



PEER-REVIEW REPORT

Name of journal: *World Journal of Clinical Cases*

Manuscript NO: 69750

Title: Effect of prior malignancy on the prognosis of gastric cancer and somatic mutation

Provenance and peer review: Unsolicited manuscript; Externally peer reviewed

Peer-review model: Single blind

Reviewer’s code: 03731081

Position: Peer Reviewer

Academic degree: MD

Professional title: Professor

Reviewer’s Country/Territory: Russia

Author’s Country/Territory: China

Manuscript submission date: 2021-07-10

Reviewer chosen by: AI Technique

Reviewer accepted review: 2021-07-11 18:54

Reviewer performed review: 2021-07-14 09:12

Review time: 2 Days and 14 Hours

Scientific quality	<input type="checkbox"/> Grade A: Excellent <input checked="" type="checkbox"/> Grade B: Very good <input type="checkbox"/> Grade C: Good <input type="checkbox"/> Grade D: Fair <input type="checkbox"/> Grade E: Do not publish
Language quality	<input checked="" type="checkbox"/> Grade A: Priority publishing <input type="checkbox"/> Grade B: Minor language polishing <input type="checkbox"/> Grade C: A great deal of language polishing <input type="checkbox"/> Grade D: Rejection
Conclusion	<input type="checkbox"/> Accept (High priority) <input checked="" type="checkbox"/> Accept (General priority) <input type="checkbox"/> Minor revision <input type="checkbox"/> Major revision <input type="checkbox"/> Rejection
Re-review	<input type="checkbox"/> Yes <input checked="" type="checkbox"/> No
Peer-reviewer	Peer-Review: <input checked="" type="checkbox"/> Anonymous <input type="checkbox"/> Onymous



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statements

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

SPECIFIC COMMENTS TO AUTHORS

This article is a good analysis of the relationship between malignant neoplasms of various organs. This manuscript uses a lot of statistical material in the SEER dataset. The authors described an interesting phenomenon of prolongation of the survival time of patients with gastric cancer with previous malignant neoplasms. The authors analyzed the causes of gastric cancer, mutation genes TTN, TP53, MUC16, ARID1A, LRP1B and CNTNAP2.