Correction: New Maximum Likelihood Estimators for Eukaryotic Intron Evolution
Hung D. Nguyen, Maki Yoshihama, Naoya Kenmochi
DOI: 10.1371/journal.pcbi.0010079
In PLoS Computational Biology, volume 1, issue 7:
An $n_i$ was omitted from Equation 6. The correct equation is as shown below:

$$L(\theta, \lambda, \alpha_k, \beta_k | \text{data}) = \left( \prod_{i=0}^{2^{N-1}} \frac{p_i^{n_i}}{n_i!} \right) \times P!$$

This correction note may be found online at DOI: 10.1371/journal.pcbi.0020028.
Published March 31, 2006.
Citation: Correction: New maximum likelihood estimators for eukaryotic intron evolution. PLoS Comput Biol 2(3): e28.

Correction: Structure Modeling of All Identified G Protein–Coupled Receptors in the Human Genome
Yang Zhang, Mark E. DeVries, Jeffrey Skolnick
DOI: 10.1371/journal.pcbi.0020013
In PLoS Computational Biology, volume 2, issue 2:
The URL provided for the GPCR model database in the published article is no longer active. The database is now located at http://cssb.biology.gatech.edu/skolnick/files/gpcr/gpcr.html.

This correction note may be found online at DOI: 10.1371/journal.pcbi.0020029.
Published March 31, 2006.
Citation: Correction: Structure modeling of all identified G protein–coupled receptors in the human genome. PLoS Comput Biol 2(3): e29.