



# BAISHIDENG PUBLISHING GROUP INC

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## ESPS PEER-REVIEW REPORT

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

**ESPS manuscript NO:** 26551

**Title:** Hepatitis C Virus (HCV) infection: an epidemiology up-date of HCV genotypes distribution

**Reviewer's code:** 00225318

**Reviewer's country:** Spain

**Science editor:** Ze-Mao Gong

**Date sent for review:** 2016-04-18 11:09

**Date reviewed:** 2016-05-03 18:50

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 checked="" type="checkbox"/> Grade B: Minor language polishing	<input type="checkbox"/> The same title	<input type="checkbox"/> High priority for publication
<input checked="" 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 checked="" type="checkbox"/> No	<input checked="" type="checkbox"/> Major revision
		BPG Search:	
		<input type="checkbox"/> The same title	
		<input type="checkbox"/> Duplicate publication	
		<input type="checkbox"/> Plagiarism	
		<input checked="" type="checkbox"/> No	

### COMMENTS TO AUTHORS

The introduction provides a good, generalized background of the topic that quickly gives the reader an appreciation of the wide search of updated information regarding the main subject. However, to make the introduction more substantial, the author may wish to provide references and technical explanation of the current commercial methods available to genotype. I think the motivations for this study it's clear enough, but it will improve if the author may wish to add the usefulness of the correct determination of the HCV genotype and subtype regarding the new DAA, providing examples of mutation associated with resistance. The searching methodology is appropriate for the study, although the author should probably provide more information about how the genotype and viral load was determined in each region. Maybe the author would benefit of an increase in data to analyze if he uses the result regarding of what he calls "non-representative population", and compare it with the rest of the population. Conclusions are sound, and give the reader a correct appreciation of the main subject in the article,



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**Title:** Hepatitis C Virus (HCV) infection: an epidemiology up-date of HCV genotypes distribution

**Reviewer's code:** 02528622

**Reviewer's country:** United States

**Science editor:** Ze-Mao Gong

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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
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<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

Very interesting paper. The analysis performed by the authors maximized the extraction of meaningful information from available literature. While subgenotyping information could not be assessed, the current HCV genotype distribution is relevant for the implementation of adequate control measures aimed to prevent virus spread.