

## ESPS PEER-REVIEW REPORT

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

**ESPS manuscript NO:** 28704

**Title:** Development of a fast, low-cost diagnostic strategy based on real-time PCR and FRET probe melting curve analysis to identify single point mutations in highly variable genomes such as hepatitis C virus

**Reviewer's code:** 00504141

**Reviewer's country:** Ireland

**Science editor:** Yuan Qi

**Date sent for review:** 2016-07-13 16:40

**Date reviewed:** 2016-08-09 19:44

CLASSIFICATION	LANGUAGE EVALUATION	SCIENTIFIC MISCONDUCT	CONCLUSION
<input type="checkbox"/> Grade A: Excellent	<input checked="" type="checkbox"/> Grade A: Priority publishing	Google Search:	<input checked="" type="checkbox"/> Accept
<input checked="" 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 checked="" type="checkbox"/> No	<input type="checkbox"/> Minor revision
<input type="checkbox"/> Grade E: Poor		BPG Search:	<input type="checkbox"/> Major revision
		<input type="checkbox"/> The same title	
		<input type="checkbox"/> Duplicate publication	
		<input type="checkbox"/> Plagiarism	
		<input checked="" type="checkbox"/> No	

## COMMENTS TO AUTHORS

Well written paper. Nicely laid out, with sufficient detail to be repeated by others. One suggestion. As the authors indicate that they are preparing similar work for the NS5A mutations why combine the two pieces of research into one unified paper. The technique is a nice adaption of current RP-PCR protocols with detect Q80K which can be easily adapted to other parts of the genome and perhaps other viruses.

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**Title:** Development of a fast, low-cost diagnostic strategy based on real-time PCR and FRET probe melting curve analysis to identify single point mutations in highly variable genomes such as hepatitis C virus

**Reviewer's code:** 03478959

**Reviewer's country:** Brazil

**Science editor:** Yuan Qi

**Date sent for review:** 2016-07-13 16:40

**Date reviewed:** 2016-08-13 02:55

CLASSIFICATION	LANGUAGE EVALUATION	SCIENTIFIC MISCONDUCT	CONCLUSION
<input type="checkbox"/> Grade A: Excellent	<input checked="" type="checkbox"/> Grade A: Priority publishing	Google Search:	<input checked="" type="checkbox"/> Accept
<input checked="" 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 checked="" type="checkbox"/> No	<input type="checkbox"/> Minor revision
<input type="checkbox"/> Grade E: Poor		BPG Search:	<input type="checkbox"/> Major revision
		<input type="checkbox"/> The same title	
		<input type="checkbox"/> Duplicate publication	
		<input type="checkbox"/> Plagiarism	
		<input checked="" type="checkbox"/> No	

## COMMENTS TO AUTHORS

The paper presents a large scientific contribution to the diagnosis of hepatitis C. A new strategy to detect single point mutations by real-time PCR was presented. This technology can be Implemented in routine clinical laboratories to be a technique of low-cost and sensitive. The paper is well written. The technique is a good adaption of current RT-PCR protocols with detect Q80K which can be easily adapted to other parts of the HCV genome and perhaps other viral types.