

Note:

This work is part of the FANTOM5 project. Data downloads, genomic tools and co-published manuscripts are summarized here <http://fantom.gsc.riken.jp/5/>.

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Article Type: Direct Submission; Collection Codes: , ,
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Publisher: PNAS; Journal: PNAS:Proceedings of the National Academy of Sciences;

Copyright: , ; Volume: 110; Issue: 0; Manuscript: 201312717; Month: ; Year: 2013

DOI: 1312717110; TOC Head: ; Section Head:

Article Type: Direct Submission; Collection Codes: , ,

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