

CORRECTION

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Correction to: Nationwide genetic testing towards eliminating Lafora disease from Miniature Wirehaired Dachshunds in the United Kingdom

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Following the publication of the original article [1], the authors identified an error in Fig. 1. The correct figure is given below and the original article has been corrected.

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Reference

1. Ahonen S, et al. Nationwide genetic testing towards eliminating Lafora disease from Miniature Wirehaired Dachshunds in the United Kingdom. *Canine Genet Epidemiol*. 2018;5:2.

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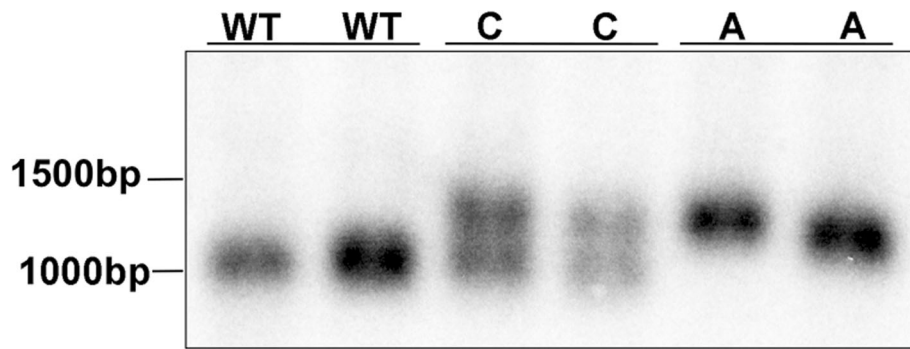


Fig. 1 Genotyping is performed using Southern blot. Southern blot is used for genotyping as PCR based methods cannot reliably distinguish the different genotypes due to the type of the mutation. The affected dogs (**a**) are homozygous for the dodecamer repeat expansion mutation with multiple dodecamer repeats, carrier (**c**) dogs have the normal and mutated allele and clear dogs (WT) have three copies of the repeat [8]