Errata for Population Genomics with R

Page 14: (row describing psmcr) sequantially ∧→ sequentially

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

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

Page 70: (third item in the list near the bottom of the page) the operation can reversed \rightsquigarrow the operation can <u>be</u> reversed

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

Page 233: (line 10) by(xsel) \rightsquigarrow by(x)

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

Page 303: the scales of the axes of Fig. 9.16 are wrong because of several small bugs in psmcr 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).

May 10, 2025

