List of revisions in Manuscript ID: 88568

Title: Circulating MicroRNA Expression and Nonalcoholic Fatty Liver Disease in Adolescents with Severe Obesity

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All changes in the manuscript are highlighted in yellow, both, in the Table and below and in the manuscript. All line numbers refer to the final submitted version of the manuscript.

Editor's:

I have reviewed the Peer-Review Report, full text of the manuscript, and the relevant ethics documents, all of which have met the basic publishing requirements of the World Journal of Clinical Cases, and the manuscript is conditionally accepted. I have sent the manuscript to the author(s) for its revision according to the Peer-Review Report, Editorial Office's comments and the Criteria for Manuscript Revision by Authors. Before its final acceptance, please provide and upload the following important documents: Biostatistics Review Certificate, a statement affirming that the statistical review of the study was performed by a biomedical statistician; Institutional Review Board Approval Form or Document, the primary version (PDF) of the Institutional Review Board's official approval, prepared in the official language of the authors' country; Signed Informed Consent Form(s) or Document(s), the primary version (PDF) of the Informed Consent Form that has been signed by all subjects and investigators of the study, prepared in the official language of the authors' country; STROBE Statement, an important document related to manuscript writing of observational/case control/retrospective cohort studies. Before final acceptance, uniform presentation should be used for figures showing the same or similar contents; for example, "Figure 1Pathological changes of atrophic gastritis after treatment. A: ...; B: ...; D: ...; D: ...; E: ...; F: ...; F: ...; G: ...". Please provide decomposable Figures (in which all components are movable and editable), organize them into a single PowerPoint file. Please authors are required to provide standard three-line tables, that is, only the top line, bottom line, and column line are displayed, while other table lines are hidden. The contents of each cell in the table should conform to the editing specifications, and the lines of each row or column of the table should be aligned. Do not use carriage returns or spaces to replace lines or vertical lines and do not segment cell content. Please check and confirm whether the figures are original (i.e. generated de novo by the author(s) for this paper). If the picture is 'original', the author needs to add the following copyright information to the bottom righthand side of the picture in PowerPoint (PPT): Copyright ©The Author(s) 2023. Before final acceptance, when revising the manuscript, the author must supplement and improve the highlights of the latest cutting-edge research results, thereby further improving the content of the manuscript. To this end, authors are advised to apply a new tool, the Reference Citation Analysis (RCA). RCA is an artificial intelligence technology-based open multidisciplinary citation analysis database. In it, upon obtaining search results from the keywords entered by the author, "Impact Index Per Article" under "Ranked by" should be selected to find the latest highlight articles, which can then be used to further improve an article under preparation/peer-review/revision. Please visit our RCA database for more information at: https://www.referencecitationanalysis.com/.

#	Editor's Comment	Requirement	Response
1	Biostatistics Review Certificate	A statement affirming that the statistical review of the study was performed by a biomedical statistician.	Thank you for this information. The required document is provided.
2	Institutional Review Board Approval Form or Document.	The primary version (PDF) of the Institutional Review Board's official approval, prepared in the official language of the authors' country.	Thank you for this information. Our research (manuscript ID: 88568) is secondary analysis of de-identified data from the Teen-LABS study, which means we don't have Institutional Review Board Approval Form or Signed Informed Consent Form. Instead, we can provide a letter of IRB approvement, and informed consent statement.
3	Signed Informed Consent Form(s) or Document(s), the primary version (PDF) of the Informed Consent Form	The primary version (PDF) of the Informed Consent Form that has been signed by all subjects and investigators of the study, prepared in the official language of the authors' country.	Thank you for this information. Our research (manuscript ID: 88568) is secondary analysis of de-identified data from the Teen-LABS study, which means we don't have Institutional Review Board Approval Form or Signed Informed Consent Form. Instead, we can provide a letter of IRB approvement, and informed consent statement.
4	STROBE Statement	An important document related to manuscript writing of observational/case control/retrospective cohort studies.	Thank you for this information. The required document is provided.
5	Uniform figure presentation	Uniform presentation should be used for figures showing the same or similar contents; for example, "Figure 1Pathological changes of atrophic gastritis after treatment. A:; B:; C:; D:; E:; F:; G:". Please provide decomposable Figures (in which all components are movable and editable), organize them into a single PowerPoint file.	Thank you for this information, we revised our figures accordingly. We additionally provided a PowerPoint file for your convenience.
6	Table format	Standard three-line tables, that is, only the top line, bottom	Thank you for this information, we

		line, and column line are displayed, while other table lines	revised our tables accordingly.
		are hidden;	
		The contents of each cell in the table should conform to the	
		editing specifications, and the lines of each row or column	
		of the table should be aligned;	
		Do not use carriage returns or spaces to replace lines or	
		vertical lines and do not segment cell content.	
7	Copyright of figures	Please check and confirm whether the figures are original	Thank you for acknowledging this
		(i.e. generated de novo by the author(s) for this paper). If	information. These figures are original
		the picture is 'original', the author needs to add the	and copyright information are added.
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Reviewer 1:

So far, many studies have reported the relationship between microRNA and NAFLD, but there is poor consistency in the global published research and evaluation of human liver miRNA expression. There is limited research on the human liver. This study identified new circulating miRNAs and analyzed their expression in different pathological features of NAFLD, which have mechanisms to promote or alleviate the progression of NAFLD. This is a new exploration and has good innovation. This study is of great significance for the diagnosis and treatment of NAFLD. As a contributor to the pathogenesis of human NAFLD, novel miRNAs are expected to serve as biomarkers for the noninvasive diagnosis and staging of NAFLD or hepatocellular carcinoma, or as targets for drug therapy, thereby preventing or reversing disease progression. The novel miRNA discovered in this study provides a new direction for targeted therapy of NAFLD. Due to the different types and quantities of miRNA expression at different stages of NAFLD, as well as differences in gender and whether obesity is present (such as obesity with NAFLD or lean individuals with NAFLD), further research is needed. Due to the relatively small sample size of this study, gender stratification studies were not conducted. Additionally, it would be better if specific miRNAs that reflect disease progression or deterioration could be identified.

#	Reviewers Comment	Response	Changes in manuscript
1	Due to the relatively small sample size of this study, gender stratification studies were not conducted.	Thank you for this comment. We acknowledge that the limited sample size in our study, with a notable disproportion in gender distribution. Specifically, we only had 36 male participants. As a result, we did not conduct stratification analysis by sex. We expanded our discussion of sample size issue in manuscript.	Discussion: Line: 464-467 Given our specific focus on adolescents with obesity, who are at high risk of developing NAFLD ^{(Loomis et al., 2016; Marchesini et al., 2003; Xanthakos et al., ²⁰¹⁵⁾, and the limitations arising from our small sample size, a validation for generalizability is imperative. The limitation in sample size is an inherent consequence of our methodological choice to employ liver biopsy for NAFLD diagnosis. While liver biopsy ensures accurate and definitive diagnosis, its invasiveness and cost present challenges in expanding the participant pool^(Spengler & Loomba, 2015). Studies with larger and more diverse populations would facilitate more robust and conclusive findings regarding NAFLD-miRNA associations.}
2	Additionally, it would be better if specific miRNAs that reflect disease progression or deterioration could be identified.	Thank you for your valuable feedback and suggestions. We added a discussion of miRNA that may reflect NAFLD progressions.	Discussion: Line: 386-394: Additionally, our analysis revealed decreased levels of miR-1296-5p, miR-1301-5p, miR-199b-5p, miR-411-5p, and miR-6885-3p in NASH compared

	to NAFL, while these levels were elevated in
	NAFLD compared to individuals without
	NAFLD. These miRNAs also demonstrated
	predominantly negative associations with fibrosis,
	lobular inflammation and ballooning (Figure 2).
	Notably, the downregulation of miR-411-5p
	aligned with a recent study by Wan et al.(Wan et
	al., 2022), which reported decreased expression of
	serum miR-411-5p in persons with NASH.
	Collectively, the distinct expression patterns
	observed across various NAFLD features suggest
	that these miRNAs may serve as potential
	biomarkers for NAFLD progression.

Reviewer 2:

Specific Comments to Authors: This study indicates the differential expression of circulating miRNAs in adolescent NAFLD, suggesting that they may become diagnostic and prognostic biomarkers for NAFLD. However, there are two shortcomings. Firstly, the sample size is small and research needs to be conducted in a larger and more diverse populations. The second issue is that there has been no molecular mechanism validation of differentially expressed circulating miRNAs through cytology or animal experiments.

#	Reviewers Comment	Response	Changes in manuscript
1	Firstly, the sample size is small and research needs	Thank you for your comment. We expanded our	Discussion:
	to be conducted in a larger and more diverse populations.	discussion of sample size limitations in this study.	Line: 464-467
			Given our specific focus on adolescents with
			obesity, who are at high risk of developing NAFLD ^{(Loomis et al., 2016; Marchesini et al., 2003; Xanthakos et al.,}
			²⁰¹⁵⁾ , and the limitations arising from our small
			sample size, a validation for generalizability is
			imperative. The limitation in sample size is an
			inherent consequence of our methodological
			choice to employ liver biopsy for NAFLD
			diagnosis. While liver biopsy ensures accurate and
			definitive diagnosis, its invasiveness and cost
			present challenges in expanding the participant
			pool ^(Spengler & Loomba, 2015) . Studies with larger and
			more diverse populations would facilitate more
			robust and conclusive findings regarding
			NAFLD-miRNA associations.

2	The second issue is that there has been no	Thank you for your insightful comment and	Result:
	molecular mechanism validation of differentially	suggestion. While we acknowledge this limitation,	Table 2
	expressed circulating miRNAs through cytology	we reviewed experimental studies and expanded	Table 5.
	or animal experiments.	our discussion on potential mechanisms	Discussion:
		associated with the identified miRNAs in NAFLD.	Line: 307 300
		Additionally, we have provided a summary of	Line. 397-399
		current findings on miRNA pathways in NAFLD	An increasing body of research have investigated
		based on experimental research, presented in	the associations between NAFLD and miRNA
		Table 3.	expression, yet little is known about mechanisms
			underlying the dysregulation of circulating
			miRNA in NAFLD patients.
			Line: 409-434
			Experimental studies provide valuable
			insights into the molecular mechanisms
			underlying associations between NAFLD and
			miRNA expression while minimizing
			confounding variables intrinsic to human
			observational studies. Particularly, miR-122, a
			highly expressed hepatic miRNA in hepatocytes,
			is associated with NAFLD progression by
			regulating lipid metabolism ^[23] . For example, Long
			et al. ^[74] revealed that miR-122 inhibited liver
			kinase B1/AMP-activated protein kinase signaling
			pathway, which further induced hepatic
			lipogenesis and steatosis in NAFLD. Additionally,
			the inhibition of miR-122-5p may suppress the
			inflammation and oxidative stress damage in
			NAFLD ^[75] . Given the observed upregulation of
			circulating miR-122 and downregulation of
			hepatic miR-122 in NASH patients ^[30, 36] , the
			elevated circulating miR-122-5p across NAFLD
			teatures in our study might be released by
			hepatocytes, as a protective mechanism against
			NAFLD progression ^[29] . Furthermore, we
			identified downregulation of plasma miR-146a-5p,

	miR-181a-5p, and miR-22-3p in individuals with
	NAFLD, which is supported by experimental
	studies of miRNA. For example, miR-146a
	targeted complex subunit 1 (MED1) to attenuate
	lipid accumulation and alleviate NAFLD
	progression in mice ^[76] . Additionally, miR-181a
	was found to downregulate peroxisome
	proliferator-activated receptor-a (PPARa) and
	mediate lipid metabolisms in NAFLD in human
	liver cells ^[77] . Moreover, miR-22 is a pivotal
	regulator of lipid and glucose metabolism, playing
	a crucial role in mitigating NAFLD progression in
	mice ^[78] . For example, miR-22 inhibited sirtuin-1
	and regulated gluconeogenesis in NAFLD ^[79] . We
	also observed increased expression of miR-125b-2-
	3p in both NAFLD and lobular inflammation,
	while studies indicated that miR-125b promoted
	the NF-ĸB-mediated inflammatory response in
	NAFLD ^[80] . Furthermore, we observed positive
	associations between liver fibrosis and expression
	of miR-34a-5p and miR-375, while experimental
	research suggested that both miR-34a-5p and miR-
	<mark>375 could alleviate liver fibrosis^[81, 82]. Together</mark>
	these experimental data support a plausible
	biological mechanism of NAFLD-miRNA
	association (Table 3).

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