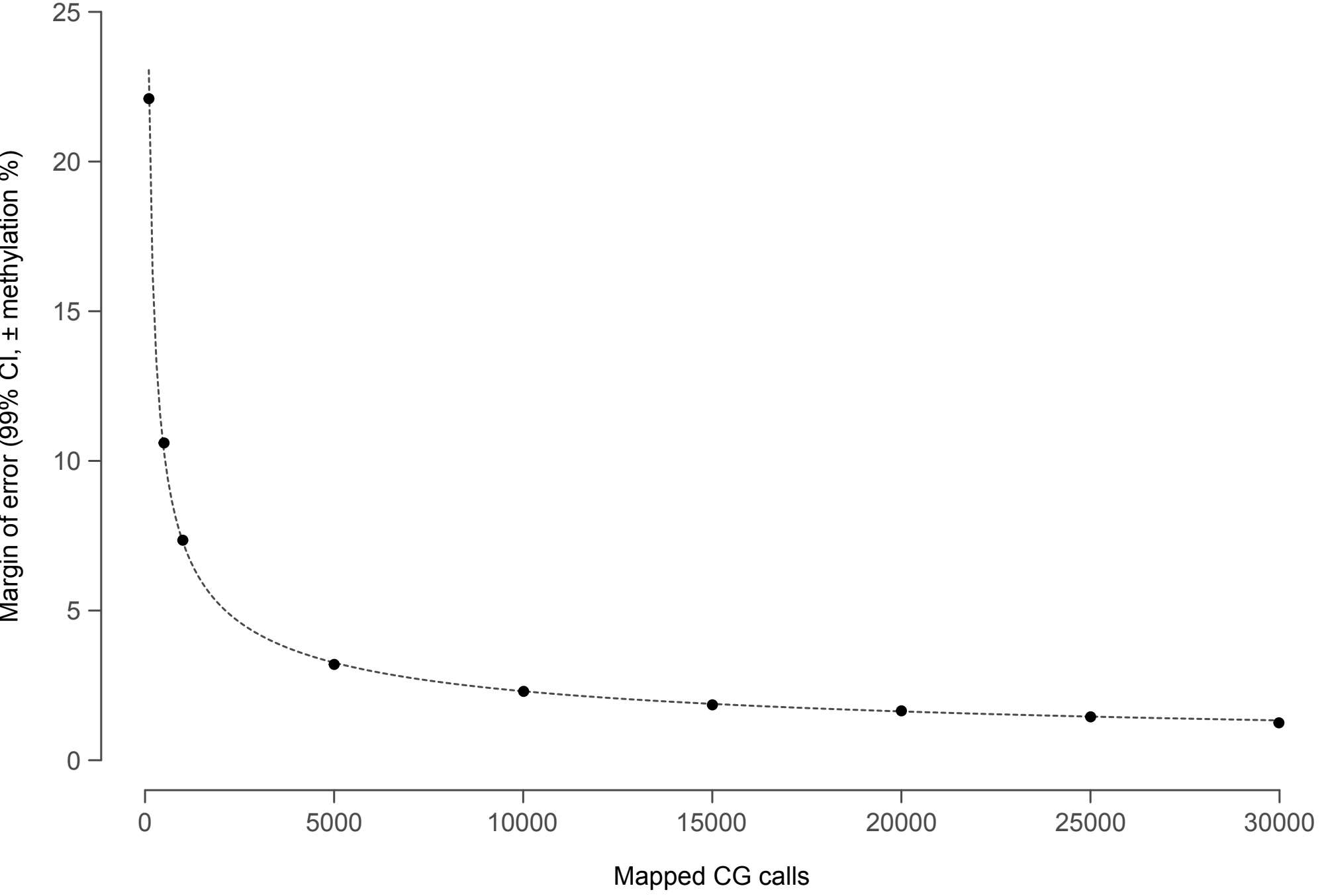
Margin of error (99% CI, ± methylation %)
25
20
15
10
5
0
0
5000
10000
15000
20000
25000
30000
Mapped CG calls