

Manuscript number... 2011-05-06400A
 Author name... Rayner
 Article type... Letter

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- Abbreviations and symbols defined on first use
- Italic font used for variables, genes and non-English languages; bold font for vectors
- Equations and special characters in Word in 'normal text' or Symbol font (Word Equation Editor avoided where possible)
- Authored in Compatibility Mode if using Word 2007 (i.e., as a Word 97-2003 document; saved as .doc format, NOT .docx)
- Brief title provided for each figure, table and box
- Figure legends no more than 100 words each (300 words each for contributions without methods sections)
- Author contribution statement included (a statement of responsibility that specifies the contribution of every author; see http://www.nature.com/authors/editorial_policies/authorship.html)
- Declaration and listing of any competing interests included in manuscript in the Author Information section (<http://www.nature.com/nature/authors/policy/competing.html>)
- Database accession numbers included for sequences determined in this paper
- Structure data in Nature standard tables

For TeX users

Word or rtf formats are preferred, unless there is very extensive use of special characters or equations. Such formats enable on-screen subediting and the preparation of a 'pre-proof', when the editing can be checked before formal proofs are prepared. We have found that this minimizes the chance of errors being inadvertently introduced.

- Word conversion not practicable

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- Separate text file ('SI Guide') provided with a brief description (file format and content) of each item
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Additional checklist for *Nature* Biology Letters

Manuscript number: 2011-05-06400A

Author name: Rayner et al.

Title: Inhibition of miR33a/b in non-human primates...

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| Formatting: | Yes |
|---|-----|
| The paper is within the 4 page length limit for Letters (please use the provided macro to help you determine the length of your paper). | ✓ |
| The methods summary is less than 300 words and the full methods are provided in the SAME word document as the main text, after the figure legends (the numbering of references used only in the full methods section continues from the main references). | ✓ |
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Please confirm that your manuscript meets each of the relevant criteria below, by ticking either "yes" or "not applicable".

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|---|-----|-----|
| All small RNA sequences are provided. | ✓ | |
| Microarray data are in MIAME format. | | ✓ |
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| <i>Nature</i> standard table has been used for X-ray crystallography/NMR structural data. | | ✓ |
| Accession numbers are provided for protein structure/cryoEM structure/ microarray/gene sequences/small RNA sequences. | ✓ | |