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## Recent Work

### Title

Correction to: Ancient polymorphisms contribute to genome-wide variation by long-term balancing selection and divergent sorting in *Boechera stricta*.

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CORRECTION

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# Correction to: Ancient polymorphisms contribute to genome-wide variation by long-term balancing selection and divergent sorting in *Boecheira stricta*

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**Correction to: *Genome Biol* (2019) 20:126**  
<https://doi.org/10.1186/s13059-019-1729-9>

Following publication of the original article [1], the authors reported that the Availability of data and materials section required updating. The updated text reads as follows:

The short reads of each genotype have been deposited under GenBank accession numbers SRP054739, SRP134283-SRP134373, SRP134393-SRP134433, SRP134436-SRP134479, SRP134481-SRP134572 and SRP134581-SRP134671. All SNPs used in population genetic analyses, locations of all accessions, and custom scripts are available in the Dryad Data Archive at <https://doi.org/10.5061/dryad.574pc6n> [66]. Seeds from these accessions are available from the Arabidopsis Biological Resource Center.

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1. Wang B, et al. Ancient polymorphisms contribute to genome-wide variation by long-term balancing selection and divergent sorting in *Boecheira stricta*. *Genome Biol.* 2019;20:126. <https://doi.org/10.1186/s13059-019-1729-9>.

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