Correction to ‘RAD sequencing and genomic simulations resolve hybrid origins within North American Canis’

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The data associated with Rutledge et al. [1] are now available on the Dryad Digital Repository at http://dx.doi.org/10.5061/dryad.pr318 [2].

References

1. Rutledge LY, Devillard S, Boone JQ, Hohenlohe PA, White BN. 2015 RAD sequencing and genomic simulations resolve hybrid origins within North American Canis. Biol. Lett. 11, 20150303. (doi:10.1098/rsbl.2015.0303)

2. Rutledge LY, Devillard S, Boone JQ, Hohenlohe PA, White BN. 2015 Data from: RAD sequencing and genomic simulations resolve hybrid origins within North American Canis. Dryad Digital Repository. (http://dx.doi.org/10.5061/dryad.pr318)