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### **Title**

Correction: Spodoptera frugiperda Smith (Lepidoptera: Noctuidae) in Cameroon: Case study on its distribution, damage, pesticide use, genetic differentiation and host plants

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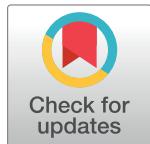
Peer reviewed

## CORRECTION

# Correction: *Spodoptera frugiperda* Smith (Lepidoptera: Noctuidae) in Cameroon: Case study on its distribution, damage, pesticide use, genetic differentiation and host plants

**Apollin Fotso Kuate, Rachid Hanna, Armand R. P. Doumtsop Fotio, Albert Fomumbod Abang, Samuel Nanga Nanga, Sergine Ngatat, Maurice Tindo, Cargele Masso, Rose Ndembah, Christopher Suh, Komi Kouma Mokpokpo Fiaboe**

The images for Figs 3 and 4 are incorrectly switched. The image that appears as Fig 3 should be Fig 4, and the image that appears as Fig 4 should be Fig 3. The figure captions appear in the correct order.



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## OPEN ACCESS

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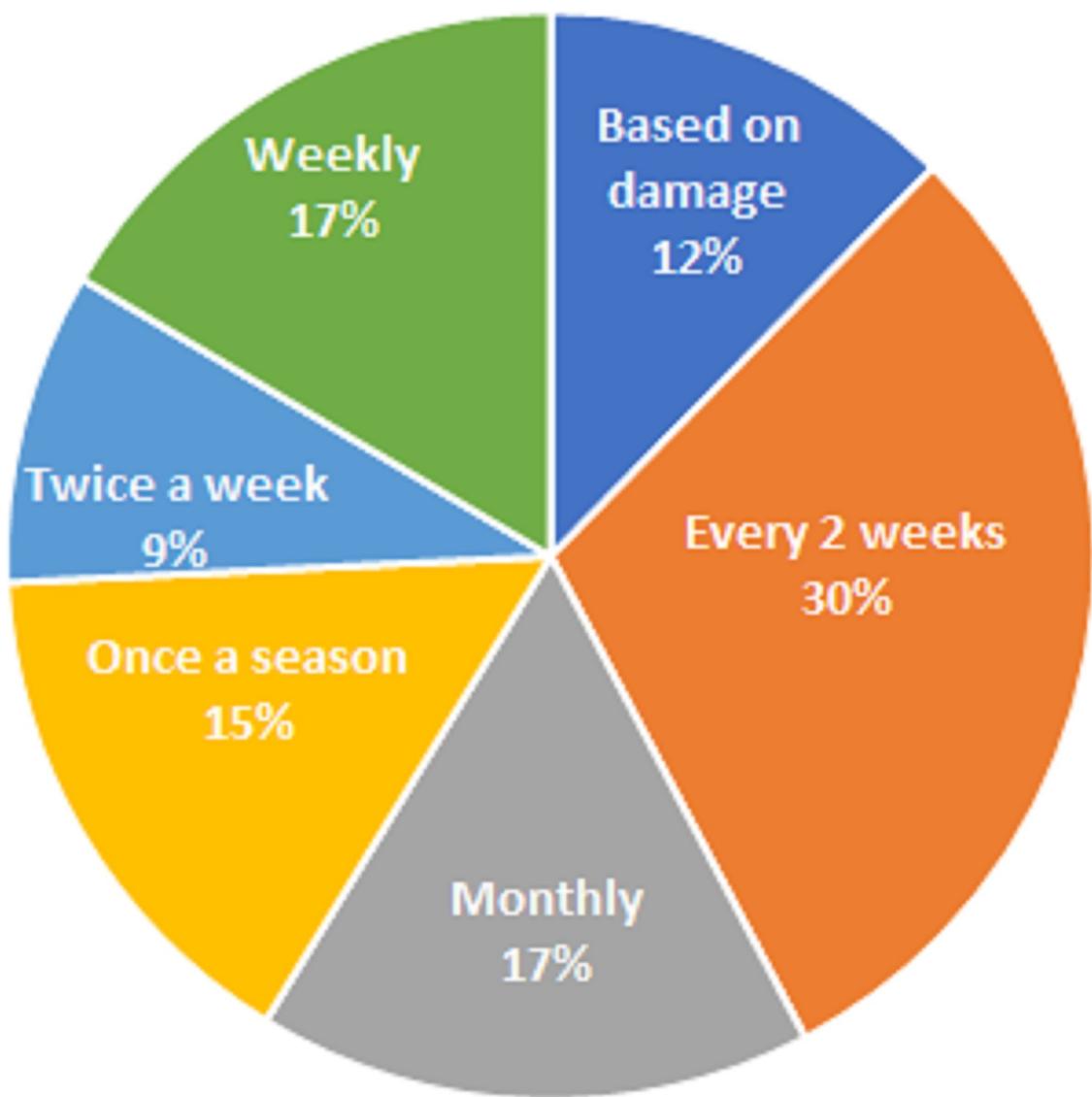


Fig 3. Pesticide application frequency to control *S. frugiperda* on maize by farmers.

<https://doi.org/10.1371/journal.pone.0217653.g001>

FAW Consensus sequence	1	70
FAW Haplotype 2 (corn race)	AACATTATTTTATTTGGAAATTGAGCAGGRATAGTAGGTACTCTTAAGTTATTAATTCGAGCT	
FAW Haplotype 1 (rice race)	.....G.....	
	.....A.....	
71		140
FAW Consensus sequence	GAATTAGGRRACTCCAGGATCTTAATTGGAGATGATCAAATTATAACTATTGTAACAGCYCATGCCT	
FAW Haplotype 2 (corn race)	.....G.....T.....	
FAW Haplotype 1 (rice race)	.....A.....C.....	
141		210
FAW Consensus sequence	TTATTATAATTTTTATAGTTACCCWATTATAATTGGAGGATTGAAATTGACTGTACCTTAAT	
FAW Haplotype 2 (corn race)	.....T.....	
FAW Haplotype 1 (rice race)	.....A.....	
211		280
FAW Consensus sequence	ATTAGGAGCYCCTGATATAGCTTCCCACGTATAAATAATAGTTTGACTTTACCCCCATCTTA	
FAW Haplotype 2 (corn race)	.....C.....	
FAW Haplotype 1 (rice race)	.....T.....	
281		350
FAW Consensus sequence	ACTTTATTAATTTCTAGTAGCATTGAGAAAATGGAGCAGGAACCTGGATGAACAGTTACCCCCCTCT	
FAW Haplotype 2 (corn race)	.....	
FAW Haplotype 1 (rice race)	.....	
351		420
FAW Consensus sequence	CCTCTAATATTGCTCATGGGGTAGTCAGTAGATTAGCTATTTCTACTTCATTAGCTGGAATTTC	
FAW Haplotype 2 (corn race)	.....	
FAW Haplotype 1 (rice race)	.....	
421		490
FAW Consensus sequence	ATCTATTTAGGAGCTATTAACTTATTACYACTATTATAATACGATTAATAATTATCATTTGAT	
FAW Haplotype 2 (corn race)	.....T.....	
FAW Haplotype 1 (rice race)	.....C.....	
491		560
FAW Consensus sequence	CAAATACCTTATTTATTGAGCTGTAGGTATTACYGCATTYTTATTATTATCTTACCTGTTAG	
FAW Haplotype 2 (corn race)	.....T.....C.....	
FAW Haplotype 1 (rice race)	.....C.....T.....	
561		630
FAW Consensus sequence	CYGGAGCTATTACTATATTACTGATCGAAATYAAATACATCATTTCGATCCTGCAGGWGGAGG	
FAW Haplotype 2 (corn race)	.C.....T.....T.....	
FAW Haplotype 1 (rice race)	.T.....C.....A.....	
631	658	
FAW Consensus sequence	TGATCCTATTCTTATCACACATTATT	
FAW Haplotype 2 (corn race)	.....	
FAW Haplotype 1 (rice race)	.....	

Fig 4. Identity plot of the COI sequences of *S. frugiperda* haplotypes from Cameroon.<https://doi.org/10.1371/journal.pone.0217653.g002>

## Reference

1. Foto Kuate A, Hanna R, Doumtsop Fotio ARP, Abang AF, Nanga SN, Ngatat S, et al. (2019) *Spodoptera frugiperda* Smith (Lepidoptera: Noctuidae) in Cameroon: Case study on its distribution, damage, pesticide use, genetic differentiation and host plants. PLoS ONE 14(4): e0215749. <https://doi.org/10.1371/journal.pone.0215749> PMID: 31034480