

1 **Title:** Assessment of bias and precision among simple closed population mark-
2 recapture estimators

3

4 **Author:** Kyle Dettloff¹

5

6 ¹National Oceanic and Atmospheric Administration, National Marine Fisheries Service,
7 Southeast Fisheries Science Center, 75 Virginia Beach Drive, Miami, FL 33149, USA

8 Email: kyle.dettloff@noaa.gov

9

10 **Keywords:** bias correction; closed population; mark-recapture; Monte Carlo simulation;
11 Schumacher-Eschmeyer

12 Abstract

13

14 Mark-recapture methods have been heavily studied and employed in fisheries and other
15 wildlife sciences over the past century to approximate population sizes for animal species of
16 interest. This paper focuses on the comparative statistical performance through simulation of
17 common closed population mark-recapture estimators, including those of Lincoln-Petersen,
18 Chapman, Chao, Schnabel, and Schumacher-Eschmeyer. A new bias-adjusted version of the
19 Schumacher-Eschmeyer estimator is proposed and is shown to exhibit superior performance at
20 small sample sizes in comparison to the original estimator. Simulation results indicate that
21 Chapman's method outperforms all other two-visit methods and that bias-adjusted versions of
22 Schnabel and Schumacher-Eschmeyer differ slightly depending on bias or precision, but both
23 perform well. Minimum sample sizes such that resulting estimates are approximately unbiased
24 are proposed to advise practitioners on the most appropriate use of these estimators for simple
25 closed population mark-recapture data.

26

27 1. Introduction

28 Reliable population size estimates are fundamental to understanding the ecology and
29 conservation needs of animal populations. Procedures to derive these estimates differ
30 fundamentally from that of traditional statistical survey methods in that there is no predetermined
31 frame of individuals from which an investigator can sample. The first solution to this problem in
32 the ecological literature dates back to 1896 when fishery scientist G.C.J. Petersen invented a
33 brass tag to attach to fish, recognizing that population size could be estimated when sufficient
34 numbers of these tags had been recaptured. The same method was later derived by ornithologist
35 Frederick Lincoln (1930) to estimate the size of North American waterfowl populations using
36 banding data. These approaches are collectively known as the Lincoln-Petersen method, the
37 simplest and most well-known metric to estimate the size of a closed population, meaning a
38 negligible effect of births, deaths, and movement during the study period. Animals are captured,
39 marked, and released on the first visit, and, after being allowed to mix with the full population,
40 randomly sampled without replacement on a second visit, noting the number of marked
41 individuals present from the first sample in the second sample. This two-visit method was
42 subsequently expanded to apply to k visits by Schnabel (1938) and further modified by
43 Schumacher and Eschmeyer (1943) to be formulated as a linear regression. More recently,
44 Brittain and Böhning (2009) used empirical data to compare the performance of the Chao (1987,
45 1989) estimator, which relaxes the assumption of independence between visits, with the more
46 common Chapman (1951) estimator, a bias-adjusted version of the two-visit Lincoln-Petersen.

47 Though more sophisticated model-based methods have been proposed to relax the
48 assumptions of independence between captures (Otis et al. 1978; Huggins 1989, 1991) and
49 closed populations (Seber 1982), this paper focuses strictly on the theoretical performance of
50 simple closed population estimators for which identification of individuals is not required. While
51 sampling complexities such as variation in capture probabilities and uncertain sampling area are
52 known to impact bias and variance in mark-recapture studies (Amstrup et al. 2005, McNair et al.
53 2018), this study aims to assess the validity of these existing metrics under conditions in which
54 all assumptions are known to be met, allowing one to select a superior metric from a purely
55 statistical basis, all other sampling considerations equal. For a study evaluating the performance

56 of various mark-recapture models on field data with known reference population sizes, including
57 those allowing individual heterogeneity in capture probabilities, see Grimm et al. (2014).

58 These simple estimators remain relevant within fisheries and conservation biology,
59 returning over 1,650 articles since 2016 with reference to closed population mark-recapture in a
60 Google Scholar search. Of these, 69 contain specific reference to the Schumacher-Eschmeyer
61 estimator, which has been employed to estimate population size in a wide variety of fisheries,
62 including endangered Atlantic sturgeon (Kahn et al. 2014, 2019; Hale et al. 2016), lake trout
63 (Hansen et al. 2008), walleye (Spencer et al. 2002), American lobster (Rowe 2002), pirarucu
64 (Castello 2004), and even for estimating angler counts in a creel survey (Hansen and Van Kirk
65 2018).

66 Comparing these historically important and still common estimators within a consistent
67 framework and across a range of simulations offers clarity as to which methods are most
68 appropriate under known sampling scenarios, with the equal benefit of indicating which
69 estimators should be confidently discontinued from use in favor of superior alternatives.
70 Consequently, these simulations are also able to suggest approximate minimum sample sizes
71 needed to generate reliable estimates of population size among the best performing methods.

72
73

74 **2. Materials and methods**

75

76 A collection of common closed population mark-recapture estimators prevalent in the
77 ecological literature were selected and compared in performance through simulation.
78 Comparisons were conducted for both two-visit methods and multi-visit methods. All methods
79 analyzed assume:

- 80 1. Closed populations (no change in population size between sampling events)
- 81 2. Independence between visits (marking does not influence the probability of recapture)
- 82 3. Independence between individuals (complete mixing occurs between sampling events)
- 83 4. Individuals are sampled without replacement
- 84 5. No marks are lost between sampling events

85 A brief overview of the various estimators is provided followed by the simulation study
86 methodology.

87

88

89 *2.1 Overview of estimators*

90

91 *2.1.1 Lincoln-Petersen estimator (2 visits)*

92 Also known as the Lincoln Index, Lincoln-Petersen (Petersen 1896, Lincoln 1930) is the
93 simplest and most intuitive of the estimators, and is the maximum likelihood estimator (MLE) of
94 N . It is given by (Eq. 1),

95

$$96 \hat{N} = \frac{Mn}{m} \tag{1}$$

97

98 where:

99 M = number of animals marked on the first visit;

100 n = total number of animals captured on the second visit;

101 m = number of marked animals recaptured on the second visit.

102 This equation implies that the proportion of marked individuals captured in the second sample
103 (m/n) is equal to the proportion of the total population (N) that has been marked in the first
104 sample (M/N). Lincoln-Petersen forms the basis for all estimators that follow.

105 106 2.1.2 Chapman estimator (2 visits)

107
108 At small sample sizes, (Eq. 1) produces biased estimates of population size (Chapman 1951).
109 Several modifications have been suggested to reduce this bias, the most common being the
110 Chapman (1951) estimator, given by (Eq. 2).

$$111 \hat{N} = \frac{(M+1)(n+1)}{(m+1)} - 1 \quad (2)$$

112
113 This estimator, based on the hypergeometric distribution, possesses finite moments, as the
114 denominator cannot be zero, which is possible in Lincoln-Petersen when $M + n < N$. A method to
115 obtain robust confidence intervals around Chapman estimates of population size is provided by
116 Sardinle (2009).
117

118 119 2.1.3 Chao estimator (2 visits)

120
121 Chao (1987, 1989) proposed an estimator that relaxes the assumption of independence in capture
122 probability between visits. Brittain and Böhning (2009) show that for the two-visit, equal capture
123 probability scenario, this estimator can be formulated as (Eq. 3).

$$124 \hat{N} = \frac{(M+n-2m)^2}{4m} + M + n - m \quad (3)$$

125
126 In the case of $M = n$, it can be seen that (Eq. 3) reduces to the Lincoln-Petersen estimator (Eq. 1).

127 128 129 2.1.4 Bayesian estimator (2 visits)

130
131 The final two-visit estimator considered is a Bayesian formulation which estimates the posterior
132 mean based on the hypergeometric distribution, analogous to the Chapman estimator. The
133 derivation is presented in Webster and Kemp (2013), and results in (Eq. 4).

$$134 \hat{N} = \frac{(M-1)(n-1)}{(m-2)} \text{ for } m > 2 \quad (4)$$

135 136 137 2.1.5 Schnabel estimator (≥ 2 visits)

138
139 Schnabel (1938) published the first mark-recapture estimator designed for more than two
140 sampling visits, generalizing the traditional two-visit Lincoln-Petersen approach. The equation is
141 formulated as a weighted average of Lincoln-Petersen estimates across the series of visits (Eq.
142 5),

$$143 \hat{N} = \frac{\sum_k M_k n_k}{\sum_k m_k} \text{ for } k \geq 2 \quad (5)$$

144
145

146 where:

147 M_k = total number of marked animals in the population prior to visit k ;

148 n_k = total number of animals captured on visit k ;

149 m_k = total number of marked animals recaptured on visit k .

150

151 All individuals captured on each visit are marked and released into the population, with no need
152 to distinguish between marks made on different visits. Note that the estimator becomes
153 equivalent to Lincoln-Petersen (Eq. 1) in the case of $k = 2$ visits.

154 An improved small sample bias correction given in (Eq. 6) was proposed by Chapman
155 (1952), noting that each m_k is approximately Poisson distributed with parameter $M_k n_k / N$. This
156 correction has been recommended by multiple sources, and its performance is evaluated here
157 alongside the original estimator.

158

159
$$\hat{N} = \frac{\sum_k M_k n_k}{\sum_k m_k + 1} \quad (6)$$

160

161 2.1.6 Schumacher-Eschmeyer estimator (≥ 2 visits)

162

163 A similar estimator which employs the same sampling methodology as Schnabel to handle
164 multiple recapture events was proposed by Schumacher and Eschmeyer (1943), taking the form
165 of (Eq. 7).

166

167
$$\hat{N} = \frac{\sum_k M_k^2 n_k}{\sum_k M_k m_k} \text{ for } k \geq 2 \quad (7)$$

168

169 The logic behind this formula is that the proportion of marked individuals on the k^{th} visit (m_k/n_k)
170 plotted against the number of individuals previously marked (M_k) should be linear and pass
171 through the origin with a slope of N^{-1} under the basic assumptions outlined above. (Eq. 7) uses
172 linear regression techniques to estimate N based on this rationale.

173 Following the bias-adjusted formulation of the Schnabel estimator in (Eq. 6), a similar
174 small sample bias correction to the Schumacher-Eschmeyer equation is proposed here as (Eq. 8)
175 based on the Chapman correction (Eq. 2).

176

177
$$\hat{N} = \frac{\sum_{k=2}^k (M_k + 1)^2 (n_k + 1)}{\sum_k M_k (m_k + 1)} - 2 \quad (8)$$

178

179 The performance of this estimator is evaluated alongside the original through simulation in the
180 present study.

181

182

183 *2.2 Simulation study*

184

185 A series of Monte-Carlo simulations were run using the R programming language
186 (v4.2.0; R Core Team, 2022) to evaluate the behavior of each estimator outlined above. Code is
187 provided in Supporting Information for readers to run these simulations under their own selected
188 parameters.

189

190 2.2.1 Estimator performance

191
192 Fixed populations of known size N were generated and randomly sampled without replacement
193 using the *sample* function, according to the capture and marking methodology for each
194 procedure. For each estimator, this was repeated 10,000 times for all sample sizes M and n
195 increasing from very small to large for both cases of $M = n_{2\dots k}$ and $M \neq n_{2\dots k}$. Estimates \hat{N} and
196 $SE(\hat{N})$ were calculated by taking the sample mean and standard deviation, respectively, from all
197 simulations producing finite values. The resulting estimates were plotted against the geometric
198 means of the marked individuals and the total numbers of captured individuals during subsequent
199 visits to evaluate how each estimator's bias and variance changes with increasing sample size.
200 Likewise, the scaled root-mean-square error (RMSE) (9) of each estimator was plotted over the
201 range of sample sizes to visualize performance in terms of a metric that combines the effect of
202 bias and variance.

$$204 \quad RMSE_{scaled} = \frac{\sqrt{var(\hat{N}) + bias(\hat{N})^2}}{N} \quad (9)$$

206 2.2.2 Minimum sample sizes

207
208 The Chapman estimator has been shown to be exactly unbiased when the sum of the sample sizes
209 is at least as large as the population size, or $M + n \geq N$ (Robson and Regier, 1964; Wittes 1972).
210 A less stringent condition for the estimator to be approximately unbiased, with negative bias less
211 than 2%, was noted by Robson and Regier (1964) in cases when:

$$213 \quad \sqrt{Mn} \geq 2\sqrt{N} \quad (10)$$

214
215 That is, when the geometric mean of the marks and captures is at least twice the square root of
216 the population size. The derivation of the degree of bias in the estimator leading to this
217 approximate threshold is provided in Chapman (1951) using Stirling's formula with the
218 hypergeometric distribution. Based on the results of 2.2.1, the bias-corrected version of Schnabel
219 (Eq. 6) was analyzed across various simulation scenarios, suggesting a generalization of (Eq. 10)
220 produces a similarly acceptable rule of thumb for the minimum sample size needed relative to
221 the population size for resulting estimates to be approximately unbiased (Eq. 11).

$$223 \quad \sqrt[k]{M \prod_{i=2}^k n_i} \geq \frac{2\sqrt{N}}{k-1} \quad (11)$$

224
225 Percent error, calculated according to (Eq. 12), was used to evaluate stability in bias at the
226 proposed approximate minimum sample sizes, based on 100,000 simulations with known
227 population sizes N ranging from 10^2 to 10^6 .

$$229 \quad Pct. Error = 100 \left(\frac{\hat{N} - N}{N} \right) \quad (12)$$

230
231 Similar exploration was conducted for the bias-adjusted Schumacher-Eschmeyer (Eq. 8).

232
233

234 **3. Results**

235

236 *3.1 Estimator performance*

237

238 While simulations were conducted for a wide range of population sizes, results below are only
239 presented for the case of $N = 1,000$, as similar patterns held across all sizes evaluated.

240

241 *3.1.1 Two-visit methods*

242

243 Fig. 1 displays simulation results for the Lincoln-Petersen, Chapman, Chao, and Bayesian
244 estimators in the case of $M < n$, or fewer individuals marked than recaptured. All four are biased
245 low at very small sample sizes, which can be easily confirmed using simple arithmetic on
246 hypothetically small M and n . This is due to the fact that either there will be no recaptures and
247 thus no valid estimate, or if there is an unlikely recapture, the resulting estimate will be a gross
248 underestimate of population size. While it is already well known that Chapman is an
249 improvement over Lincoln-Petersen in terms of bias, it was seen clearly here that Lincoln-
250 Petersen becomes biased high for an intermediate range of sample sizes after exhibiting this
251 known low bias at very small sample sizes. Lincoln-Petersen also exhibits a higher variance than
252 Chapman even at sample sizes where Chapman is already approximately unbiased. Likewise, the
253 Chao and Bayesian estimators all eventually become biased high before approaching
254 approximately unbiased states at larger sample sizes. The Chao estimator retains a high bias for
255 much longer than any of the others in this case, while also being more variable than Lincoln-
256 Petersen. The Bayesian estimator clearly has higher bias and variance at small sample sizes than
257 Lincoln-Petersen or Chao. In contrast, the Chapman estimator never becomes biased high, and
258 approaches the value of the true population size much faster than the other three estimators while
259 also exhibiting lower variance. Similar outcomes were observed with $n < M$ and $n = M$, so
260 simulation results are only presented for $M < n$ to illustrate a scenario where the Lincoln-
261 Petersen and Chao estimates differ.

262 Fig. 2 demonstrates how the RMSE for each method changes with increasing sample size
263 plotted on the x-axis as the geometric mean of M and n for a hypothetical population of size $N =$
264 $1,000$. The Bayesian estimator has the highest RMSE at low sample sizes, eventually
265 approaching that of the other methods at larger sample sizes, while the Chao estimator retains a
266 much higher RMSE even as the sample size increases. It is clear again that the Chapman
267 estimator exhibits superior performance, having a much lower RMSE than the others at sample
268 sizes even above the point at which the extreme downward-bias present in all methods begins to
269 disappear (represented by the dashed vertical line in Fig. 2).

270

271 *3.1.2 Multi-visit methods*

272

273 Simulation results are presented for both the original and bias-adjusted variations of Schnabel
274 and Schumacher-Eschmeyer with $M = n_2 = n_3$ (Fig. 3). For brevity, results are only displayed for
275 $k = 3$ visits, as similar scaled patterns were observed under increased k . As with the two-visit
276 methods, all estimates are biased low at very small sample sizes with negative bias decreasing as
277 sample size increases. Similar to Lincoln-Petersen, both of the original unadjusted estimators
278 eventually become biased high before turning approximately unbiased at larger sample sizes.
279 Between the original versions, the unadjusted Schnabel estimator possesses noticeably lower

280 variability and slightly less small sample bias than the unadjusted Schumacher-Eschmeyer (Figs.
281 3 and 4).

282 Performance was very similar between the adjusted versions, however, offering
283 improvements in both bias and variability. In each, bias approaches zero as the sample size
284 increases without ever becoming positive. At small sample sizes, the adjusted Schnabel becomes
285 unbiased slightly faster, while the adjusted Schumacher-Eschmeyer is less variable, resulting in a
286 lower RMSE for the adjusted Schumacher-Eschmeyer within a small range of sample sizes
287 before the two become effectively equivalent at larger sample sizes (Fig. 4). While of limited
288 practical impact, simulation results revealed that taking the ceiling of the adjusted Schnabel
289 estimator provides a more appropriate estimate of population size, with slightly reduced bias
290 across all sample sizes. The adjusted Schumacher-Eschmeyer estimator presented in (Eq. 8)
291 exhibited the fastest reduction in bias at small sample sizes while remaining exactly unbiased at
292 large sample sizes among a variety of alternate formulations considered. That is, it was seen to
293 satisfy that:

294

$$295 E_N[\hat{N}] \rightarrow N \text{ as } \sqrt[k]{M \prod_{i=2}^k n_i} \rightarrow N \quad (13)$$

296

297 All patterns were essentially the same in the case of $M > n_2 = n_3$, and the only apparent
298 difference when $M < n_2 = n_3$ is even higher variability in the unadjusted Schumacher-Eschmeyer
299 estimator at small sample sizes. Notice that the slightly lower RMSE of the adjusted
300 Schumacher-Eschmeyer is driven by lower variability at small sample sizes, even though the
301 absolute bias of the adjusted Schnabel decreases slightly faster (Fig. 5).

302 3.2 Minimum sample size

303

304 Simulations confirmed the minimum sample sizes (Eq. 10) suggested by Robson and Regier
305 (1964) for the Chapman estimator to produce approximately unbiased estimates. This threshold
306 is represented by a dashed vertical line in Fig. 2. Relative bias (Eq. 11) at these minimum
307 suggested sample sizes was less than 2% for cases $M = n$, $M < n$, and $M > n$ across all population
308 sizes N from 10^2 to 10^6 .

309 Likewise, the approximate minimum sample size rule (Eq. 11) proposed for the bias-
310 adjusted Schnabel estimator, following the logic in (Eq. 10) and confirmed based on examination
311 of simulation results, reveals that the geometric mean of the sample sizes across all visits should
312 be at least twice the square root of the population size divided by one less the number of visits.
313 Relative bias at this sample size was observed to generally range between approximately -10%
314 and -3% for population sizes N ranging from 10^2 to 10^6 , respectively, increasing slightly with the
315 number of visits (Table 1). A corresponding simple rule of thumb to achieve a consistent bias
316 threshold relative to population size was not readily apparent for the bias-corrected Schumacher-
317 Eschmeyer estimator.

318 4. Discussion

319

320 The methods described in this paper have been well studied over the decades since they
321 have been published and are still widely encountered today, including in introductory ecology
322 textbooks (Krebs 1999) and statistical programming packages (Nelson, 2023, >100 K total
323
324

325 downloads; Ogle et al., 2023, >550 K total downloads). The comparison of these multiple
326 common methods side by side under known conditions provides insight into the exact behavior
327 of the estimators across a complete range of sample sizes.

328 It is clear that among the two-visit methods, the Chapman estimator is superior in terms
329 of both bias and variance. The estimator of Bailey (1951, 1952), originally proposed for
330 sampling with replacement and not presented here due to its similarity to Chapman, behaves
331 much the same but retains a slightly larger downward bias across all sample sizes, which can be
332 inferred from a simple examination of the equation. The Chao and Bayesian estimators can
333 easily be eliminated as viable options for data of this type in that the Chao estimator possesses a
334 large upward bias when $M \neq n$ and the Bayesian estimator possesses much higher bias and
335 variance at small sample sizes than the other methods. The unidirectional bias of the Chapman
336 estimator is appealing, but it should be noted that a strong downward bias is still present at very
337 small sample sizes, and therefore it is not recommended to be used when sample sizes are
338 smaller than that approximated by (Eq. 10). This recommendation is paradoxical in that the
339 approximate minimum sample size depends on an estimate of the population size, which one is
340 aiming to estimate to begin with. Therefore, it should be seen as a way of avoiding inaccurate
341 estimates from absurdly small sample sizes based on an educated guess of the order of
342 magnitude of the size of the population being sampled.

343 Likewise among multi-visit methods, the bias-adjusted estimators of Schnabel and
344 Schumacher-Eschmeyer clearly exhibited superior performance at small sample sizes, with no
345 reduction in performance at large sample sizes. Krebs (1999) suggested thresholds for when the
346 small sample adjusted Schnabel should be used, however based on the present simulation results
347 it is recommended that the adjusted estimators are used in place of the originals in all scenarios.
348 The bias-adjusted form of the Schumacher-Eschmeyer estimator presented here can be especially
349 impactful on population size estimates of species with low sample sizes and numbers of
350 recaptures (e.g., only two recaptured Atlantic sturgeon among 17 marked in a study by Kahn et
351 al. (2014)). Application can also extend to calculating more robust closed population estimates
352 for threatened species either directly or indirectly impacted by fishing pressure, such as Indo-
353 Pacific humpback dolphin (Zhou et al. 2007), or juvenile marine turtle species observed to
354 exhibit high fidelity to distinctive foraging areas (Wildermann et al. 2019).

355 Deviation from assumptions in field scenarios can lead to additional limitations of these
356 estimators to accurately estimate population size that extend beyond small sample bias. First and
357 foremost, if a population is assumed to be closed but in fact immigration and emigration are
358 occurring, population size may be overestimated since there are fewer marked individuals
359 present to be captured at any given time. Conversely, large numbers of individuals with low
360 capture probabilities within a closed population are assumed to lead to underestimation of
361 population size among all estimators evaluated (Amstrup et al. 2005). However, Grimm et al.
362 (2014) observed that a multiple Lincoln-Petersen estimator (with Chapman correction), similar to
363 the multi-visit methods evaluated here but with data from all subsequent visits pooled, performed
364 well on field data even with heterogeneous capture probabilities among individuals. This led
365 them to conclude it is a viable estimator of minimum population size provided a sufficient
366 number of individuals with high catchability are sampled, as capture probability increases and
367 heterogeneity decreases over multiple capture periods. Additionally, Seber (1982) noted the
368 regression-based Schumacher-Eschmeyer is expected to be the most robust multi-visit method
369 regarding violations of assumptions at the expense of loss in efficiency, which was observed in
370 the present study in terms of higher small sample variability when assumptions were exactly met.

371 This lends support to favoring the adjusted Schumacher-Eschmeyer both in terms of theoretical
372 performance and robustness to violations of assumptions.

373 A generalization of (Eq. 10), (Eq. 11) can provide a rough approximation of the
374 minimum sample size necessary to avoid a strong downward theoretical bias in estimates from
375 the adjusted multi-visit methods. Note, however, that the downward bias at this precise threshold
376 will be slightly larger for the adjusted Schnabel (Table 1) than for the two-visit Chapman
377 estimator with the trade-off of decreased variability at equal total sample sizes, and larger still for
378 the adjusted Schumacher-Eschmeyer (Fig. 5). This may be something to consider when selecting
379 an estimator depending on the relative risks associated with over vs. under-estimating a given
380 population size, noting that the approximate sample size provided by (Eq. 11) should be seen
381 even more as a lower bound when using the adjusted Schumacher-Eschmeyer.

382 Otis et al. (1978) and Evans, Kim, and O'Brien (1996) outline the problems with using
383 traditional normal approximation based methods to construct confidence intervals for mark-
384 recapture estimates, especially at small sample sizes, which fail to capture asymmetry around
385 estimates and are known to result in coverage below nominal levels. Buckland and Garthwaite
386 (1991) give general approaches to obtain confidence intervals for recapture data using parametric
387 bootstrapping techniques, which are shown to provide more robust estimates. Tyers (2021)
388 implements a bootstrap method shown to be robust for two-visit data, in which data m for the
389 second sampling event are resampled with replacement using a binomial distribution with size
390 parameter n and probability parameter m/n . These or similar bootstrap techniques can be
391 extended to the multi-visit methods outlined here, which also have the advantage of being valid
392 for the exact form of the estimator used.

393 Other sources have provided rules of thumb as to the number of recaptures needed to
394 achieve approximately unbiased estimates for closed population mark-recapture methods (Bailey
395 1951, Ricker 1975, Krebs 1999), which may simplify sampling in certain scenarios. However, all
396 results presented here treat the sample sizes as fixed and the number of recaptures as random, in
397 accordance with the procedures evaluated. It should also be noted that the present evaluation of
398 performance relates to the intrinsic theoretical bias of the estimators when assumptions are
399 perfectly met, and should not be extended to sources of bias or variability arising from sampling
400 methodology.

401

402 **Acknowledgements**

403 I would like to thank five anonymous reviewers for constructive suggestions to improve this
404 manuscript.

405 **Funding**

406

407 This research did not receive any specific grant from funding agencies in the public, commercial,
408 or not-for-profit sectors.

409

410 **References**

411 Amstrup, S.C., McDonald, T.L., Manly, B.F.J., 2005. Handbook of capture-recapture analyses.
412 Princeton University Press. Princeton, New Jersey, USA.

413
414 Bailey, N.T., 1951. On estimating the size of mobile populations from recapture data. *Biometrika*
415 38(3-4), 293-306. <https://doi.org/10.2307/2332575>
416
417 Bailey, N.T., 1952. Improvements in the interpretation of recapture data. *J. Anim. Ecol.* 21(1),
418 120-127. <https://doi.org/10.2307/1913>
419
420 Brittain, S., Böhning, D., 2009. Estimators in capture-recapture studies with two sources. *Adv.*
421 *Stat. Anal.* 93(1), 23-47.
422
423 Buckland, S.T., Garthwaite, P.H., 1991. Quantifying precision of mark-recapture estimates using
424 the bootstrap and related methods. *Biometrics* 47(1), 255-268.
425 <https://doi.org/10.2307/2532510>
426
427 Castello, L. 2004. A method to count pirarucu *Arapaima gigas*: fishers, assessment, and
428 management. *N. Am. J. Fish. Manag.* 24(2), 379-389. <https://doi.org/10.1577/M02-024.1>
429
430 Chao, A., 1987. Estimating the population size for capture-recapture data with unequal
431 catchability. *Biometrics* 43(4), 783-791. <https://doi.org/10.2307/2531532>
432
433 Chao, A., 1989. Estimating population size for sparse data in capture-recapture experiments.
434 *Biometrics* 45(2), 427-438. <https://doi.org/10.2307/2531487>
435
436 Chapman, D.G., 1951. Some properties of the hypergeometric distribution with applications to
437 zoological sample censuses. *Univ. Calif. Publ., Stat.* 1(7), 131-160.
438
439 Chapman, D.G., 1952. Inverse, multiple and sequential sample censuses. *Biometrics* 8(4), 286-
440 306. <https://doi.org/10.2307/3001864>
441
442 Evans, M.A., Kim, H.M., O'Brien, T.E, 1996. An application of profile-likelihood based
443 confidence interval to capture: recapture estimators. *J. Agric. Biol. Environ. Stat.* 1(1),
444 131-140. <https://doi.org/10.2307/1400565>
445
446 Grimm, A., Gruber, B., Henle, K., 2014. Reliability of different mark-recapture methods for
447 population size estimation tested against reference population sizes constructed from field
448 data. *PLoS One* 9(6), e98840. <https://doi.org/10.1371/journal.pone.0098840>
449
450 Hale, E.A., Park, I.A., Fisher, M.T., Wong, R.A., Stangl, M.J., Clark, J.H, 2016. Abundance
451 estimate for and habitat use by early juvenile Atlantic sturgeon within the Delaware River
452 Estuary. *Trans. Am. Fish. Soc.* 145(6), 1193-1201.
453 <https://doi.org/10.1080/00028487.2016.1214177>
454
455 Hansen, J.M., Van Kirk, R.W., 2018. A mark-recapture-based approach for estimating angler
456 harvest. *N. Am. J. Fish. Manag.* 38(2), 400-410. <https://doi.org/10.1002/nafm.10038>
457

458 Hansen, M.J., Horner, N.J., Liter, M., Peterson, M.P., Maiolie, M.A., 2008. Dynamics of an
459 increasing lake trout population in Lake Pend Oreille, Idaho. *N. Am. J. Fish. Manag.*
460 28(4), 1160-1171. <https://doi.org/10.1577/M07-149.1>
461

462 Huggins, R.M., 1989. On the statistical analysis of capture-recapture experiments. *Biometrika*
463 76, 133-140. <https://doi.org/10.1093/biomet/76.1.133>
464

465 Huggins, R.M., 1991. Some practical aspects of a conditional likelihood approach to capture
466 experiments. *Biometrics* 47, 725-732. <https://doi.org/10.2307/2532158>
467

468 Kahn, J.E., Hager, C., Watterson, J.C., Russo, J., Moore, K., Hartman, K., 2014. Atlantic
469 Sturgeon annual spawning run estimate in the Pamunkey River, Virginia. *Trans. Am.*
470 *Fish. Soc.* 143(6), 1508-1514. <https://doi.org/10.1080/00028487.2014.945661>
471

472 Kahn, J.E., Hager, C., Watterson, J.C., Mathies, N., Hartman, K.J., 2019. Comparing abundance
473 estimates from closed population mark-recapture models of endangered adult Atlantic
474 sturgeon. *Endanger. Species Res.* 39, 63-76. <https://doi.org/10.3354/esr00957>
475

476 Krebs, C.J, 1999. *Ecological methodology*. 2nd ed. Harper & Row, New York, New York, USA.
477

478 Lincoln, F.C., 1930. Calculating waterfowl abundance on the basis of banding returns. U.S.
479 Department of Agriculture Circular 118, 1-4.
480

481 McNair, J.N., Ruetz III, C.R., Carlson, A., Suh, J, 2018. Reducing effects of dispersal on the bias
482 of 2-sample mark-recapture estimators of stream fish abundance. *PLoS ONE* 13(8),
483 e0200733. <https://doi.org/10.1371/journal.pone.0200733>
484

485 Nelson, G.A, 2023. *fishmethods: Fishery Science Methods and Models*. R package version 1.11-
486 3. URL <https://CRAN.R-project.org/package=fishmethods>
487

488 Ogle, D.H., Doll, J.C., Wheeler, P., Dinno, A., 2023. *FSA: Fisheries Stock Analysis*. R package
489 version 0.9.4. URL <https://github.com/fishR-Core-Team/FSA>
490

491 Otis, D.L., Burnham K.P., White, G.C., Anderson D.R, 1978. Statistical inference from capture
492 data on closed animal populations. *Wildl. Monogr.* 62, 1-135.
493

494 Petersen, C.G.J, 1896. The yearly immigration of young plaice into the Limfjord from the
495 German Sea. Report of the Danish Biological Station to the Board of Agriculture 6, 1-48.
496

497 R Core Team, 2022. *R: A language and environment for statistical computing*. R Foundation for
498 Statistical Computing, Vienna, Austria. URL <https://www.R-project.org/>
499

500 Ricker, W.E., 1975. Computation and interpretation of biological statistics of fish populations.
501 *Bull. Fish. Res. Board Can.* 191, 1-382.
502

503 Robson, D.S., Regier, H.A., 1964. Sample size in Petersen mark-recapture experiments. *Trans.*
504 *Am. Fish. Soc.* 93(3), 215-226. [https://doi.org/10.1577/1548-](https://doi.org/10.1577/1548-8659(1964)93[215:SSIPME]2.0.CO;2)
505 [8659\(1964\)93\[215:SSIPME\]2.0.CO;2](https://doi.org/10.1577/1548-8659(1964)93[215:SSIPME]2.0.CO;2)
506

507 Rowe, S., 2002. Population parameters of American lobster inside and outside no-take reserves
508 in Bonavista Bay, Newfoundland. *Fish. Res.* 56(2), 167-175.
509 [https://doi.org/10.1016/S0165-7836\(01\)00321-6](https://doi.org/10.1016/S0165-7836(01)00321-6)
510

511 Sadinle, M., 2009. Transformed logit confidence intervals for small populations in single
512 capture-recapture estimation. *Commun. Stat. Simul. Comput.* 38(9), 1909-1924.
513 <https://doi.org/10.1080/03610910903168595>
514

515 Schnabel, Z.E., 1938. The estimation of the total fish population of a lake. *Am. Math. Mon.*
516 45(6), 348-352. <https://doi.org/10.2307/2304025>
517

518 Schumacher, F.X., Eschmeyer, R.W., 1943. The estimation of fish populations in lakes or ponds.
519 *J. Tenn. Acad. Sci.* 18, 228-249.
520

521 Seber, G.A.F., 1982. The estimation of animal abundance and related parameters. 2nd ed.
522 Macmillan, New York, New York, USA.
523

524 Spencer, S.C., Colby, P.J., Momot, W.T., Fruetel, M. 2002. Response of a walleye population to
525 pulse fishing in Henderson Lake, Ontario. *N. Am. J. Fish. Manag.* 22(3), 842-851.
526 [https://doi.org/10.1577/1548-8675\(2002\)022<0842:ROAWPT>2.0.CO;2](https://doi.org/10.1577/1548-8675(2002)022<0842:ROAWPT>2.0.CO;2)
527

528 Tyers, M., 2021. recapr: Two event mark-recapture experiment. R package version 0.4.4. URL
529 <https://CRAN.R-project.org/package=recapr>
530

531 Webster, A.J., Kemp, R. 2013. Estimating omissions from searches. *Am. Stat.* 67(2), 82-89.
532 <https://doi.org/10.1080/00031305.2013.783881>
533

534 Wildermann, N.E., Sasso, C.R., Stokes, L.W., Snodgrass, D., Fuentes, M.M., 2019. Habitat use
535 and behavior of multiple species of marine turtles at a foraging area in the Northeastern
536 Gulf of Mexico. *Frontiers in Marine Science* 6, 155.
537 <https://doi.org/10.3389/fmars.2019.00155>
538

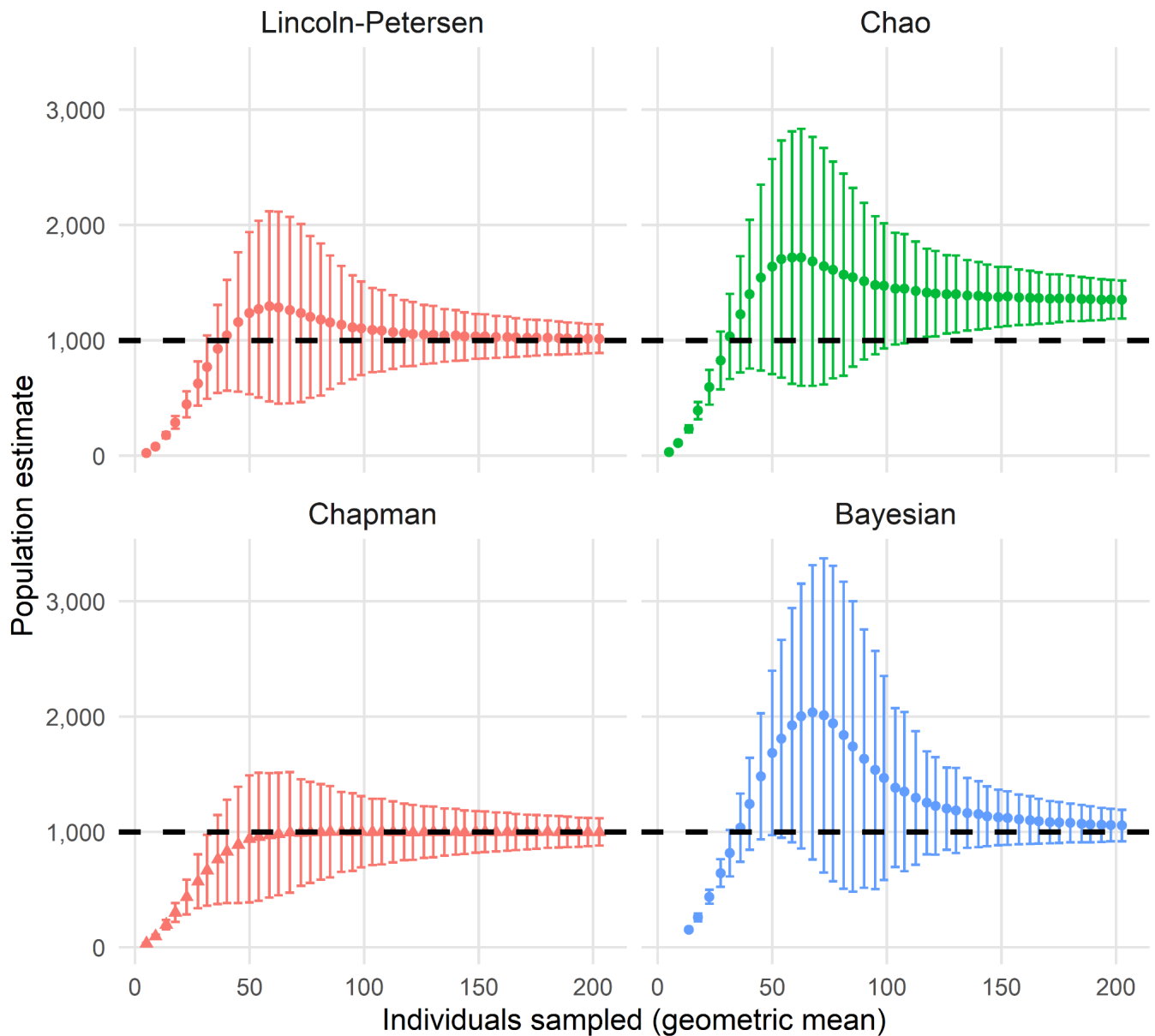
539 Wittes, J.T., 1972. Note: on the bias and estimated variance of Chapman's two-sample capture-
540 recapture population estimate. *Biometrics* 28(2), 592-597.
541 <https://doi.org/10.2307/2556173>
542

543 Zhou, K., Xu, X., Tian, C. 2007. Distribution and abundance of Indo-Pacific humpback dolphins
544 in Leizhou Bay, China. *N. Z. J. Zool.* 34(1), 35-42.
545 <https://doi.org/10.1080/03014220709510061>

546 Fig. 1. Two-visit simulated estimates of population size \hat{N} \pm standard errors across sample
547 sizes, denoted as the geometric mean over all visits, from a population of $N = 1,000$ individuals,
548 with M (i.e., n_1) $< n_2$.

$N = 1,000$, visits = 2

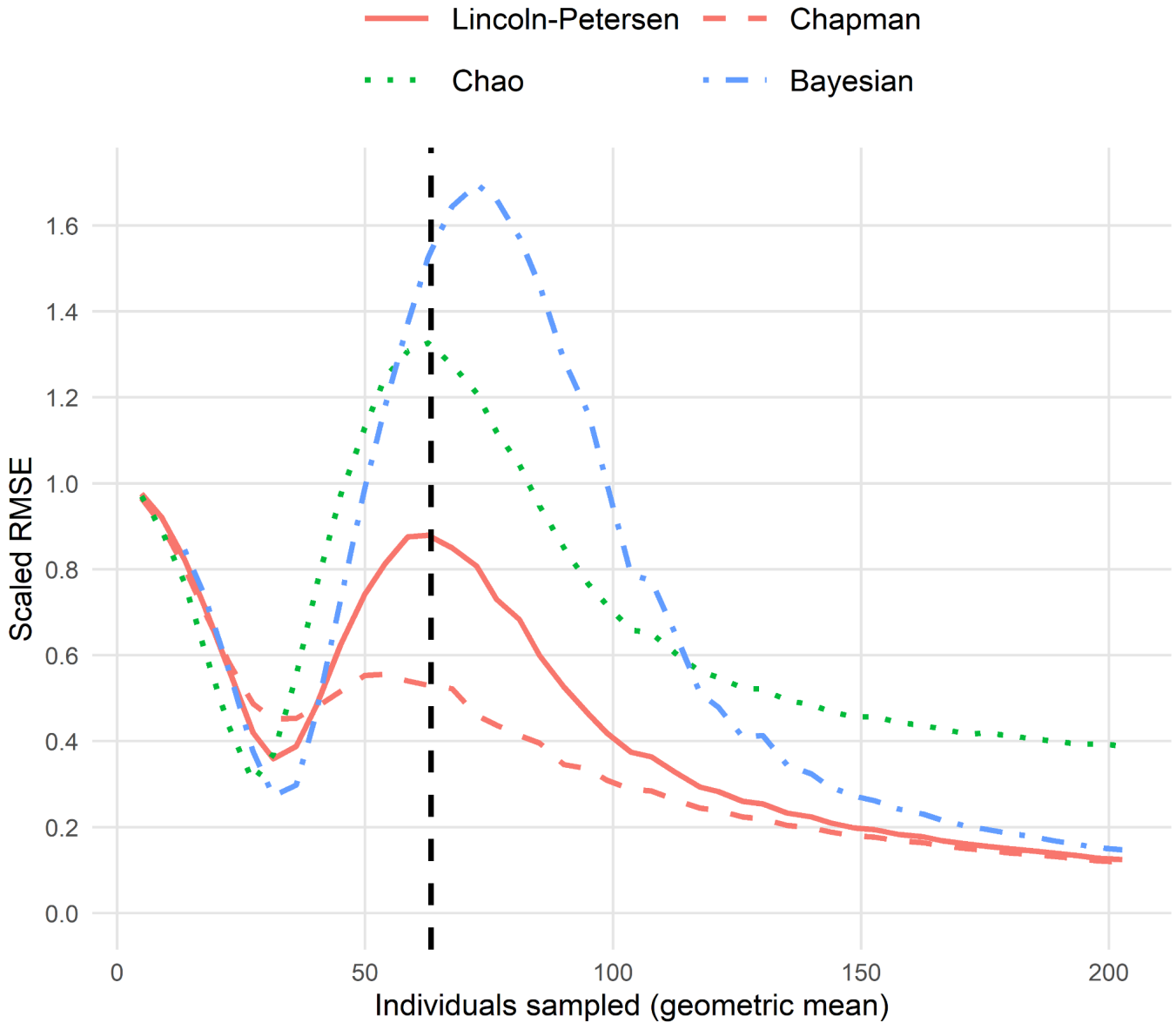
$M/\Sigma n = 0.25$



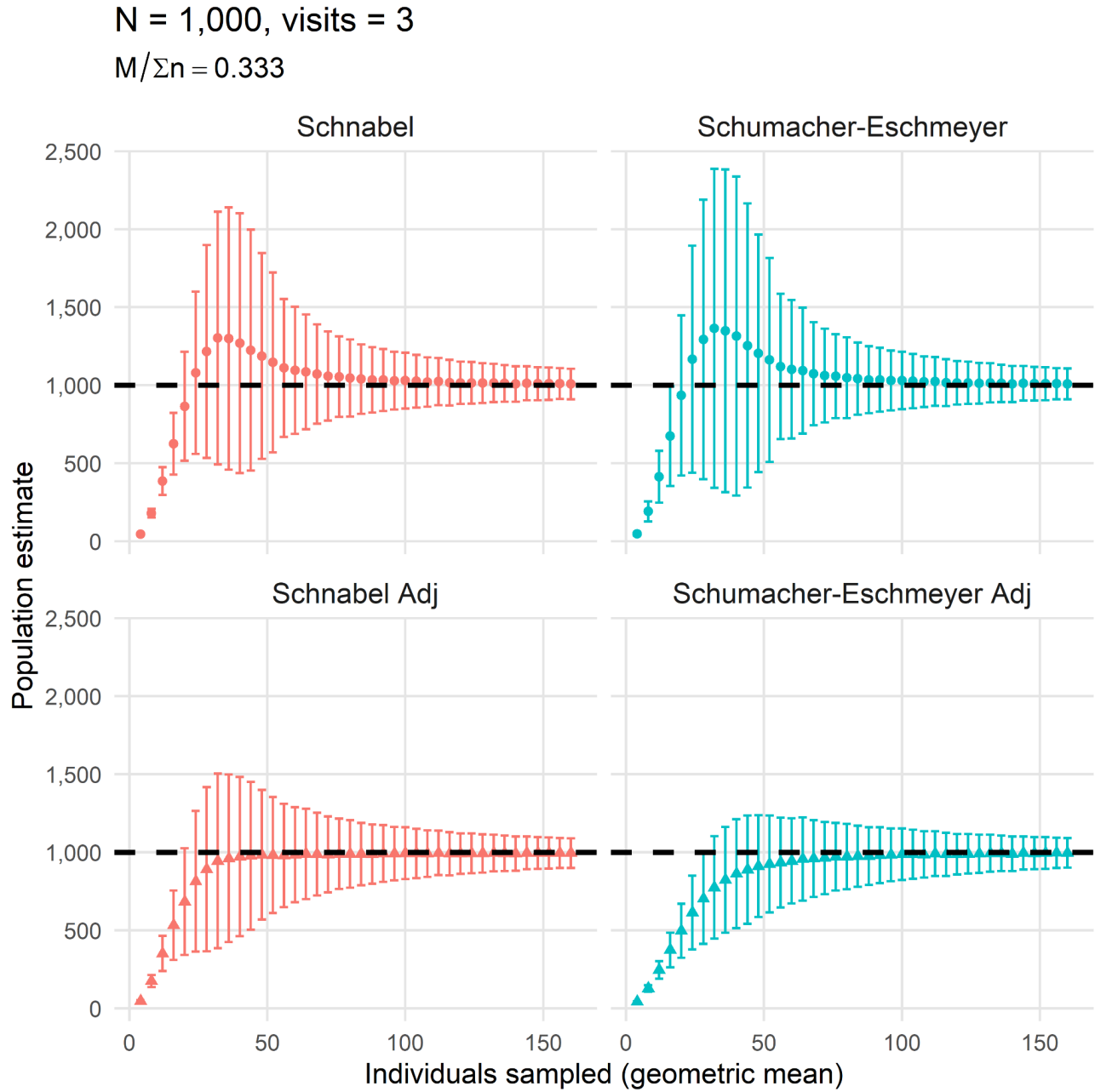
549 Fig. 2. Scaled root mean square error (RMSE) of the two-visit estimators across sample sizes,
550 denoted as the geometric mean over all visits, from a population of $N = 1,000$ and M (i.e., n_1) <
551 n_2 . Dashed vertical line represents the sample size at which the Chapman estimator becomes
552 approximately unbiased.

$N = 1,000$, visits = 2

$M/\Sigma n = 0.25$



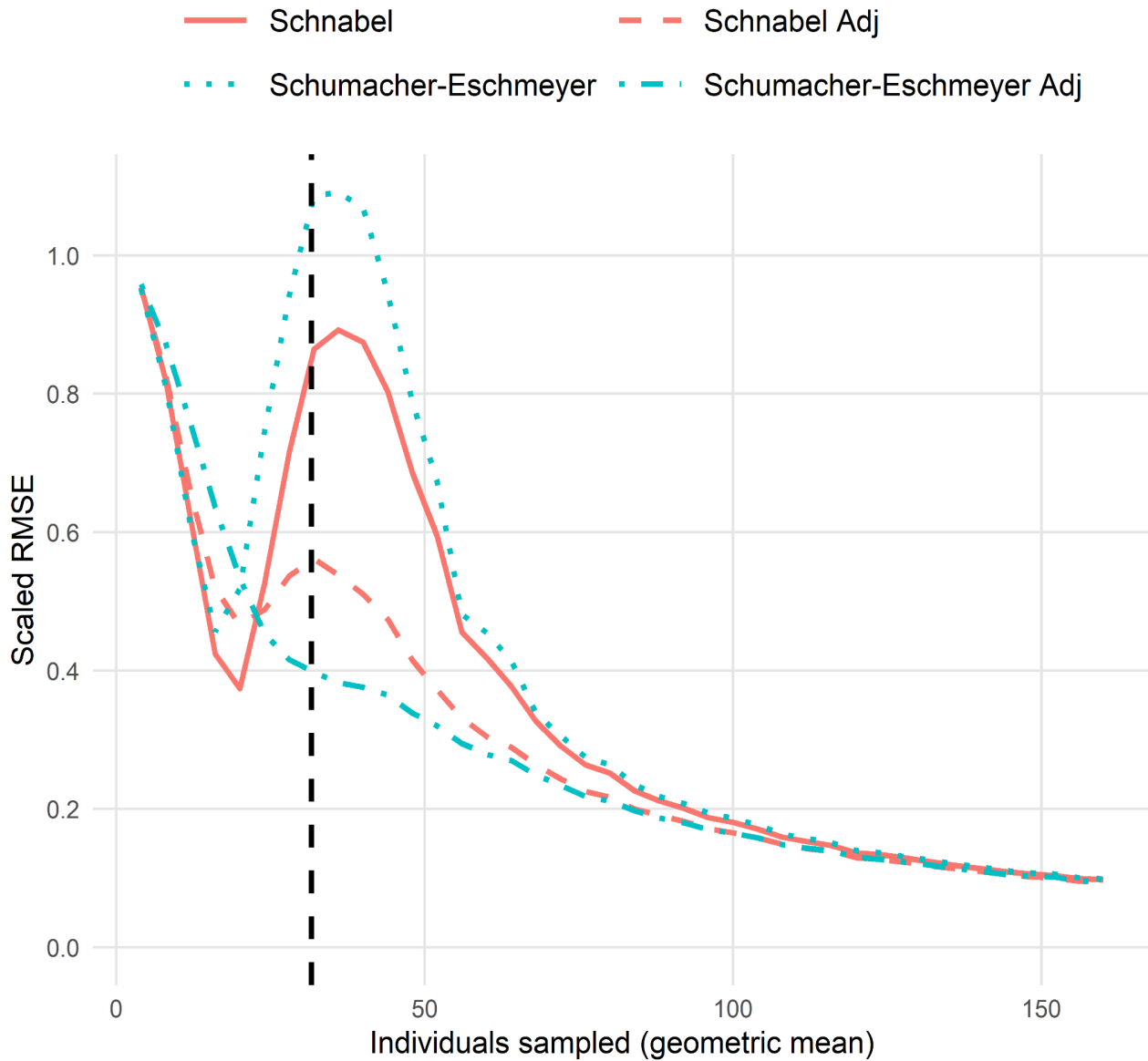
553 Fig. 3. Multi-visit ($k = 3$) simulated estimates of population size \hat{N} +/- standard errors across
554 sample sizes, denoted as the geometric mean over all visits, from a population of $N = 1,000$
555 individuals, with M (i.e., $n_1 = n_2 = n_3$).



556 Fig. 4. Scaled root mean square error (RMSE) of the multi-visit estimators across sample sizes,
557 denoted as the geometric mean over all visits, from a population of $N = 1,000$ and M (i.e., $n_1 =$
558 $n_2 = n_3$). Dashed vertical line represents the sample size at which the adjusted Schnabel estimator
559 becomes approximately unbiased.

$N = 1,000$, visits = 3

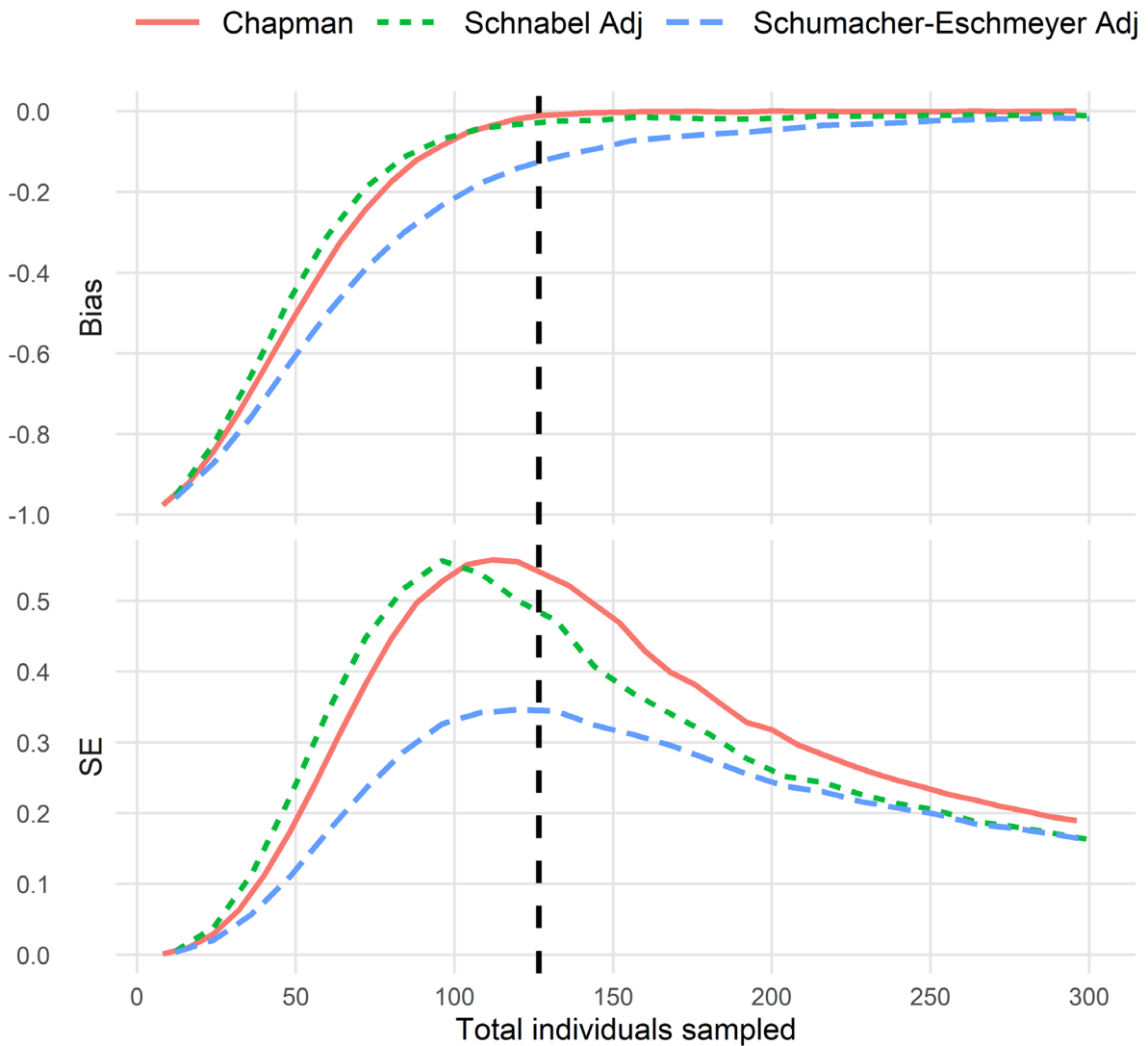
$M/\Sigma n = 0.333$



560 Fig. 5. Comparison of relative bias and standard error (SE) of the Chapman estimator with those
 561 of the adjusted multi-visit estimators across total sample size over all visits ($k = 2$ for Chapman, k
 562 $= 3$ for multi-visit estimators), from a population of $N = 1,000$ and M (i.e., $n_1) = n_2 (= n_3)$.
 563 Dashed vertical line represents the sample size at which the 2-visit Chapman estimator becomes
 564 approximately unbiased (Eq. 10). Note the Chapman estimator possesses slightly less downward
 565 bias at this threshold than the adjusted Schnabel for an equal number of total individuals sampled
 566 spread over 3 visits, with the advantage of lower variability in the adjusted Schnabel. The
 567 adjusted Schumacher-Eschmeyer possesses the lowest variability among methods at the expense
 568 of taking longest to become unbiased.
 569

$N = 1,000$, visits = 2, 3

$M = \Sigma n/k$



N	$k = 3$		$k = 4$		$k = 5$	
	\bar{N}	Pct. Error	\bar{N}	Pct. Error	\bar{N}	Pct. Error
100	89	-10.6	89	-10.8	87	-13.2
1,000	935	-6.5	912	-8.8	904	-9.6
10,000	9,394	-6.1	9,293	-7.1	9,138	-8.6
100,000	94,792	-5.2	93,062	-6.9	91,832	-8.2
1,000,000	948,038	-5.2	930,035	-7.0	916,636	-8.3

570
571 Tbl. 1. Relative bias in the adjusted Schnabel estimator (Eq. 6) for M (i.e., n_l) = $n_{2...k}$ at
572 recommended minimum sample sizes from (Eq. 11) for $k = 3, 4$ and 5 visits at various population
573 sizes N , based on 100,000 simulations.