CORRIGENDA

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Pages G872–G884: R. A. Weisiger, S. Pond, and L. Bass. “Hepatic uptake of protein-bound ligands: extended sinusoidal perfusion model.” Page G879 (2nd column, 1st line): the sentence that reads “As for the standard model, the only fitted parameter was $P_m$ (resulting value 0.248 ± 0.051 cm/s)” should read “As for the standard model, the only fitted parameter was $P_m$ (resulting value 0.082 ± 0.004 cm/s)” The only effect of this error on the results and conclusions of this study is that the best-fit line in Fig. 3 is very slightly different in shape, with a correlation coefficient ($r^2$) that still exceeds 0.999. The corrected figure is provided below.

![Corrected Figure 3](http://ajpgi.physiology.org/)

Fig. 3. Fit of standard and extended models to date for steady oleate uptake. Solid circles show mean oleate uptake velocity by the perfused rat liver for the current study. Oleate and albumin concentrations were varied at a fixed 1:10 molar ratio. Standard error bars are shown where they are larger than the symbols used. Solid line shows predictions of the extended sinusoidal perfusion model and broken line shows predictions of the standard (unextended) sinusoidal perfusion model. In both cases, $P_m$ is treated as an unknown variable and all other parameter values are as in Table 1. For fitting, each point was weighted in inverse proportion to its magnitude.