

Erratum to: A new resource of single nucleotide polymorphisms in the Japanese eel *Anguilla japonica* derived from restriction site-associated DNA

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Published online: 4 August 2016
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Erratum to: Ichthyol Res DOI 10.1007/s10228-016-0518-7

The original publication of this article contains an erroneous description, which is corrected as stated below.

In the first paragraph of “Results”, the fourth sentence reads, “A preliminary analysis using our own perl script found 37,952 positions of *Sbf*I recognition sites in the reference genome ... 79,836 *Sbf*I-based RAD loci in the *A. japonica* genome (twice the number of *Sbf*I recognition sites)”. However, the number of *Sbf*I recognition sites in the reference genome should be 39,918.

The online version of the original article can be found under doi:[10.1007/s10228-016-0518-7](https://doi.org/10.1007/s10228-016-0518-7).

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