Received XXXX

(www.interscience.wiley.com) DOI: 10.1002/sim.0000

Statistics in Medicine

Likelihood-based analysis of outcome dependent sampling designs with longitudinal data: supplementary materials

Leila R. Zelnick^{a*}, Jonathan S. Schildcrout^b, and Patrick J. Heagerty^c

^a *Department of Medicine, University of Washington, Seattle, WA, 98195, USA*

^b *Department of Biostatistics, Vanderbilt School of Medicine, Nashville, TN, 37203, USA*

^c *Department of Biostatistics, University of Washington, Seattle, WA, 98195, USA*

⇤*Correspondence to: Department of Medicine, University of Washington, Seattle, WA, 98195, USA. Email: lzelnick@uw.edu*

Appendix A: Likelihood-based estimator using subsampled subjects only and with covariates

The likelihood SO,WC (Equation 4) differs from that of likelihood SO,NC (Equation 3) by the second term in Equation 5. Without loss of generality, suppose M_i is a binary marker. Define $a_0 = AC_0(M_i = 0, T_i; \theta)$ and $a_1 = AC_0(M_i = 0; T_i; \theta)$ 1, T_i ; θ). Under a complete and balanced design, $P(M_i = m_i, T_i; \Gamma) = P(M_i = m_i; \Gamma)$ and $\mathbb{E}_{M_i}[\Gamma_i[AC_0(M_i, T_i)] =$ $\mathbb{E}_{M_i}[AC_0(M_i, \mathcal{T}_i)]$, and the *i*th person's ascertainment correction contribution to the likelihood (Equation 5 in main text) can be written as follows:

$$
\frac{AC_0(M_i, T_i; \theta) \cdot P(M_i = m_i, T_i; \Gamma)}{\mathbb{E}_{M_i}[AC_0(M_i, T_i; \theta)]} = \frac{AC_0(M_i, T_i; \theta) \cdot P(M_i = m_i; \Gamma)}{\mathbb{E}_{M_i}[AC_0(M_i, T_i; \theta)]}
$$
\n
$$
= \frac{a_1^m a_0^{1-m} \cdot p^m (1-p)^{1-m}}{p a_1 + (1-p) a_0}
$$
\n
$$
= \frac{(a_1 p)^m (a_0 (1-p))^{1-m}}{p a_1 + (1-p) a_0}
$$
\n
$$
= \frac{\left(\frac{a_1}{a_0}p\right)^m (1-p)^{1-m}}{\frac{a_1}{a_0}p + 1 - p}
$$
\n
$$
= \frac{\frac{a_0}{a_1 p}}{\frac{a_0}{a_1 p}} \cdot \frac{\left(\frac{a_1}{a_0}p\right)^m (1-p)^{1-m}}{\frac{a_1}{a_0}p + 1 - p}
$$
\n
$$
= \left(\frac{1}{1 + \frac{a_0}{a_1} \cdot \frac{1-p}{p}}\right)^m \left(\frac{\frac{a_0}{a_1} \cdot \frac{1-p}{p}}{1 + \frac{a_0}{a_1} \cdot \frac{1-p}{p}}\right)^{1-m}
$$
\n
$$
= \xi^m (1 - \xi)^{1-m},
$$

where $\xi =$ $\begin{pmatrix} 1 \end{pmatrix}$ $1 + \frac{a_0}{a_1} \cdot \frac{1-p}{p}$! . For balanced and complete designs, a_0 and a_1 do not vary across subjects, so this term's contribution to likelihood SO,WC (Equation 4) can be seen as a reparameterization of the marginal distribution of *M*, which contains no information about θ . Thus, adding this term to the likelihood *SO, NC* (Equation 3) will add no information to the resulting inference, and in fact will yield the same estimate, up to the chosen tolerance of the Newton-Raphson algorithm.

L.R. Zelnick et al.

Supplementary Table 1

Supplementary Table 1. Percent bias and relative efficiency for likelihood-based ODS estimators, under low subjectto-subject heterogeneity. Results shown summarize 1000 replications with $N = 1000, \beta = (\beta_0, \beta_T, \beta_M, \beta_{M \times T}) =$ $(10, -0.25, -0.75, 0.5), \sigma_{b_0}^2 = 4, \sigma_{b_1}^2 = 0.25, \sigma_e^2 = 1$, and $\rho = 0$. $N_S = 250$ subjects were subsampled on average. Percent bias defined as the $100 \times$ the difference between estimator mean and parameter value, divided by parameter value, and relative efficiency for an estimator is defined as the ratio of variances between a random sample of $N_S = 250$ and the estimator. NA for percent bias indicates that percent bias is undefined.

Supplementary Table 2

Supplementary Table 2. Percent bias and relative efficiency for likelihood-based ODS estimators, under high subjectto-subject heterogeneity. Results shown summarize 1000 replications with $N = 1000, \beta = (\beta_0, \beta_T, \beta_M, \beta_{M \times T}) =$ $(10, -0.25, -0.75, 0.5), \sigma_{b_0}^2 = 4, \sigma_{b_1}^2 = 4, \sigma_e^2 = 4$, and $\rho = 0$. $N_S = 250$ subjects were subsampled on average. Percent bias defined as the $100 \times$ the difference between estimator mean and parameter value, divided by parameter value, and relative efficiency for an estimator is defined as the ratio of variances between a random sample of $N_S = 250$ and the estimator. NA for percent bias indicates that percent bias is undefined.

Supplementary Table 3

Supplementary Table 3. Percent bias and relative efficiency for time-specific predicted means and predicted difference in means, under high subject-to-subject heterogeneity. Results shown summarize 1000 replications with $N = 1000$, $\beta =$ $(\beta_0, \beta_T, \beta_M, \beta_{M \times T}) = (10, -0.25, -0.75, 0.5), \sigma_{b_0}^2 = 4, \sigma_{b_1}^2 = 4, \sigma_e^2 = 4, \text{ and } \rho = 0. N_S = 250 \text{ subjects were subsampled}$ on average. Percent bias defined as the $100 \times$ the difference between estimator mean and parameter value, divided by parameter value, and relative efficiency for an estimator is defined as the ratio of variances between a random sample of $N_S = 250$ and the estimator.

Supplementary Table 4

Supplementary Table 4. Percent bias and relative efficiency for time-specific predicted means and predicted difference in means, under low subject-to-subject heterogeneity. Results shown summarize 1000 replications with $N =$ 1000, $\beta = (\beta_0, \beta_T, \beta_M, \beta_{M \times T}) = (10, -0.25, -0.75, 0.5), \sigma_{b_0}^2 = 4, \sigma_{b_1}^2 = 0.25, \sigma_e^2 = 1$, and $\rho = 0$. $N_S = 250$ subjects were subsampled on average. Percent bias defined as the $100 \times$ the difference between estimator mean and parameter value, divided by parameter value, and relative efficiency for an estimator is defined as the ratio of variances between a random sample of $N_S = 250$ and the estimator.

L.R. Zelnick et al.

Supplementary Table 5

Supplementary Table 5. Baseline characteristics of Cystic Fibrosis Foundation Registry cohort ($N = 3,141$).

Values are mean (SD), except as noted.

Supplementary Figure 1

 \circ 50 å å $\frac{4}{10}$ CF subject-specific slopes c.c \tilde{c} 통 $\frac{0}{0}$ $\ddot{}$ Ş $\frac{1}{\alpha}$ $\overline{\mathbf{2}}$ 1 3 4 CF subject-specific intercepts œ o opocoo o

Supplementary Figure 1. Subject-specific intercepts and slopes resulting from regressing a CF patient's *FEV*¹ longitudinal outcome on time. The distribution of intercepts and slopes do not appear to be clearly inconsistent with bivariate normality.