

*Supplementary Information*

Non-invasive genotyping and spatial mark-recapture methods to estimate European pine marten density in forested landscapes

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**Supplement S1** MMX and MMY probe sequences.

MMX and MMY probe sequences are reversed from the text provided in (Mullins et al., 2009) and are as follows: MMX, 5'-VIC-CCTGGTCTGAAAAC-T-MGB-3' and MMY 5'-6FAM-TGTGTCTCTCTGTCAAMGB-3'.