Erratum to: Hierarchical spatial genetic structure in a distinct population segment of greater sage-grouse

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Unfortunately, Fig. 5 has been published incorrectly (the polygons were missing) in the original publication. The corrected Fig. 5 is published below.

The online version of the original article can be found under doi:[10.1007/s10592-014-0618-8](https://doi.org/10.1007/s10592-014-0618-8).

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Fig. 5 Factorial components analysis of Bi-State and neighboring populations of greater sage-grouse using 14 microsatellite loci. Samples in the *red polygon* represent the individuals in the neighboring populations of Churchill and Lassen. Samples in the *blue polygon* represent a subset of individuals from the PN sampling locale (PNa). Samples in the *green polygon* represent the remaining samples in the Bi-State. The individuals within the *green polygon* that are *pink* are from the WM sampling locale (Color figure online)