## Errata I

Yang 2014 Molecular Evolution: A Statistical Approach

| Page | Incorrect | Correct | Notes |
| :---: | :---: | :---: | :---: |
| p. 3 table 1.1 F84 row A column C | $\beta \pi_{\text {T }}$ | $\beta \pi_{\mathrm{C}}$ |  |
| p. 67 eq. 2.38 | $V_{0}$ | $U_{0}^{-1}$ |  |
| p. 105 figure 4.2 last number for node 6 | 0.004344 | 0.000444 |  |
| p. 106 line - 23 | $0.004344=0.003070$ | $0.000444=0.002976$ |  |
| p. 123 line 15 | Yang, Lauder, and Lin (1995b; | Yang, Lauder, and Lin (1995c; |  |
| p. 138 line - 12 | $\begin{aligned} & \pi_{1}=x_{1} / s, \pi_{2}=x_{2} / s, \pi_{3} \\ & =x_{3} / \mathrm{s} \end{aligned}$ | $\begin{aligned} & \pi_{1}=\mathrm{e}^{x_{1}} / \mathrm{s}, \pi_{2}=\mathrm{e}^{x_{2}} / \mathrm{s}, \\ & \pi_{3}=\mathrm{e}^{x_{3}} / \mathrm{s} \end{aligned}$ |  |
| p. 182 1.-7 | simple and composite | null and alternative |  |
| p. 188 below eq. 2.15 | for all $n$. | for all $x$. |  |
| p. 196 in Eq. 6.33 | $\exp \{-$ | $\exp \{$ | delete the minus sign |
| p. 196 line -12 | $\left(L_{1}+L_{1}\right)$ | $\left(L_{0}+L_{1}\right)$ |  |
| p. 197 line 16 | Bayesians | Bayesian |  |
| p. 198 line 2 | have the same K-L divergence from the true model | are all correct and encompass the true model |  |
| p. 204 below eq. 6.66 | $V=-H^{-1}$ | $V=-(n H)^{-1}$ |  |
| p. 207 line -3 | (13.24) | (A.24) |  |
| p. 209 fig. 6.8b |  |  | The four lines with points should be moved to the right to align with the x axis. |
| p. 2281.6 | ( $\alpha+1$ ) | ( $\alpha-1$ ) |  |
| p. 2301.18 | with $x>7$ | with $\theta>7$ |  |
| p. 232 above eq. 7.37 | $E \Lambda^{n} E^{-T}$ | $E \Lambda^{n} E^{-1}$ | change -T into -1 |
| p.233, 3 lines above eq. <br> 7.40 . | gives the efficiency | gives the variance |  |
| p. 234 line -6, | k | K | twice |
| p. 238 line -7 | $E \Lambda^{n} E^{-T}$ | $E \Lambda^{n} E^{-1}$ | change $-T$ into -1 |
| p. 244 lines -14 \& -13 | $\rho_{k}+\rho_{k+1}$ | $\rho_{2 k}+\rho_{2 k+1}$ |  |
| p. 282 paragraph 1 | If the branch lengths (and other parameters) are from the posterior for the |  | This statement is true only in the case where $\alpha$ in equation (8.21) is $<1$ for all possible $t$ |

current tree, the
branch lengths in the new tree (when the algorithm moves to the new tree through a cross-tree move) will automatically be from the stationary (posterior)
distribution for the new tree.
p.294, lines 2-3
p.294, first line within eq.
8.29
p. 310 fig. 9.1 b
p. 329 line -7
p. 337 below eq. 9.47
p. 335 line -3
p. 336 paragraph 1 in Sec
9.4.3.2
p. 338 paragraph above Sec
9.4.3.3
p. 338 paragraph 1 in Sec
9.4.3.3
p. 3561.2
p. 422 1.-3
p. 480
p. 196 in Eq. 6.33
p. 196 line - 12
the proposal ratio will be one. Otherwise the proposal ratio will be the ratio of the uniform and exponential densities for the age of node $a$ :
the proposal ratio will be the ratio of the uniform densities for the age of node $a$ at the source and target. Otherwise it will be the ratio of the uniform and exponential densities.
$\frac{1 /\left(t_{u}-\max \left\{t_{x}, t_{v}\right\}\right)}{1 /\left(t_{b}-\max \left\{t_{x}, t_{y}\right\}\right)}$
MRCA $\rightarrow$

## $\tau \mathrm{s}$

species
Liu et al. (2010a)
Liu et al. (2010b)

Liu et al. (2010b)

Liu et al. (2010a)
(Zhang et al. 2011)
if $j \leq i, \operatorname{set} j=j+1$
Steel, M.
$\exp \{-$
$\left(L_{1}+L_{1}\right)$
and $\boldsymbol{t}^{\prime}$ but is in general incorrect. In the latter case, use of withinchain moves will benefit the mixing efficiency for estimating posterior model (tree) probabilities.

Label the mother node of node $a$ as $u$ and the daughter node as $v$. See attached figure.
arrow should point to the node in the gene tree
remove space
twice
twice, and reorder references
delete the minus sign

| p. 198 line 2 | have the same K-L <br> divergence from the <br> true model | are all correct and <br> encompass the true <br> model |
| :--- | :--- | :--- |
| p. 233 line 3 below Eq. <br> 7.40 | $E=1 /\left(2 \pi_{2}-1\right)$ | $E=1 /\left(1-2 \pi_{2}\right)$ |
| p. 257 table 7.6 first <br> column | $B_{01}$ | $B_{10}=1 / B_{01}$ |
| p. 361 line -3 | habits | habitats |
| p. 422 line -3 | if $j \leq i$, | if $j \geq i$, |

Thanks to Utkarsh J Dang, Chi Zhang, Tianqi Zhu for corrections.

We decided to set symbols for vectors in bold italic $(\boldsymbol{x}, \boldsymbol{y}, \boldsymbol{t})$ rather than bold ( $\mathbf{x}, \mathbf{y}, \mathbf{t}$ ). However this rule is not consistently applied. The following lists the page numbers (and number of occurrences) where the symbols should be corrected.
p. 102 (1 time), p. 103 (6 times), p. 104 (2), p. 105 (1), p. 106 (2), p. 107 (1), p. 111 (15), p. 112 (8), p. 114 (4), p. 115 (5), p. 117 (2), p. 120 (3), p123 (1), p127 (6), p. 128 (17), p. 130 (13), p. 131 (2), p. 139 (1), p. 172 (1), p. 178 (3), p. 226 (10), p. 267 (2), p. 268 (12), p. 269 (1), p. 281 (2), p. 329 (1), p. 382 (2), p. 408 (4), p. 447 (1).

