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Correction to: Genetic characterization of Addison's disease in Bearded Collies

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Following publication of the original article [1], it was noted that due to a typesetting error the caption of Fig. 1 was mistakenly captured within the main body of the article.

Furthermore, the layout of the online versions of Tables 1 and 2 have been updated to improve the presentation of the genotype comparisons.

The correct Fig. 1 with the caption has been included in this Correction article, and the original article has been updated.

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Reference
1.   Gershony LC, Belanger JM, Hytönen MK, et al. Genetic characterization of Addison's disease in Bearded Collies. BMC Genomics. 2020;21:833. https://doi.org/10.1186/s12864-020-07243-0.

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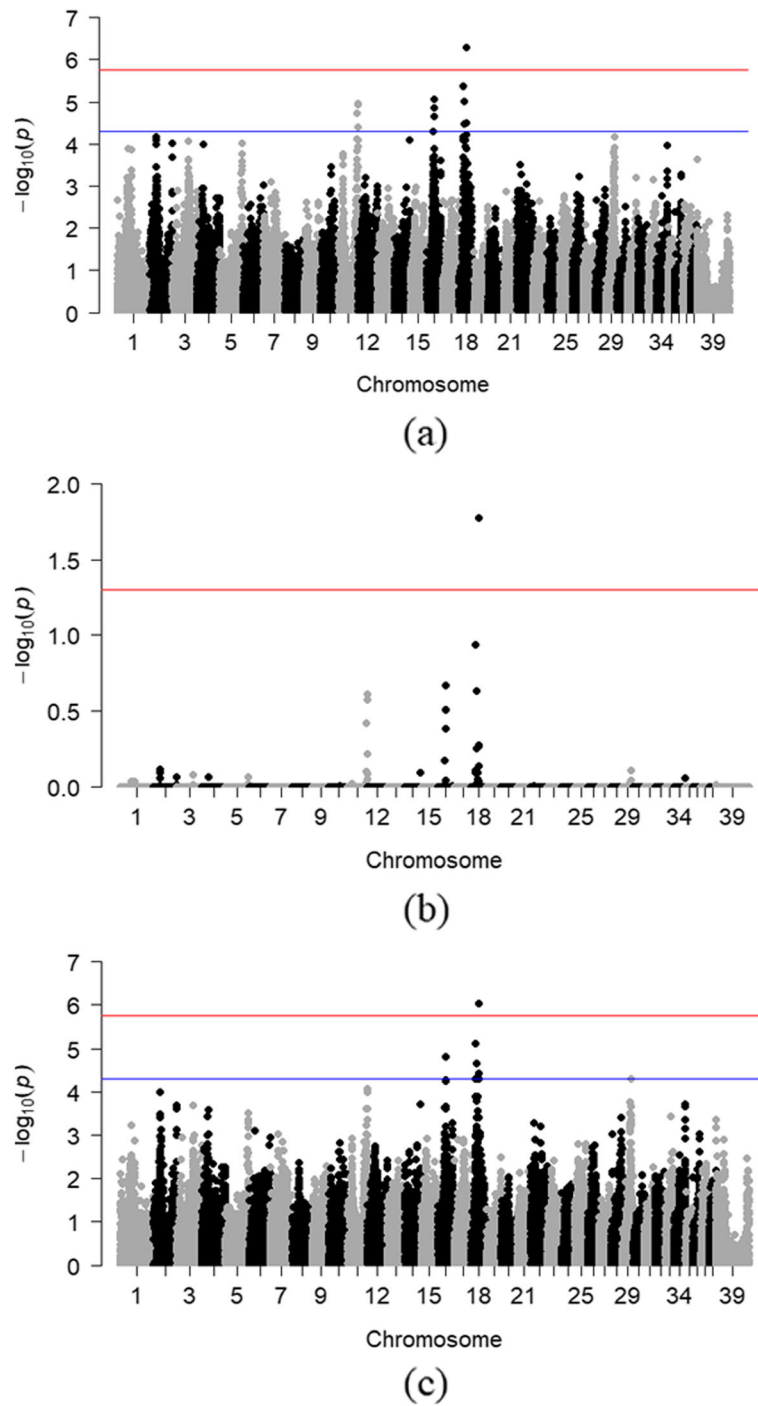Full list of author information is available at the end of the article

Fig. 1 Manhattan Plots. Chi-square based allelic association (**a**) and association testing after 100,000 max(T) permutations (**b**) in PLINK for 103 unrelated Bearded Collies (41 cases, 62 healthy controls; λGC = 1.2280). **c** Association testing using GEMMA's univariate linear mixed model approach to account for population substructure of the same dataset (λGC = 1.0328). The blue and red lines indicate suggestive (*p* < 0.00005; −log10[*p*-value] ≥ 4.3) and Bonferroni-adjusted genome-wide significance threshold (−log10[*p*-value] ≥ 5.75), respectively