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Genome-wide association study identifies a putative variant in MYO3B and common variants in PTPRT associated with bovine tuberculosis

Citation for published version:

Bermingham, M, Genome-wide association study identifies a putative variant in MYO3B and common variants in PTPRT associated with bovine tuberculosis: Front cover of Heredity February 2014, 2014, Digital or Visual Products.

Link: Link to publication record in Edinburgh Research Explorer

Document Version: Preprint (usually an early version)

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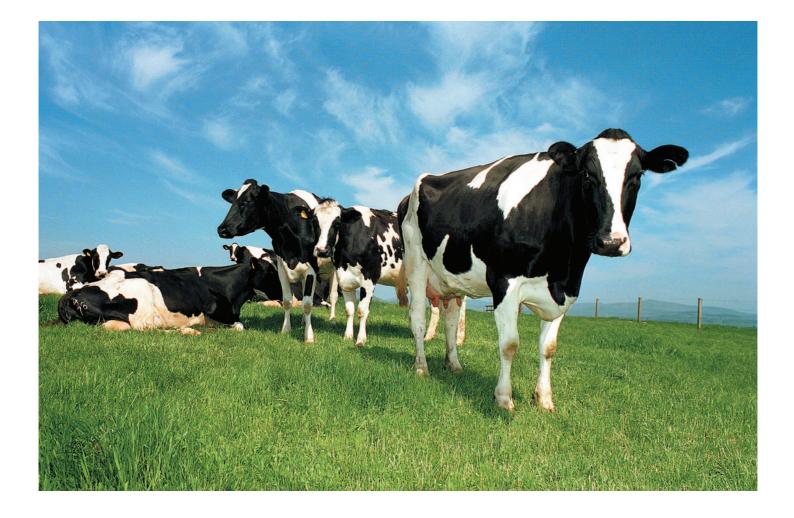
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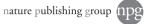


Volume 112 Number 5 May 2014 www.nature.com/hdy



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