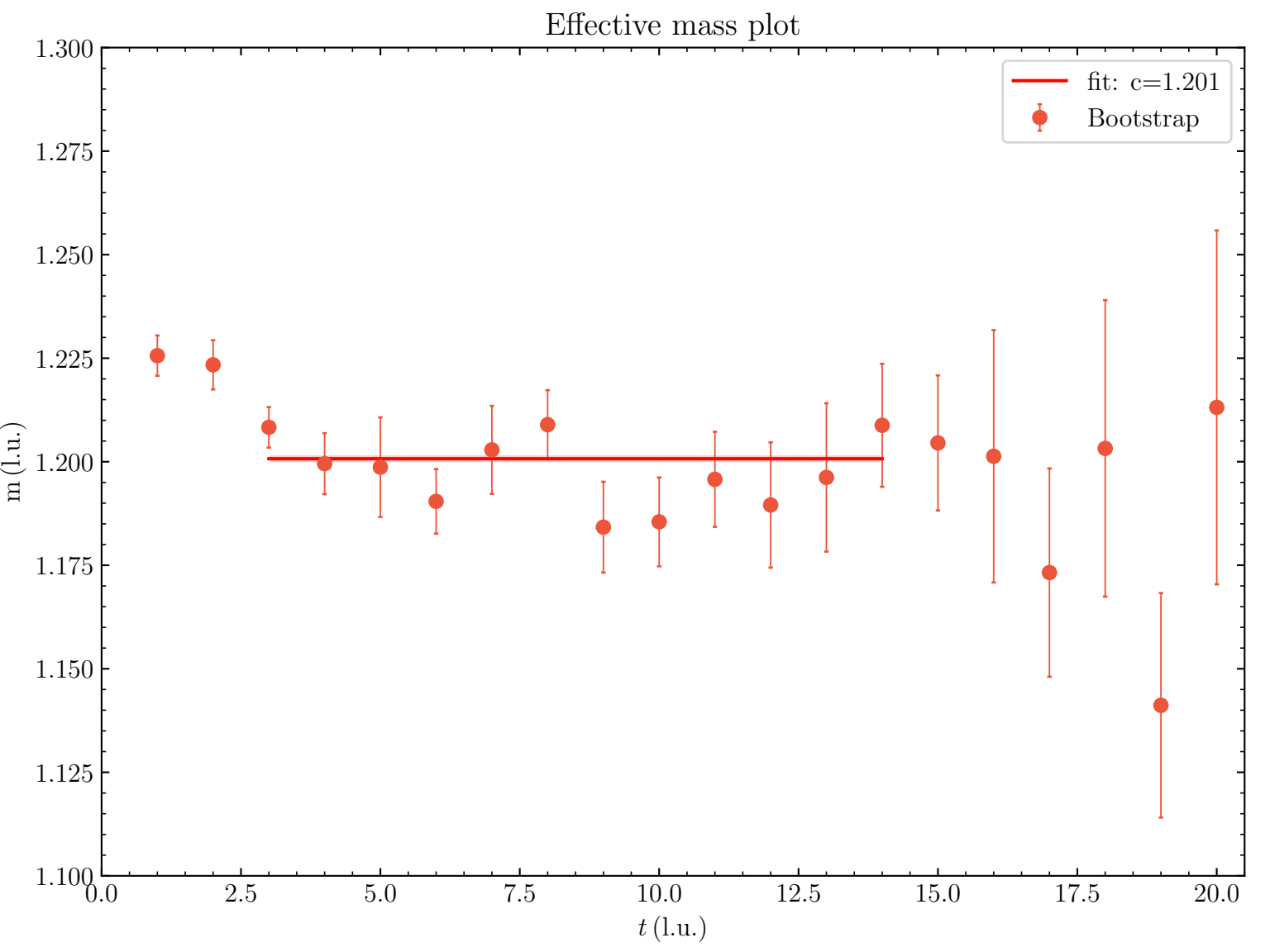
Effective mass plot
fit: c=1.201
Bootstrap
m (l.u.)
t (l.u.)
1.300
1.275
1.250
1.225
1.200
1.175
1.150
1.125
1.100
0.0
2.5
5.0
7.5
10.0
12.5
15.0
17.5
20.0