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

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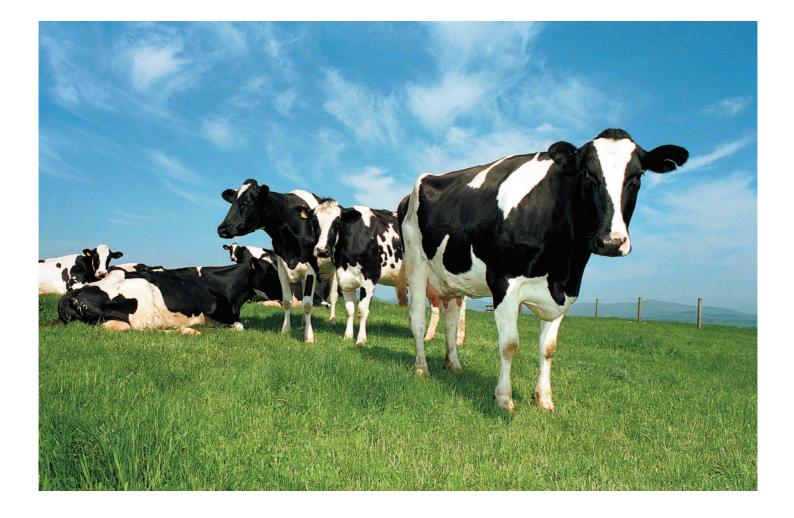
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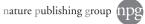


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