

ORIGINAL ARTICLE**EVALUATION OF AN ANTIGEN-ANTIBODY “COMBINATION” ENZYME LINKED IMMUNOSORBENT ASSAY FOR DIAGNOSIS OF HEPATITIS C VIRUS INFECTIONS****Odari E.O^{1, 2}, Budambula N.L.M², Nitschko H¹****ABSTRACT**

BACKGROUND: Development of “combination” assays detecting in parallel, within a single test, Hepatitis C Virus (HCV) antigens and antibodies, not only reduces the window period in HCV-infection but also costs. Reduction of costs is important for developing countries where money and personal resources are limited.

METHODS: We compared the Monolisa® HCV Antigen-Antibody Ultra (Bio-Rad Laboratories Limited, Marnes La Coquette, France) with the AXSYM HCV version 3.0 (Abbot Diagnostics, Germany)-the latter assay detecting only antibodies to HCV. Seventy three HCV-PCR positive and negative samples were tested.

RESULTS: Although the two assays showed comparable results, two samples from a bone marrow transplant (BMT) patient of viral loads 7.8×10^5 and 8.9×10^6 IU/mL could not be detected by the Monolisa® HCV Antigen-Antibody Ultra assay. Failure to detect the two samples with viral loads considered above threshold of detection for antigen proteins suggested a lack of sensitivity by this assay to discover viral capsid protein in patient samples. Genotyping of these samples revealed genotype 1b, a HCV-subtype which is widespread and should thus be easily detected.

CONCLUSION: We conclude that although this assay depicts high sensitivity and specificity in detecting antibodies to HCV, it seems not to add further benefit in our study population to detect HCV infections by enhanced sensitivity due the potential contingency to trace viral capsid antigens.

KEY WORDS: Ag-Ab Combination assay, Hepatitis C Virus, ELISA, Monolisa HCV Ag-Ab Ultra

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INTRODUCTION

Hepatitis C virus (HCV) remains a major health care burden globally. It is currently estimated that 170 million people suffer from HCV chronic infections worldwide (1). The highest prevalence rates are still reported in Africa, Eastern Mediterranean, South East Asia and the Western Pacific. True incidence rates across the world continue to fluctuate. In developing countries exact incidence rates have been difficult to calculate given the asymptomatic, often latent nature of the disease prior to clinical presentation. Further prevalence rates across the developed world are changing since more countries are now

aware of transfusion-related HCV. Many regions still use blood donors to report the prevalence of HCV infection. Using blood donors may underestimate the true prevalence since they represent generally a highly selected population.

Studies show intravenous drug use as the leading risk factor for the spread HCV (2, 3) while others include transfusion of blood and blood products as well as hemodialysis (4-7) and sexual route of transmission (8-10). Therefore, the complexity and uncertainty related to the geographic distribution, chronic HCV, determination of its associated risk factors and

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evaluation of cofactors accelerating its progression, underscore the difficulties in global prevention and control of HCV.

Contrary to industrialized countries where anti-HCV screening and mini-pool nucleic acid amplification testing (NAT) is done on all collected blood hence a reduction in transmission, transfusion related transmission remains critical for many developing countries. The fact that HCV shows no clear clinical signs during the long window period further complicates the risk of transmission through other routes to the general population. Infection with HCV in Human Immunodeficiency Virus (HIV) infected could lead to an increased rate of progression to acquired immune-deficiency syndrome (AIDS), hamper immune reconstitution of patients and aggravate the risk of hepa-toxicity especially in patients on HIV treatment. Indeed, HCV infection has been reported to increase the number of complications in persons co-infected with HIV (11). With the increasing number of cases of HIV/HCV co-infection in Africa (12-16), there is a need for early HCV diagnosis so as to enhance optimum treatment and management of HIV patients. This would reduce possible complications resulting from dual infections.

Apart from early detection of HCV infection, combined antigen/antibody ELISA assays would be used for detection in the immune-suppressed or immune-deficient patients including HIV positive or patients with a history of bone marrow transplantation (BMT). Such patients show increased risk of serious veno-occlusive disease (VOD) of the liver (17-21). With recovery of cellular immunity after transplantation, severe hepatitis and fulminant liver failure may occur (22). In the absence of humoral or immediate cell mediated response in these patients, a rapid viral replication would lead to the production of various antigens. Today with BMT history of over 30 years, infection in long term survivors requires regular HCV follow-up assays and viral load tests.

This study aimed at evaluating the diagnostic performance of *Monolisa® HCV Ag-Ab ULTRA* (Bio-Rad) ELISA kit for simultaneous detection of HCV Antibodies and capsid antigens patients samples.

MATERIALS AND METHODS

Study design: The study involved both retrospective and prospective laboratory-based analysis of 73 (47 positive and 26 negative) archived plasma samples between May and September, 2012. The positive samples included a panel of 7 samples obtained at different time points from a BMT patient. Samples had been stored at -20°C before analysis and were used with permission from the virology department of the Max von Pettenkofer-Institute in Munich, Germany, where testing of the same samples was done.

Test principle for *Monolisa® HCV Ag-Ab ULTRA* detection kit: *Monolisa® HCV Ag-Ab ULTRA* assay is an enzyme immunoassay designed to detect both the capsid antigen and antibodies in serum or plasma. A micro-plate is coated with monoclonal antibodies against the capsid protein of HCV and two recombinant proteins produced in *E. coli*; one from the NS3 region (genotype 1 and 3a), and a recombinant antigen from the non-structural region NS4. A mutated peptide from the capsid structural region is also immobilized on the plate. Two conjugates are involved; conjugate 1 (R6) is a mouse biotinylated monoclonal antibody against the HCV capsid (it does not react with the HCV capsid mutated peptide coated on the micro-plate), Conjugate 2 (R7) is a mouse peroxidase-labeled antibody to human IgG and peroxidase-labeled streptavidin.

Test evaluations and sample characterization: The samples were first tested by “antibody only” ELISA kits (AxSYM HCV version 3.0 (Abbott Diagnostics, Germany) and Ortho® HCV 3.0 with Enhanced SAve (Ortho-Clinical Diagnostics, United Kingdom)) before amplification by PCR. Samples giving conflicting results by the ELISA kits were tested by western blot (INNO-LIA HCV score, Innogenetics® N.V, Belgium). Plasma HCV RNA quantification was performed by real time PCR using an in-house assay with internal positive controls. Based on the results, samples were categorized as; (1) HCV-RNA positive, (2) antibody positive-HCV-RNA negative, (3) HCV antibody negative-HCV-RNA negative, and (4) HCV-RNA positive BMT serial samples obtained over a period of 276 days.

Sensitivity: The 47 positive samples (31 HCV-RNA positive, 9 antibody positive- HCV-RNA

negative and 7 HCV-RNA positive serial samples) were used to evaluate the sensitivity of the assay. The 31 HCV-RNA positive samples were of genotypes 1 to 4 and a “mixed” genotype, with viral loads from 4×10^4 IU/mL to 29×10^6 IU/mL, whereas the 7 serial samples were of genotype 1b measuring from 3×10^3 to 10×10^6 IU/mL. Genotyping of plasma samples was determined by Restriction Fragment Length Polymorphism (RFLP), and where RFLP failed, the samples were sequenced and blasted in the HCV database for determination. Genotypes determined included 1a (4/31), 1b (7/31), 2a (5/31), 2b (1/31), 3a (6/31), 4 (7/31) and “mixed” [1a, 3b and 4] (1/31).

Specificity: Specificity was evaluated using 26 negative samples selected from a group of patients with no known risk of HCV infection who attended the hospital for regular checkups.

Test procedures: Test procedure for *Monolisa® HCV Ag-Ab ULTRA* was performed following the instructions of the manufacturer. Briefly, the diluted washing solution and the working antigen positive control were freshly prepared. The following components were added into the plate; 100µL of conjugate 1(R6) was added into each well, 50µL of negative control serum (R3) in well A1, 50µL of antibody positive control serum (R4) into B1, C1 and D1, 50µL of working antigen positive control into well E1 and 50µL of patient samples into in well F1 and the succeeding wells. The materials were mixed for 5 seconds covered with an adhesive film and incubated for 1 hour at 37°C. After incubation the plates were washed five times using 400µL of washing solution before addition of 100µL of conjugate-2 solution (R7) and a further incubation at 37°C for 30 minutes. After another 5-step wash, 80µL of freshly prepared enzymatic development solution was

added followed by a 30 minutes’ incubation in the dark at room temperature. The reaction was finally stopped using 100µL stopping solution (R10) and the optical density (OD) measured at 450/620 nm using a “Sunrise” ELISA reader.

The presence or absence of antibodies to HCV or/and HCV capsid antigen was determined by comparing the recorded absorbance for each sample with the calculated cut-off value. The cut-off value was determined by dividing the mean of the OD readings for the three positive controls by 4. Readings below the cut-off were considered non-reactive; samples below the cut-off value by less than 10% were retested. Samples above the cut-off values were considered initially reactive and retested in duplicate before a final interpretation was made. Results were compared with those of AxSYM HCV version 3.0 for the same samples.

Data analysis: Data were analyzed using GraphPad prism 5 statistical software. Pearson’s correlation (R^2) was used to correlate viral load against OD for ELISA assays. Sensitivity and specificity of the assays were tested using PCR and Western Blot Assays as the gold standards.

RESULTS

Overall performance of *Monolisa® HCV Ag-Ab ULTRA* assay kit (sensitivity, specificity and predictive values).

As shown in table 1, the two assays equally detected all the 31 HCV-RNA positive samples, and for each assay 4 out of 7 serial samples. Results, however, varied within HCV-RNA negative-antibody positive and the HCV-RNA negative-antibody negative samples.

Table 1: Suspected Hepatitis C samples tested with AXSYM and *Monolisa Ag/Ab* ELISA kits (n=73)

Test samples Sample type	Total	AXSYM		<i>Monolisa® HCV Ag-Ab</i>	
		Reactive	Non-Reactive	Reactive	Non- Reactive
PCR positive samples	31	3(100%)	0	31(100%)	0
Antibody positive- PCR negative	9	9(100%)	0	8(88.9%)	1(11.1%)
HCV negative samples	26	1(3.8%)	25(96.2)	0	26(100%)
Bonemarrow transplant patient	7	4(57.1%)	3(42.9%)	4(57.1%)	3(42.9%)
Total	73	45	28	43	30

Sensitivity and Specificity: In the first case, the overall sensitivity and specificity was calculated

for the two assays using the 73 samples. As shown in table 2 a higher sensitivity was realized with

AxSYM HCV version 3.0, and a likelihood ratio of 24.34, compared to *Monolisa® HCV Ag-Ab ULTRA assay* kit. Since false negative results only occurred in the BMT panel, a second analysis was done excluding this panel mainly to observe the performance of the assay on individual samples devoid of the effect of the serial samples. There was a general improvement in sensitivity in both assays, once more a likelihood ratio of 26.0 was

realized for AxSYM HCV version 3.0. Table 2 shows sensitivity and specificity results for the two ELISA assays. An improved sensitivity results was obtained for both assays when the samples from the BMT patient were excluded from the analysis. Likelihood ratios of 24.3 and 26.0 were obtained in the general and partial analyses respectively.

Table 2: Sensitivity, specificity and predictive values of *Monolisa Ag/Ab* and AxSYM HCV version 3.0 ELISA assays

ELISA assay	95% confidence interval			
	Sensitivity %	Specificity %	PPV %	NPV %
<i>Monolisa® HCV Ag-Ab ULTRA assay</i>				
Overall Analysis	91.5(43/47) (79.6 - 97.6)	100(26/26) (86.8 - 100)	100 (91.8 - 100)	86.7 (69.3 - 96.2)
Partial analysis**	97.5(39/40) (86.8 - 99.9)	100 (86.8 - 100)	100 (91-100)	96.3 (81 - 99.9)
AxSYM HCV version 3.0				
Overall Analysis	93.6 (82.5 - 98.7)	96.2 (80.4 - 99.9)	97.8 (88.2 - 99.9)	89.3 (71.8 - 99.7)
Partial analysis**	100(40/40) (91.2 - 100)	96.1 (80.4 - 99.9)	97.6 (87.1 - 99.9)	100 (86.3 - 100)

** 7 samples in the BMT sample panel excluded from the analysis

Viral load versus optical density: Evaluation of *Monolisa® HCV Ag-Ab ULTRA assay* and AxSYM ELISA kits was further done with samples of different viral loads. The samples included those below the theoretical threshold of 260,000 IU/mL (approximately 13.7 pg of total HCV core antigen)(23, 24) calculated for *Monolisa® HCV Ag-Ab ULTRA assay* kit (25). No

significant correlation was observed between viral loads and the optical density; $r(32) = .097$; $p = .60$ for *Monolisa® HCV Ag-Ab ULTRA assay* (Fig. 1) and $r(32) = .082$; $p = .66$ for AxSYM HCV version 3.0 (Fig. 2). The finding confirmed the earlier observation during the study where some samples of 8.9×10^6 IU/mL and 7.8×10^5 IU/mL were not detected by the two assays.

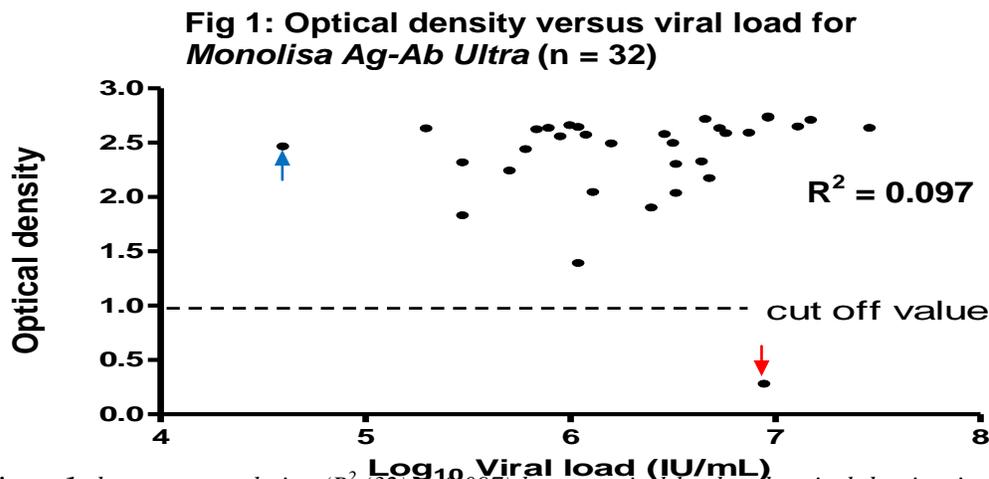


Figure 1 shows no correlation ($R^2 (32) = 0.097$) between viral load and optical density signals for *Monolisa Ag-Ab Ultra* assay. One sample of approximately 10×10^6 IU/mL was not detected (red arrow) since OD fell below the cut off value, whereas one sample below 1×10^5 IU/mL (blue arrow) was detected.

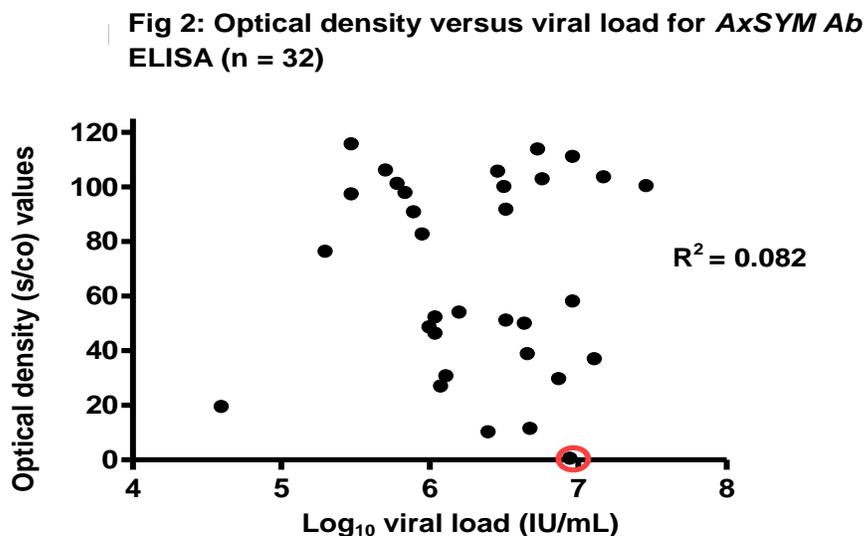


Figure 2 shows no correlation ($R^2 (32) = 0.082$) between viral load and optical density signals for *AxSYM, antibody ELISA*. One sample of approximately 10×10^6 IU/mL (circled in a red ring) was not detected since the OD was below the cut off value ($s/co = 1.0$), whereas on sample below 1×10^5 IU/mL was detected.

Samples from a Bone Marrow Transplant (BMT) patient: Since immune reconstitution after bone marrow transplantation has been studied (26-33) and found to vary between 6 months to over 2 years, detection of antibodies during this period remains minimal. The BMT samples were used to evaluate the efficacy in antigen detection by *Monolisa® HCV Ag-Ab ULTRA* assay. This capacity would be important if this assay would be used in immunocompromised patients. The BMT

samples were of genotype 1b, a genotype that is less responsive to alpha-interferon therapy compared to genotypes 2 and 3 (34), and is mainly found in the Americas, Europe and parts of Asia. All the detections were only observed in the pre-BMT samples (Fig. 3) with no detection on samples collected up to 276 days (about 9 months) post transplant. The ODs post transplant was below the cut-off-limits (0.5 and 1) of the 2

assays. The trends of detections were also similar

in both assays (Fig 3a and 3b).

Fig. 3a Monolisa® HCV Ag-Ab ULTRA on BMT patient

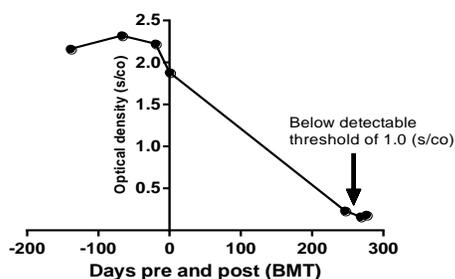


Fig. 3b AxSYM version 3.0 on BMT patient

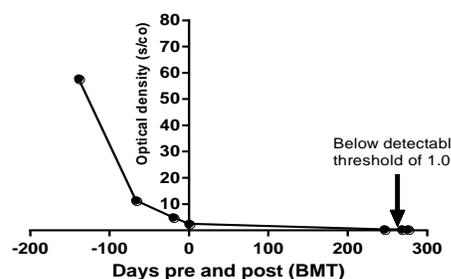


Fig. 3 Detection of HCV in plasma of a BMT patient

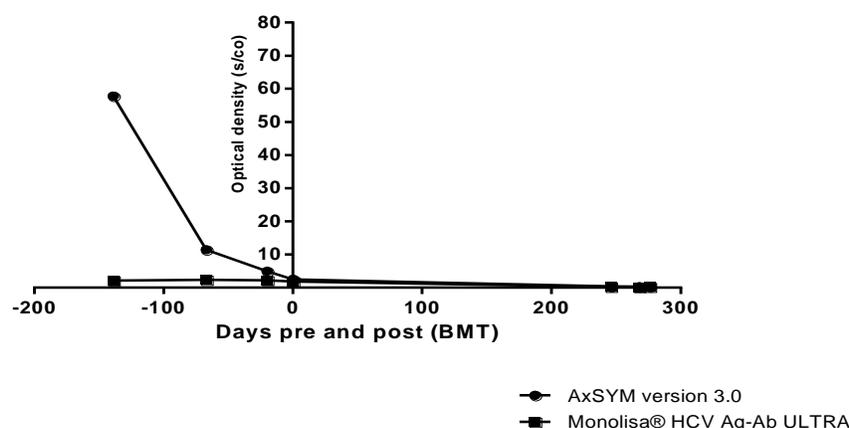


Figure 3 shows results of ELISA tests from plasma of a BMT patient obtained pre and post bone marrow transplant. All samples obtained before bone marrow transplant were positively detected (Figure 3a and 3b), whereas all samples collected more than 200 days after transplantation were not detected by the 2 assays.

DISCUSSION

ELISA assays combining antibody and antigen detections in a single kit have been developed and introduced in the field of HIV diagnosis (35). Introduction of similar assays for detecting acute HCV infection, which previously relied on classical serological methods (36), offers advantages over the existing antibody “only” assays mainly on early detection and detection in immunocompromised population.

Despite the anticipated advantages of “combination” assays, we report the failure of *Monolisa® HCV Ag-Ab ULTRA* to detect three PCR positive samples of genotype 1b from a BMT patient, despite the high viral loads in 2 samples. A similar study (25) also reported failure of *Monolisa® HCV Ag-Ab ULTRA* assay to detect a

genotype 1b (8.19×10^6 IU/mL) and a genotype 3a (7.76×10^6 IU/mL) samples. Further a genotype 4 HCV positive sample could not be detected by this assay despite the fact that the virus had a normal core sequence (37). These findings of failure to detect samples with high viral loads have a significant implication for this assay designed to detect both antigen and antibody proteins.

Arguably, HCV as a member of *Flaviviridae* family has an icosahedral capsid of T3 or T4 symmetry with 180 or 240 core protein subunits. Since one RNA genome contains 240 subunits of core protein of a molecular mass 19×10^3 Da, one picogram (pg) of virus core corresponds to 1.3×10^5 HCV genome molecules. Some studies even indicate lower ratios of viral particles per pg of proteins, for example in one study the ratio never exceeded 2×10^4 IU/mL (6×10^4

genomes/pg)(38), whereas another study estimated an average 8×10^3 IU/mL per pg of core proteins (5×10^3 IU/mL to 1.2×10^4 IU/mL per pg of proteins) (39). The latter findings were supported by our in-house results (unpublished data) which showed an antigen detection limit of 1.5pg/mL for the core antigen corresponding to 1.6×10^4 IU/mL. Based on these calculations, we note that less than 15% of core proteins are associated with HCV particles containing RNA and 240 core subunits, whereas the remaining 85% of the protein are associated with structures lacking HCV RNA. These structures have previously been described (40). The latter argument would explain the results realized in figure 1 where 2 samples with viral loads below the theoretical threshold were easily detected by this assay. These samples also showed higher optical signals compared to some samples with higher viral loads.

We however note that a similar trend of results shown in figure 1 was also realized in figure 2 where the 2 samples were again detected by the “antibody only” kit. The similarity in the trend of detection by the 2 different assays pointed towards the fact that the optical signals for these 2 samples were generated by the antibodies as opposed to antigens in the samples. This argument is strengthened by the fact that, upon generation of antibodies by the immune system, viral loads normally decline. The above argument therefore supports the view that *Monolisa® HCV Ag-Ab ULTRA* assay, as composed, is inclined to antibody detection than core antigens.

The two samples with viral loads above the threshold limit which were not detected by *Monolisa® HCV Ag-Ab ULTRA* assay similarly elicit various arguments about the assay. Some studies have failed to demonstrate the presence of antigens in RNA positive sera, for example, in one study,(41) antigens were not detected in sera despite measuring between 10^5 IU/mL and 8.2×10^5 IU/mL. In the latter study, some samples below 8.5×10^5 IU/mL were detected. Other studies have also demonstrated that analytical sensitivity of antigen assays does not constantly correspond the HCV- RNA titer (42). In the latter study 14 sera 10^5 and 10^6 IU/mL remained negative, while 12 sera below 10^5 IU/mL tested positive to antigen. Similar results as obtained by *Monolisa® HCV Ag-Ab ULTRA* assay were also realized by AxSYM assay, a clear indication that

the samples contained antibodies below the detectable limits possibly due to immunosuppression as a result of BMT. The fact that *Monolisa® HCV Ag-Ab ULTRA* assay detected all the BMT samples before transplantation, rules out the presence of antigens but only points towards antibody detection

Our observations therefore suggest that the diagnostic performance of this “combination” assay is inclined towards antibody than antigen detection, if any. Similar findings with “combo” assays in HIV diagnosis have been reported (43). A closer look at the results obtained by Laura Dean’s group (44) in their evaluation of *Monolisa® HCV Ag-Ab ULTRA* assay kit using different panels tend to suggest the same for this assay.

The overall sensitivity of 91.5% (with a higher sensitivity of 97.5% when BMT samples were excluded) and a specificity of 100% for *Monolisa® HCV Ag-Ab ULTRA* assay kit correspond to those reported in similar studies (42, 45-49).The ability of this assay to distinguish as negative a sample which had been falsely detected as antibody positive-HCV-RNA negative by AxSYM HCV version 3.0 ELISA but categorized as “indeterminate” by Ortho® HCV 3.0 with Enhanced SAve (and only resolved by western blot as negative), points towards its perfect discriminatory ability, hence specificity of the assay. Indeed, the mean absorbance value within the negative samples of 0.083 (0.033-0.219) was less than 30% of its cut off value of 0.327. This performance is comparable to those of ELISA kits routinely used in screening for blood-transmitted infections such as hepatitis B virus (HBV) and HIV infections. Such discriminatory ability has also been reported by Ansaldi (41), with a specificity of 100% for this same assay platform and by Dean’s group (44). It is, however, important to note that the results obtained in this study were not from sero-conversion panels, where many of *Monolisa® HCV Ag-Ab ULTRA* assay kit studies has been based, and as such these results cannot be used conclusively to predict the performance of this assay kit among sero-conversion samples. Further limitation of the study included low number of samples, hence the need for an increased number of samples. Although samples were also selected to reflect global distribution of genotypes, there is a need to

use a large number of samples mainly from immunocompromised patients of African origin, since eventually; this assay would be more beneficial to limited resource settings such as in Sub-Saharan Africa.

In conclusion, this study emphasizes, that although *Monolisa® HCV Ag-Ab ULTRA* assay kit presents an impressive sensitivity and specificity, the protein detection is more of antibody- than antigen-based. Even though we could not intensively test for antigen sensitivity, our results did not suggest any antigen sensitivity. In their analysis of window period samples, Laperche's group suggested the implementation of this assay in countries where only HCV antibody screening is the efficient measure of improving blood safety (50). A closer look at their results shows a low sensitivity of *Monolisa® HCV Ag-Ab ULTRA* assay in RNA positive-antibody negative samples where only 1/3 of such samples were detected among blood donors and less than ½ detected in hemodialysis panel (50). We, therefore, suggest that introducing this assay to replace currently available "antibody only" detection kits should be done with caution, unless it is based on cost implications. Further, this assay cannot be used to replace Nucleic Acid based Techniques (NAT) or core antigen-only-based assays in routine blood transfusion. Finally, *Monolisa® HCV Ag-Ab ULTRA* assay cannot be used alone for HCV diagnosis in immunocompromised patients.

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