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Erratum to: Potential of promotion of alleles by genome editing to improve quantitative traits in livestock breeding programs

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After the publication of this work [1], we noticed that Figs. 1 and 2 were accidentally interchanged. The correct version order of Figs. 1 and 2 are provided here. The original article was corrected.

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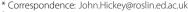
 Jenko J, Gorjanc G, Cleveland MA, Varshney RK, Whitelaw CBA, Woolliams JA, et al. Potential of promotion alleles by genome editing to improve quantitative traits in livestock breeding programs. Genet Sel Evol. 2015;47:55.

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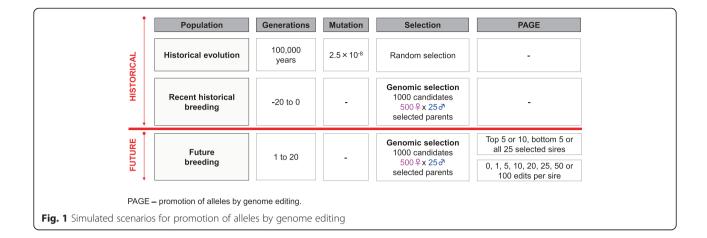
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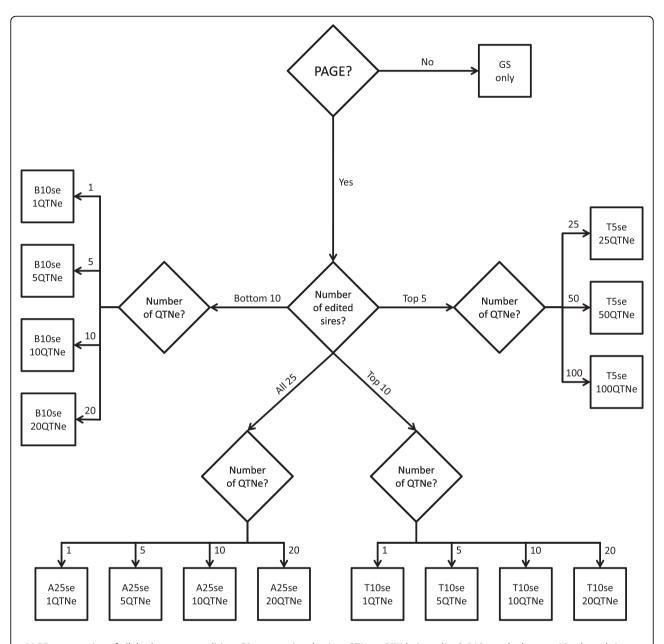




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PAGE – promotion of alleles by genome editing; GS – genomic selection; QTNe – QTN being edited; B10se – the bottom 10 selected sires edited; A25se – all of the 25 selected sires edited; T10se – the top 10 selected sires edited; T5se – the top 5 selected sires edited

Fig. 2 Overall design of the simulation