



Correction

## Correction: Ramzan F. et al. "Combining Random Forests and a Signal Detection Method Leads to the Robust Detection of Genotype-Phenotype Associations" Genes, 2020, 11, 892

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Received: 22 September 2020; Accepted: 6 October 2020; Published: 14 October 2020



The authors would like to make a correction to the published paper [1]:

- 1. In Section 2.2, paragraph 5, the authors would like to add a new citation [54] after [53], and a sentence at the end of this paragraph, "A similar technique was used in [54] to define window boundaries for the general analysis of genomic data.".
- 2. In Figure 1, the authors would like to add a sentence at the end of the figure caption, "An analogous suite of displays is shown in [54] to present the partitioning of genomic data into windows.".

The authors would like to apologize for any inconvenience caused. The changes do not affect the scientific results. The manuscript will be updated, and the original will remain online on the article webpage, with a reference to this correction.

Conflicts of Interest: The authors declare no conflict of interest.

## Reference

 Ramzan, F.; Gültas, M.; Bertram, H.; Cavero, D.; Schmitt, A.O. Combining Random Forests and a Signal Detection Method Leads to the Robust Detection of Genotype-Phenotype Associations. *Genes* 2020, 11, 892. [CrossRef] [PubMed]

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