

CORRECTION

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Correction: Transcriptome analysis and identification of chemosensory genes in the larvae of *Plagioder a versicolora*

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Correction: *BMC Genomics* 23, 845 (2022)
<https://doi.org/10.1186/s12864-022-09079-2>

Following the publication of the original article [1], it was noted that the figure legends were paired incorrectly. The figure legends for Figs. 2 and 3 were wrongly given as captions for Figs. 3 and 2 respectively.

The correct figures and captions have been included in this correction, and the original article has been corrected.

Reference

1. Wu ZR, Fan JT, Tong N, et al. Transcriptome analysis and identification of chemosensory genes in the larvae of *Plagioder a versicolora*. *BMC Genomics*. 2022;23:845. <https://doi.org/10.1186/s12864-022-09079-2>.

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		C1	C2	
PverCSP3MGVCTIFLICLVGLVLAQPDEKYYTTKYDNLIDITLTKSERLLKNYVHCLLCKGKIPDGAEL	C	C	63
PverCSP5MFSLFILSCFVIGSLSTVTEKTYTTKYDNINVEDIVKNDRLIKNYVDCLLERGGKCTPDGLEL	C	C	63
PverCSP9MKSIIIVCFVVLASVIARPDSSKYTSKYDNVLDLAIHNDRLIRNYVDCLLGKKKCKDGEEL	C	C	64
PverCSP11MFLSFLFCVIVVSLVEGCKYQSKYDNIIVDSILTNRRVLLKSVLRGILLGEGFCPPAREF	C	C	61
PverCSP12	MKRLLSASVLFVVLAVLATDRKERSKVAGYTTKYDNVLDLAIIRNERILIRNYVDCLLGKKKCTPDGEVL	C	C	70
	C3 C4			
PverCSP3	KEHLFDALLTDCSKCSEFCKKNSKKIIRYLIDNKPDPWYKELEVRYDPEGTYKAKYDEEIKDKAAE....	C	C	128
PverCSP5	KKNMFDALIEDCSKCEKCKEGSEFMMRYLIDNKPAYWELQERYDPSGSYKRYLDKAKTEVNIQPIVK	C	C	133
PverCSP9	KRILFDALKTKCAKCESEFKNNAKKVINHLIKNCRAWWELEAVYDSEGYRKEYEAEAKKEGIDLN...	C	C	131
PverCSP11	RTHIFDALSNRCARCTDAQVNIILKTSKIFMRNPNEDWEKISKKFDQPKYRASFTKFLNA.....	C	C	122
PverCSP12	KGHIKDAIENECDKCSLVCKKSVRVKGTIYQKHPTWWKELCDHFDPEAKYQKRYEKFIQDALKEVE...	C	C	137

Fig. 2 Multiple amino alignments of the predicted CSPs. Conserved cysteine (C1-C4) residues are marked with a blue box

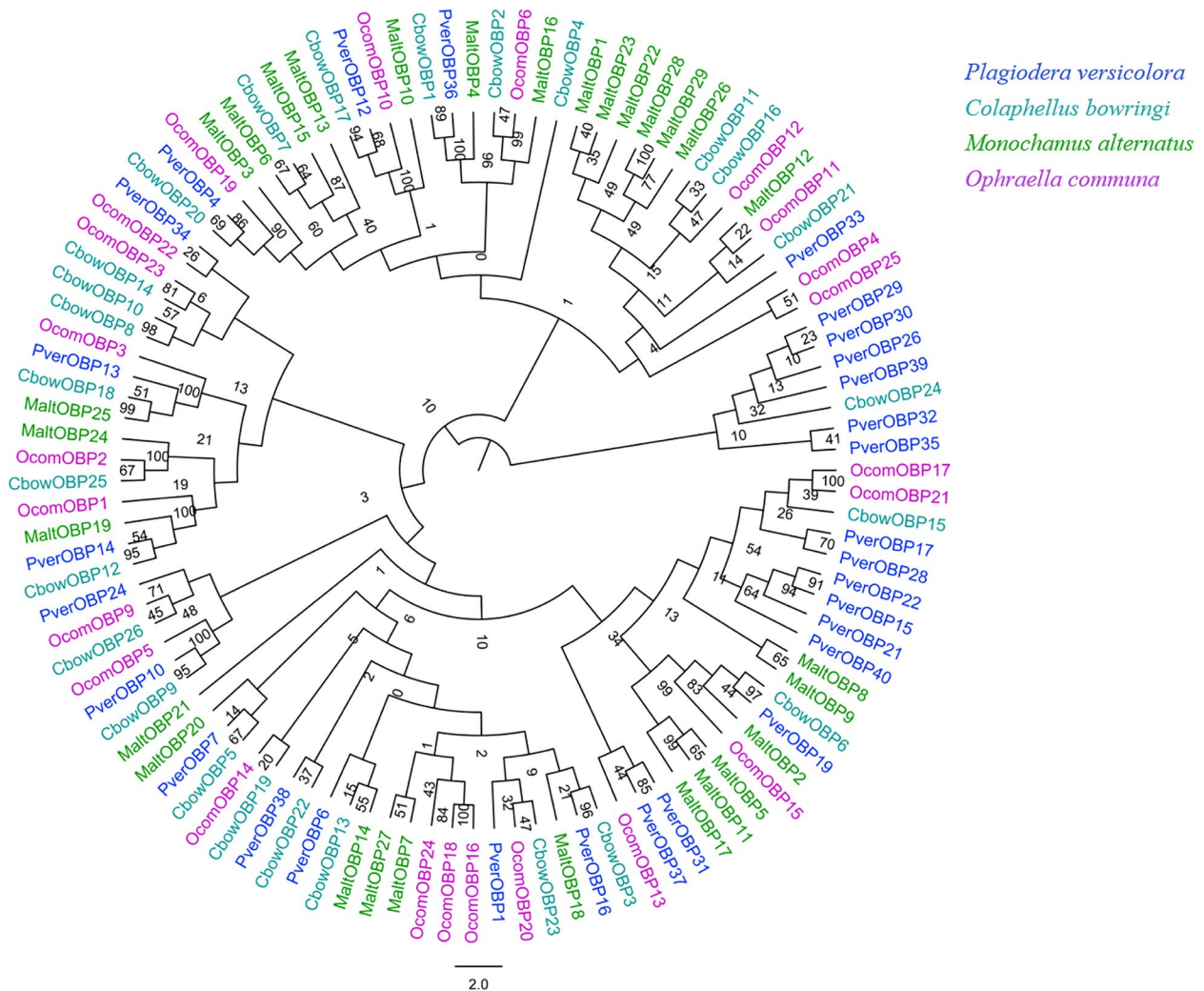


Fig. 3 Phylogenetic analysis of the OBPs (odorant-binding proteins) from four insect species: *P. versicolora* (Pver), *C. bowringi* (Cbow), *M. alternatus* (Malt), *O. communa* (Ocom). The *P. versicolora* genes are shown in blue. The values at the branch nodes represent bootstrap values based on 1000 replicates