

www.ancienttl.org · ISSN: 2693-0935

Berger,G. W., 2011. *Errata: Estimating the error in equivalent dose values obtained from SAR*. Ancient TL 29(1): 51. https://doi.org/10.26034/la.atl.2011.449

This article is published under a *Creative Commons Attribution 4.0 International* (CC BY): https://creativecommons.org/licenses/by/4.0



© The Author(s), 2011

Errata: Estimating the error in equivalent dose values obtained from SAR

G. W. Berger

Desert Research Institute, 2215 Raggio Parkway, Reno, NV 89512, USA (e-mail: glenn.berger@dri.edu)

(Received 16 April 2011)

Errata

In the paper by Berger (2010) there are some typographical/transcription errors in some of the equations, all but one error relating to the saturating-exponential (E) model. Only one of these errors (that in the matrix component I_{aa} concerning the E model) occurred also in the author's software and led to some incorrect error estimates for some of the D_E values derived from the E-model data sets, but had no effect on the computed D_E values and on the best-fit dose-response curves (DRC) for the E model.

In equation 12 of Berger (2010),

$$\Delta A = ([WU]^{t}[WU])^{-1}([WU]^{t}[WY^{*}])$$

for the iterative calculation of the best-fit parameters, the matrix W should be replaced by \sqrt{W} . In the immediately subsequent equation for wy*, brackets were inadvertently omitted during transcription. The correct equation is

$$wy^* = [y_i - a(1 - e^{-bx_i})]\sqrt{w_i}$$

In the subsequent subsection "5.2 Error in D_E ", the equations for the matrix components I_{aa} , I_{bb} and I_{ab} are incorrect. Certain parameters were inadvertently omitted during transcription. The correct expressions are as follows:

$$I_{aa} = \sum_{i} w_i (1 - e^{-bx_i})^2$$
$$I_{bb} = \sum_{i} w_i (ax_i e^{-bx_i})^2$$
$$I_{ab} = I_{ba} = \sum_{i} w_i f_i x_i e^{-bx_i}$$

Finally, at the end of the first paragraph in the subsection "6.1 Regression to obtain a, b, c" for the E+L model, the stated matrices WU and WY^* should be replaced by $\sqrt{W}U$ and $\sqrt{W}Y^*$.

The consequence of the coding error in the equation for I_{aa} (used in the calculation of errors in the D_E values shown in the paper) is as follows. In Table 1, the last two D_E values in the last column should read 0.698 \pm 0.062 (not \pm 0.058), and 28.48 \pm 0.69 (not \pm 0.68). In Table 3, the only changes (all in the last column) are: 2104 \pm 255 (not \pm 128), and 1417 \pm 668 (not \pm 148). These changes put the author's error estimates (for the E model) in Table 3 closer to those from Duller's (2007) 'curve-fitting' error estimates, and strengthen one of Berger's (2010) conclusions: that the two error-estimation schemes (Berger's and Duller's) generally produce no significantly different error estimates.

References

- Berger, G.W. (2010) Estimating the error in equivalent dose values obtained from SAR. *Ancient TL* 28, 55-66.
- Duller, G.A.T. (2007) Assessing the error on equivalent dose estimates derived from single aliquot regenerative dose measurements. *Ancient TL* **25**, 15-24.