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PEER-REVIEW REPORT

Name of journal: World Journal of Gastroenterology

Manuscript NO: 75763

Title: Associations of gut microbiota with dyslipidemia based on sex differences in

subjects from Northwestern China

Provenance and peer review: Invited Manuscript; Externally peer reviewed

Peer-review model: Single blind

Reviewer's code: 02537353 Position: Editorial Board Academic degree: BSc

Professional title: Associate Professor

Reviewer's Country/Territory: Italy

Author's Country/Territory: China

Manuscript submission date: 2022-02-18

Reviewer chosen by: AI Technique

Reviewer accepted review: 2022-02-18 07:46

Reviewer performed review: 2022-02-23 12:55

Review time: 5 Days and 5 Hours

Scientific quality	[] Grade A: Excellent [] Grade B: Very good [] Grade C: Good [Y] Grade D: Fair [] Grade E: Do not publish
Language quality	[] Grade A: Priority publishing [Y] Grade B: Minor language polishing [] Grade C: A great deal of language polishing [] Grade D: Rejection
Conclusion	[] Accept (High priority) [] Accept (General priority) [] Minor revision [Y] Major revision [] Rejection
Re-review	[]Yes [Y]No



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Peer-reviewer

Peer-Review: [] Anonymous [Y] Onymous

statements

Conflicts-of-Interest: [] Yes [Y] No

SPECIFIC COMMENTS TO AUTHORS

The aim of the authors was to investigate the associations of GM characteristics with serum lipid profiles under sexual dimorphism in a Chinese population. The topic could be interesting but the manuscript presents different critical points Major revisions 1) The authors use the characteritazion of metabolic profile of microbiota the dated database FAPROTAX. Actually there is the PICrust 2 that makes predictions starting from the reads themselves and this brings the prediction closer to reality. and rightly to define the metabolic profile the microbiota studies evaluate the fecal SCFA, why the authors did not define the SCFA signature in stool of patients discussion is very long and verbose, please short it and focused on very important points

4) The number of enrolled patients is low for the conclusions of the authors 1) Please specify in the section Study Design "206 adult individuals " the number of patients and Healthy controls. In addition, please define how you have enrolled the Healthy controls 2) Please replace subjects with patients



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Peer-review model: Single blind

Reviewer's code: 02623966 **Position:** Editorial Board

Academic degree: MD, MSc, PhD

Professional title: Attending Doctor, Research Scientist

Reviewer's Country/Territory: Greece

Author's Country/Territory: China

Manuscript submission date: 2022-02-18

Reviewer chosen by: Xin Liu

Reviewer accepted review: 2022-04-13 12:15

Reviewer performed review: 2022-04-13 12:16

Review time: 1 Hour

Scientific quality	[] Grade A: Excellent [Y] Grade B: Very good [] Grade C: Good [] Grade D: Fair [] Grade E: Do not publish
Language quality	[] Grade A: Priority publishing [Y] Grade B: Minor language polishing [] Grade C: A great deal of language polishing [] Grade D: Rejection
Conclusion	[] Accept (High priority) [Y] Accept (General priority) [] Minor revision [] Major revision [] Rejection
Re-review	[Y]Yes []No



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Peer-reviewer

Peer-Review: [] Anonymous [Y] Onymous

statements Conflicts-of-Interest: [] Yes [Y] No

SPECIFIC COMMENTS TO AUTHORS

It is an interesting manuscript. Authors succeed to present their data in a clear way adding information to the existing literature. Therefore, I have no corrections to do and the manuscript can be published unaltered.



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Manuscript NO: 75763

Title: Associations of gut microbiota with dyslipidemia based on sex differences in

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Provenance and peer review: Invited Manuscript; Externally peer reviewed

Peer-review model: Single blind

Reviewer's code: 00053786 **Position:** Editorial Board

Academic degree: FAASLD, MD, PhD

Professional title: Research Scientist

Reviewer's Country/Territory: Mexico

Author's Country/Territory: China

Manuscript submission date: 2022-02-18

Reviewer chosen by: Xin Liu

Reviewer accepted review: 2022-04-11 13:26

Reviewer performed review: 2022-04-20 16:45

Review time: 9 Days and 3 Hours

Scientific quality	[] Grade A: Excellent [Y] Grade B: Very good [] Grade C: Good [] Grade D: Fair [] Grade E: Do not publish
Language quality	[] Grade A: Priority publishing [Y] Grade B: Minor language polishing [] Grade C: A great deal of language polishing [] Grade D: Rejection
Conclusion	[] Accept (High priority) [Y] Accept (General priority) [] Minor revision [] Major revision [] Rejection
Re-review	[Y]Yes []No



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Peer-reviewer statements

Peer-Review: [Y] Anonymous [] Onymous

Conflicts-of-Interest: [] Yes [Y] No

SPECIFIC COMMENTS TO AUTHORS

Guo L et al. present a retrospective study concerning the association between gut microbiota and dyslipidemia in male and female subjects from Northwestern China. The study may be of interest to the medical-hepatologist community in regards to the sex differences shown in the features of the gut microbiota community and dyslipidemia. However, there are several concerns that need to be amended. 1) The term sex dimorphism is not correct. Commonly, sex dimorphism is about traits that are only seen in one sex but not in the other. In general, any dyslipidemia was not exclusive to one sex or the other, and dysbiosis is present in both sexes. Therefore, the title and other parts of the text should read: sex differences or gender differences where the term sex dimorphism was originally written. For example, the new title can be: Sex differences in the association of gut microbiota with dyslipidemia in subjects from Northwestern China. 2) a) Add the time of patient recruitment when this study was carried out. b) The total number of patients was 142 subjects, 81 subjects and 61 controls. Please clarify methodology section 3) Discussion: In regards to the this data in the text of the functional analysis of the metabolic pathways that were differentially activated in the study patients, it would be of great value that the authors discuss the influence of diet in the subjects vs controls, and between genders because diet composition influences GM diversity. These findings have implications for the management of obesity-related chronic diseases. Can the authors add some information about the local diet or changes in the local diet that may lead to the prevalence of dyslipidemia?