

## ESPS PEER-REVIEW REPORT

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

**ESPS manuscript NO:** 29909

**Title:** Gut microbial diversity analysis using Illumina sequencing for functional dyspepsia with liver depression-spleen deficiency syndrome and the interventional of Xiaoyaosan in rat model

**Reviewer's code:** 02535483

**Reviewer's country:** Finland

**Science editor:** Jing Yu

**Date sent for review:** 2016-08-30 12:32

**Date reviewed:** 2016-10-31 19:04

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 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 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 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

Dear Sirs, The enclosed manuscript is interesting and worth to publish. Minor lack is missing data from Xiaoyaosan, i.e what it is, which plant?, what does it contain, how it is used etc. I have never hear such medicine, thus these information should be added. Second comment is for the used database. Greengenes is quite often used but not reliable. Silva is golden standard for data analyses. Fortunately authors report data into only phyla level so the reliability of greengenes remains OK (satisfactory level).