

## Errata for *Population Genomics with R*

- Page xiv:** (fifth line from bottom) opportunity  $\nwarrow$  opportunity
- Page 14:** (row describing `psmcr`) sequantially  $\nwarrow$  sequentially
- Page 15:** (end of caption of Fig. 1.9) selection of on loci.  $\nwarrow$  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  $\nwarrow$  the file named
- Page 70:** (third item in the list near the bottom of the page)  
the operation can reversed  $\nwarrow$  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)`  $\nwarrow$  `by(x)`
- Page 245:** (near the end of the first paragraph) the  $\nwarrow$  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).

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