Correction

Correction: Gene Flow within and between Catchments in the Threatened Riparian Plant *Myricaria germanica*

The *PLOS ONE* Staff

The images for Figure 2 and Figure 3 were inadvertently swapped. Please view the correct images and legends for Figure 2 and Figure 3.

Citation: The *PLOS ONE* Staff (2014) Correction: Gene Flow within and between Catchments in the Threatened Riparian Plant *Myricaria germanica*. PLoS ONE 9(7): e103669. doi:10.1371/journal.pone.0103669

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Figure 2. Population graphs showing the genetic relationships between *Myricaria germanica* sites, with catchments shown in different colors (Inn: blue; Rhine, orange; Rhone, red; Ticino, yellow); the size of circles is proportional to effective population size.
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Figure 3. Allelic richness and population-specific inbreeding coefficients (F_{IS}) of 20 nuclear SSR in *Myricaria germanica*. A and E. Relationship with elevation. B and F. Relationship with log10-transformed census population size. D. Relationship between allelic richness and F_{IS} . C and G. Boxplots of allelic richness, grouped by affiliation to genetic clusters (see Fig. 1B). H. Boxplots of F_{IS} , grouped by affiliation to catchment. The values plotted for census population size are midpoints of the estimated intervals of population size. Lines represent linear regressions. Cluster 1: Rhine, Rhone, Ticino. Cluster 2: Inn.
doi:10.1371/journal.pone.0099400.g003

In addition, some labels were removed from Figure 3 during the production process. The publisher apologizes for this error. Please see the corrected figure above.

Reference

1. Werth S, Scheidegger C (2014) Gene Flow within and between Catchments in the Threatened Riparian Plant *Myricaria germanica*. PLoS ONE 9(6): e99400. doi:10.1371/journal.pone.0099400