



**SDI Review Form 1.6**

Journal Name:	<a href="#">Journal of Advances in Microbiology</a>
Manuscript Number:	Ms_JAMB_49827
Title of the Manuscript:	Gut Microbiota Dysbiotic Pattern and its associated Factors in a Cameroonian Cohort with and without HIV infection
Type of the Article	

**General guideline for Peer Review process:**

This journal's peer review policy states that **NO** manuscript should be rejected only on the basis of '**lack of Novelty**', provided the manuscript is scientifically robust and technically sound. To know the complete guideline for Peer Review process, reviewers are requested to visit this link:

(<http://www.sciencedomain.org/page.php?id=sdi-general-editorial-policy#Peer-Review-Guideline>)

**PART 1: Review Comments**

	Reviewer's comment	Author's comment (if agreed with reviewer, correct the manuscript and highlight that part in the manuscript. It is mandatory that authors should write his/her feedback here)
<b>Compulsory</b> REVISION comments	<p>Professional English editing is required</p> <p>The study should be a molecular study. The study does not include a sequence based assay. All the studies use next generation sequence techniques in microbiota studies. You can not show all the bacteria without using these techniques. The study cannot be published without these assays.</p>	<p>The sequencing based assay (culture-independent technique) was not carry out at the set point, due to unavailability of infrastructure and cost. This study was focused on the culture-dependent technique, and the organisms were identified with different biochemistry techniques. The study was design to captured just the culturable gut flora. Thus the pattern was focus mainly on culturable gut flora. Limitations of this study were acknowledged in the discussion section.</p> <p>Studies have published results on comparisons between culture-dependent and culture-independent techniques, of which their principal focus was to identify which technique could capture the majority of the gut flora.</p> <p>Our study focused on capturing culturable gut flora and not all the gut flora. Our current data has provided insight on culturable gut flora.</p>
<b>Minor</b> REVISION comments		
<b>Optional/General</b> comments		

**PART 2:**

	Reviewer's comment	Author's comment (if agreed with reviewer, correct the manuscript and highlight that part in the manuscript. It is mandatory that authors should write his/her feedback here)
Are there ethical issues in this manuscript?	(If yes, Kindly please write down the ethical issues here in details)	