

Errata for *Population Genomics with R*

Page xiv: (fifth line from bottom) oppotunity \rightarrow opportunity

Page 14: (row describing `psmc`) sequantially \rightarrow sequentially

Page 15: (end of caption of Fig. 1.9) selection of on loci. \rightarrow selection on other loci.

Page 17: (fifth line of Sect. 2.1.1) it will be to ~~to~~ extract DNA

Page 56: (before the last bloc of R code) the file names \rightarrow the file named

Page 70: (third item in the list near the bottom of the page)
the operation can reversed \rightarrow the operation can be reversed

Page 176: (in the paragraph after the first block of R lines) because ~~of~~ the ~~very~~ predicted

Page 233: (line 10) `by(xsel)` \rightarrow `by(x)`

Page 245: (near the end of the first paragraph) the \rightarrow then

Page 303: the scales of the axes of Fig. 9.16 are wrong because of several small bugs in `psmc` which have been fixed in version 0.1-2 of this package. Mostly, the axis scales were wrongly multiplied by 10^4 . The corrected figure is shown next page (slightly different because the bootstraps were re-run).

December 23, 2025

