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PEER-REVIEW REPORT

Name of journal: World Journal of Gastroenterology Manuscript NO: 89482 Title: Lipid metabolism-related lncRNA R Y1-817I4.1 promotes fatty acid synthesis and tumor progression in hepatocellular carcinoma Provenance and peer review: Unsolicited Manuscript; Externally peer reviewed Peer-review model: Single blind Reviewer's code: 02936529 **Position:** Editorial Board Academic degree: FRCS (Hon), MD, PhD Professional title: Professor, Surgical Oncologist **Reviewer's Country/Territory:** Brazil Author's Country/Territory: China Manuscript submission date: 2023-11-02 Reviewer chosen by: AI Technique Reviewer accepted review: 2023-11-04 02:05 Reviewer performed review: 2023-11-13 14:06

Review time: 9 Days and 12 Hours

	[] Grade A: Excellent [Y] Grade B: Very good [] Grade C:
Scientific quality	Good
	[] Grade D: Fair [] Grade E: Do not publish
Novelty of this manuscript	 [] Grade A: Excellent [Y] Grade B: Good [] Grade C: Fair [] Grade D: No novelty
Creativity or innovation of this manuscript	 [] Grade A: Excellent [Y] Grade B: Good [] Grade C: Fair [] Grade D: No creativity or innovation
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Scientific significance of the conclusion in this manuscript	 [] Grade A: Excellent [Y] Grade B: Good [] Grade C: Fair [] Grade D: No scientific significance
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	[]Yes [Y]No
Peer-reviewer statements	Peer-Review: [Y] Anonymous [] Onymous Conflicts-of-Interest: [] Yes [Y] No

SPECIFIC COMMENTS TO AUTHORS

The authors conducted the differential expression analyses in The Cancer Genome Atlas (TCGA) to identify lipid metabolism-related lncRNAs in HCC progressivos. qRT-PCR analysis was used to evaluate the expression of LMR lncRNAs and Nile red staining was used to observe intracellular lipid levels. Dual-luciferase reporter gene and RIP assays were performed to validate the interaction between RP11-817I4.1, miR-3120-3p, and ACLY. The authors concluded that 3 LMR-lncRNAs (NRAV, TMCC1-AS1, and RP11-817I4.1) were found to be predictive markers for HCC patients and were used to build risk models. Furthermore, RP11-817I4.1 knockdown reduced proliferation, migration, and invade. RP11-817I4.1 significantly increased lipid levels in HCC cells through the miR-3120-3p/ ACLY axis abs. They also concluded that LMR-lncRNAs have the capacity to predict the clinical characteristics and prognoses of patients with HCC and a new lncRNA, RP11-817I4.1, can accelerate the emergence and development of HCC. The methodology was proper and well described. The figures are illustrative and well displayed. The discussion is concise and updated. English polite is needed, mainly in concordance.



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