



**ESPS PEER-REVIEW REPORT**

**Name of journal:** World Journal of Gastroenterology

**ESPS manuscript NO:** 20637

**Title:** Mutation analysis of 13 driver genes of colorectal cancer-related pathways in Taiwanese patients

**Reviewer’s code:** 00253956

**Reviewer’s country:** United Kingdom

**Science editor:** Jing Yu

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CLASSIFICATION	LANGUAGE EVALUATION	SCIENTIFIC MISCONDUCT	CONCLUSION
<input type="checkbox"/> Grade A: Excellent	<input type="checkbox"/> Grade A: Priority publishing	Google Search:	<input type="checkbox"/> Accept
<input type="checkbox"/> Grade B: Very good	<input type="checkbox"/> Grade B: Minor language polishing	<input type="checkbox"/> The same title	<input type="checkbox"/> High priority for publication
<input type="checkbox"/> Grade C: Good	<input type="checkbox"/> Grade C: A great deal of language polishing	<input type="checkbox"/> Duplicate publication	<input type="checkbox"/> Rejection
<input type="checkbox"/> Grade D: Fair	<input type="checkbox"/> Grade D: Rejected	<input type="checkbox"/> Plagiarism	<input type="checkbox"/> Minor revision
<input type="checkbox"/> Grade E: Poor		<input type="checkbox"/> No	<input type="checkbox"/> Major revision
		BPG Search:	
		<input type="checkbox"/> The same title	
		<input type="checkbox"/> Duplicate publication	
		<input type="checkbox"/> Plagiarism	
		<input type="checkbox"/> No	

**COMMENTS TO AUTHORS**

The manuscript entitled “mutational analysis of 13 driver genes of colorectal cancer-related pathways in Taiwanese patients” by Chang et al 2015 details the use of DNA sequencing techniques applied to colorectal cancer (CRC) samples, and details the identification of novel genetic mutations, as well as characterisation of the prevalence of other mutations in 13 driver genes of CRC-related pathways. This paper will be of interest to scientists working in the CRC field and I would therefore recommend publication. However, I would suggest the following minor revisions prior to publication: 1. The authors should provide a complete list of abbreviations and/or acronyms used throughout the manuscript, including protein and gene names. 2. Figure 1 is too small to be read easily, and the material suits just inclusion in a Table. Hence these figures could be enlarged to be made readable and included as a Supplementary Figure.