

Effect of different sampling designs and methods on the estimation of secondary production: A simulation

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Web Appendix 1

This appendix describes 13 estimation techniques that were investigated in the Monte Carlo simulation study, in addition to the 7 classic methods. The notation used throughout this section is the same as in the main body of the text.

IS-2 Same as IS-1, except that

$$P_i = (B_{t_1 i \bullet} - B_{11 i \bullet}) + \sum_{t=t_1}^{T-1} (N_{(t+1)i \bullet} + N_{t i \bullet}) (W_{(t+1)i \bullet} - W_{t i \bullet}) / 2,$$

where t_1 is the sampling occasion on which $N_{t_1 i \bullet}$ is the largest, i.e., as per the suggestion of Siegismund (1982).

IS-3 Same as IS-1, but discarding values to ensure positive production, i.e., computing P_i using only the largest possible subset of times for which $W_{t i \bullet}$ increases in t , starting with $W_{1 i \bullet}$.

IS-4 Keeping all points as in IS-1, but replacing negative productions by zeros, i.e., using

$$P_i = \sum_{t=1}^{T-1} (N_{(t+1)i \bullet} + N_{t i \bullet}) \max(0, W_{(t+1)i \bullet} - W_{t i \bullet}) / 2.$$

IG-2 Same as IG-1, but replacing negative productions by zeros, i.e.,

$$P_i = \sum_{t=1}^{T-1} (B_{(t+1)i \bullet} + B_{t i \bullet}) \max \left\{ 0, \log \left(\frac{W_{(t+1)i \bullet}}{W_{t i \bullet}} \right) \right\} / 2.$$

AC-2 Same as AC-1, but discarding points prior to the largest N , i.e., for each cohort i , find the time t_i at which $N_{t_i \bullet}$ is maximized, and discard couples $(N_{t \bullet}, W_{t \bullet})$ for which $t < t_i$.

AC-3 An Allen curve obtained by merging all cohorts in one; specifically, sampling cohort i at time T_i was treated as equivalent to sampling cohort 1 at time $365(i-1) + T_i$ and the curve $\log N = \alpha + \beta W$ was fitted, which led to

$$P = \frac{e^\alpha}{\beta} (e^{\beta W_{\max}} - e^{\beta W_{\min}})$$

with W_{\max} and W_{\min} , respectively, standing for the largest and smallest weights, overall.

SF-2 A variant of SF-1 proposed by Krueger and Martin (1980), i.e.,

$$P = K \sum_{k=1}^{K-1} (N_{\bullet \bullet (k+1)} - N_{\bullet \bullet k}) \sqrt{W_{\bullet \bullet (k+1)} W_{\bullet \bullet k}}.$$

SF-3 Another variant of SF-1 mentioned by Cusson (2004), i.e.,

$$P = K \sum_{k=1}^{K-1} (N_{\bullet \bullet (k+1)} + N_{\bullet \bullet k}) (W_{\bullet \bullet (k+1)} - W_{\bullet \bullet k}) / 2.$$

GR-2 A variant of GR-1 with a von Bertalanffy adjustment curve, i.e.,

$$P = K \sum_{k=1}^{K-1} B_{\bullet \bullet k} \left\{ \hat{a} - \hat{a} \frac{L_k}{\ell(k)} \right\} \log \left\{ \frac{\hat{L}_{\max} - \ell(k+1)}{\hat{L}_{\max} - \ell(k)} \right\},$$

where $\ell(k) = (L_k + L_{k+1})/2$ and \hat{L}_{\max} is obtained by fitting a von Bertalanffy growth curve

$$L(T) = \hat{L}_{\max} \{1 - \exp \hat{\lambda} (T - \hat{T}_0)\},$$

while \hat{a} results from adjusting the linear regression curve

$$\log(W) = \hat{a} \log(L) + \hat{b}.$$

In both cases, the adjustment is based on cohort-merged data.

GR-3 A variant of GR-1 assuming length growth to be linear in time, i.e.,

$$P = I \sum_{k=1}^K B_{\bullet \bullet k} \hat{a} \hat{\lambda} \left\{ 1 - \frac{\hat{L}_{\max}}{\ell(k)} \right\}$$

where \hat{a} , $\ell(k)$, \hat{L}_{\max} , and $\hat{\lambda}$ are as defined above. The same substitution as in GR-2 was made, except that time was assumed to be a linear function of the size in length.

GR-4 A variant of GR-1 using length as a surrogate for time, i.e.,

$$P = K \sum_{k=1}^K B_{\bullet\bullet k} \frac{\hat{a}}{k},$$

where \hat{a} is the same as above.

MR-2 A variant of MR-1 assuming that length growth is linear in time, i.e.,

$$P = -\hat{z} l \sum_{k=1}^K B_{\bullet\bullet k},$$

where \hat{z} was obtained from regressing $\log(N) = \hat{z}_0 + \hat{z} T$ on the cohort-merged data (as in AC-3).

MR-3 Similar to MR-2, but using length as a surrogate for time, i.e.,

$$P = K \sum_{k=1}^K B_{\bullet\bullet k} \left\{ \frac{\hat{z}}{\hat{\lambda}(K+1-k)} \right\},$$

where \hat{z} is defined as in MR-2 and $\hat{\lambda}$ has the same meaning as in GR-2.