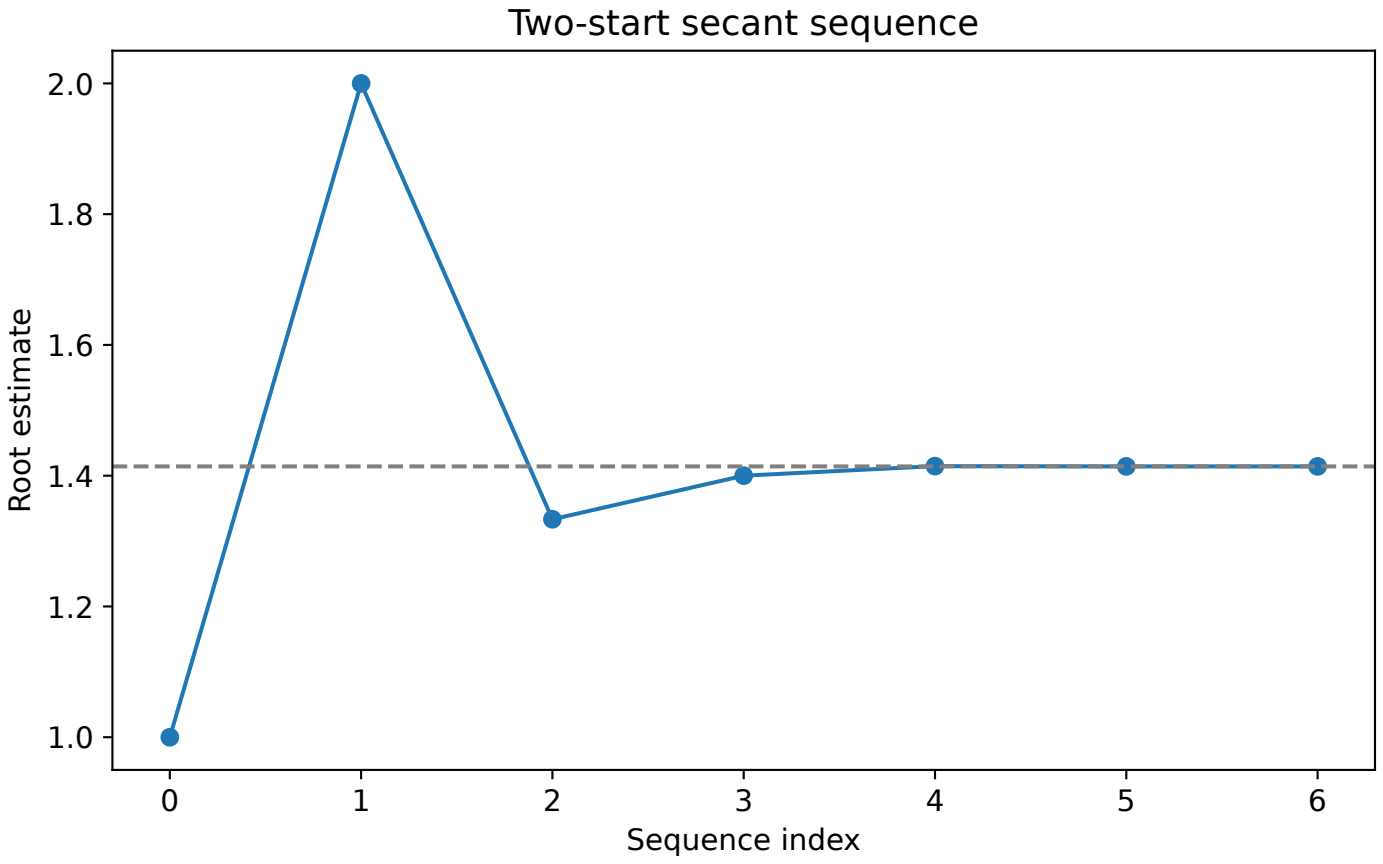
Two-start secant sequence
Root estimate
Sequence index
2.0
1.8
1.6
1.4
1.2
1.0
0
1
2
3
4
5
6