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ESPS Peer-review Report

Name of Journal: World Journal of Gastroenterology

ESPS Manuscript NO: 8936

Title: Association of MYO9B gene polymorphisms with IBD in Chinese Han population

Reviewer code: 00505564

Science editor: Gou, Su-Xin

Date sent for review: 2014-01-23 13:26

Date reviewed: 2014-01-28 23:48

CLASSIFICATION	LANGUAGE EVALUATION	RECOMMENDATION	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"/> Existed	<input type="checkbox"/> High priority for publication
<input checked="" type="checkbox"/> Grade C (Good)	<input checked="" type="checkbox"/> Grade C: a great deal of language polishing	<input type="checkbox"/> No records	<input type="checkbox"/> Rejection
<input type="checkbox"/> Grade D (Fair)		BPG Search:	<input type="checkbox"/> Minor revision
<input type="checkbox"/> Grade E (Poor)	<input type="checkbox"/> Grade D: rejected	<input type="checkbox"/> Existed	<input type="checkbox"/> Major revision
		<input type="checkbox"/> No records	

COMMENTS TO AUTHORS

ESPS 8936 Review Association of MYO9B gene polymorphisms with IBD in Chinese Han population
Jing Hu, Qiao Mei, Jian Huang, Nai-Zhong Hu, Xiao-Chang Liu, Jian-Minh Xu The paper possesses a logical presentation of facts with regard to the description of the patient cohort, the performance of experiments and analysis of data. However, this reviewer highly recommends that the paper should be read and corrected by a professional proofreader before being reconsidered for publication. Concerning the contents of the paper. It would be helpful if it contained a definition what constitutes the Han population. Also, the description of the PCR method should include the sequences of primers and probes used, as well as methodological details regarding the quantification of nucleic acids, and testing of inhibition. The method of statistical analysis should be specified under the tables to explain the origin of the p values shown. Speaking of statistical analysis: as far as I know, genetic analyses such as exploring frequencies, often contain multiple comparisons of data, so a correction of the significance levels may be needed.