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Step 1

Explore the different types of open access publishing, and select the type that works best for you and your institution and/or funder

Step 2

Research the best home for your paper

Step 3

Make sure you understand publication fees and your funding options

Step 4

Ensure you understand any open access mandates, and how to comply with funder and institutional requirements

Step 5

Select your licensing and copyright preference

Step 6

Ensure you meet journal requirements for making your data available

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- The most well-known subject-based repository for research is PubMed Central (PMC) and sister sites such as Europe PMC.

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Quick reference guides: to help you as you publish

Gold open access

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- Payment is usually taken after acceptance and before publication.

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Step 4

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Q Search WILEY . Online Library WILEY Login / Register Volume 13, Issue 11 **Ecology and Evolution** November 2023 Open Acce e10660 RESEARCH ARTICLE 🔂 Open Access 00 Integrated analysis of miRNA profiles and gut bacterial changes in Altica viridicyanea following antibiotic treatment Advertisement Yipeng Ren, Yuan Wang, Juhong Chen, Siying Fu, Wenjun Bu 🔀, Huaijun Xue 🔀 WILEY First published: 31 October 2023 | https://doi.org/10.1002/ece3.10660 Ecology and Evolution ∃ SECTIONS 🍸 PDF 🔧 TOOLS < SHARE **Call for Papers** Open science hallenge innovations & future Abstract directions seeks papers for an upcoming Special issue The gut bacteria involves in insect homeostasis by playing essential roles in host physiology, metabolism, innate immunity, and so forth. microRNAs (miRNAs) are mission Deadlin endogenous small noncoding RNAs that posttranscriptionally regulate gene expression to affect immune or metabolic processes in insects. For several non-model insects, the available knowledge on the relationship between changes in the gut bacteria and miRNA profiles is limited. In this study, we investigated the gut bacterial diversity, composition, Information ires References Related and function from Altica viridicyanea feeding on normal- and antibiotic-treated host plants using 16S rRNA amplicon sequencing; antibiotics have been shown to affect the body Metrics weight and development time in A. viridicyanea, suggesting that the gut bacteria of the Full text views: 87 () normal sample were more diverse and abundant than those of the antibiotic-fed group, and most of them were involved in various physical functions by enrichment analysis. Furthermore, we executed small RNA transcriptome sequencing using the two Details experimental groups to obtain numerous sRNAs, such as piRNAs, siRNAs, and known and © 2023 The Authors. Ecology and Evolution novel miRNAs, by data mapping and quality control, and furthermore, a total of 224 published by John Wiley & Sons Ltd. miRNAs were identified as significantly differentially expressed miRNAs, of which som This is an open access article under the terms of DEMs and their target genes participated in immune- and metabolism-related pathway the Creative Commons Attribution License, which based on GO and KEGG annotation. Besides, regarding the regulatory roles of miRNA and permits use, distribution and reproduction in any target genes, a interaction network of DEM-target gene pairs from eight immune- or medium, provided the original work is properly cited. metabolism-related signaling pathways were constructed. Finally, we discovered that DEMs from above pathways were significantly positively or negatively correlated with gut Check for updates bacterial alterations following antibiotic treatment. Collectively, the observations of this study expand our understanding of how the disturbance of gut bacteria affects miRNA profiles in A. viridicyanea and provide new valuable resources from extreme ranges for **Research Funding** future studies on the adaptive evolution in insects. Fundamental Research Funds for the Central Universities. Grant Number: **1 INTRODUCTION** 63213120 microRNAs (miRNAs), serving as important regulators of gene expression, are a type of small National Natural Science Foundation of

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