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### Authors

Chen, Bin  
Harbach, Ralph E  
Walton, Catherine  
[et al.](#)

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### Erratum

## Erratum to “Population genetics of the malaria vector *Anopheles aconitus* in China and Southeast Asia” [Infect. Genet. Evol. 12 (2012) 1958–1967]



Bin Chen <sup>a,\*</sup>, Ralph E. Harbach <sup>b</sup>, Catherine Walton <sup>c</sup>, Zhengbo He <sup>a</sup>, Daibin Zhong <sup>d</sup>, Guiyun Yan <sup>d</sup>, Roger K. Butlin <sup>e</sup>

<sup>a</sup> Institute of Entomology and Molecular Biology, College of Life Sciences, Chongqing Normal University, Chongqing 401331, China

<sup>b</sup> Department of Entomology, The Natural History Museum, London SW7 5BD, UK

<sup>c</sup> Faculty of Life Sciences, The University of Manchester, Manchester M13 9PT, UK

<sup>d</sup> College of Health Sciences, University of California at Irvine, CA 92697, USA

<sup>e</sup> Animal and Plant Sciences, The University of Sheffield, Sheffield S10 2TN, UK

When this article was published, [Table 2](#) contained a number of errors. The correct artwork is printed below.

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\* Corresponding author at: Institute of Entomology and Molecular Biology, College of Life Sciences, Chongqing Normal University, Chongqing 401331, China. Tel.: +86 2365918391; fax: +86 2365340559.

E-mail address: [c\\_bin@hotmail.com](mailto:c_bin@hotmail.com) (.)

**Table 2**  
Variation in the 639 bp sequence of the *COII* gene of *Anopheles aconitus*.

Haplotype	Sites	GenBank accession no.
	1111222222333333333344444444445555555555 12234778567701345790122335590123667799111225677 67804251951410448976847034752400381756069282706	
HAPL1 (44)	GACACTTTCTACTAACTTAATTTATATATTTGGTATAGAAAATTTAAA	AJ626951
HAPL2 (20)	.....G.....G...C...	AJ626952
HAPL3 (10)	. . T. ....G.....G...C...	AJ626953
HAPL4 (7)	.....C.....G.....G...C...	AJ626954
HAPL5 (6)	.....T.....C.....G...C...	AJ626955
HAPL6 (5)	.....C.....G.....C..A.....G...C...	AJ626956
HAPL7 (1)	.....G.....G.....G.....G...C...	AJ626957
HAPL8 (1)	.....G.....G.....G.....G...CC...	AJ626958
HAPL9 (2)	.....G.....G.....G.....G...C...	AJ626959
HAPL10 (1)	.....C...T.....G.....G...C...	AJ626960
HAPL11 (1)	.....C.....G.....A.....G...C...	AJ626961
HAPL12 (1)	. . T. ....G.....G...C...	AJ626962
HAPL13 (1)	.....C.....G...C.....G...C...	AJ626963
HAPL14 (3)	.....G.....G.....G.....G...C...	AJ626964
HAPL15 (1)	.....C.....G...G.....C..A.....G...C...	AJ626965
HAPL16 (1)	.....T.C.....G.....C..A.....G...C...	AJ626966
HAPL17 (2)	.....T.....C..C.....G...C...	AJ626967
HAPL18 (2)	.....T.....G.....G.....G...C...	AJ626968
HAPL19 (1)	. . G. . . T. ....CC..G.....C...	AJ626969
HAPL20 (1)	.....C.....G.....G...C...	AJ626970
HAPL21 (1)	.....CCC.....G.....G...C...	AJ626971
HAPL22 (1)	.....G.....G.....G...C...	AJ626972
HAPL23 (1)	.....C.....G.....G...C...	AJ626973
HAPL24 (1)	A. ....C.....C.....G...C...	AJ626974
HAPL25 (1)	. G. ....G.....G...C...	AJ626975
HAPL26 (1)	. G. ....C.....G.....G...C...	AJ626976
HAPL27 (1)	.....G..C.....G...C...	AJ626977
HAPL28 (1)	.....CG.....G.....G...C...	AJ626978
HAPL29 (2)	.....G.....C..G...C...	HQ403792. 1
HAPL30 (1)	.....C.....G.....G...C...	HQ403794. 1
HAPL31 (1)	.....G.....G.....G...C.G.	HQ403776. 1
HAPL32 (1)	.....G.....G.....G...G.	HQ403771. 1
HAPL33 (1)	.....A.....G.....G...G.	HQ403765. 1
HAPL34 (1)	. GT. ....G.....G.....G...C...	HQ403757. 1
HAPL35 (1)	. . T. ....G.....A.G...C...	HQ403772. 1
HAPL36 (3)	.....G.C.....G.....G...G.	HQ403808. 1
HAPL37 (1)	.....C.....C..A.....GGG..C...	HQ403804. 1
HAPL38 (1)	. G. ....C.....G.....G...C...	HQ403812. 1
HAPL39 (1)	.....G.G.....G.....G...G.	HQ403799. 1
HAPL40 (1)	.....G.....G.....G...G.	HQ403795. 1
HAPL41 (1)	.....T.....C.....G.....G...G.	HQ403796. 1
HAPL42 (1)	.....C.....C.....G.....G...CG..	HQ403800. 1
HAPL43 (1)	.....C.....C.....G.....G...C...	HQ403802. 1
HAPL44 (1)	. . T. ....C.....C.....G.....G...G.	HQ403806. 1
HAPL45 (1)	.....C.....G.....G.....G...G.	HQ403807. 1
HAPL46 (1)	.....G.....G.....G...T	HQ403814. 1

Frequencies of haplotypes are in parentheses and variable positions are shown in the first row. Identity with the first sequence is denoted by a dot and substitution by a different base letter.