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CONFLICT OF INTEREST DISCLOSURE

T.L. and L.N. are scientists employed by Diagnostics for the Real World Ltd (DRW); H.L, in addition to her academic position, is also the President and CEO of DRW. The other authors have no conflict of interest to disclose.

Performance of a New Rapid Test for the Detection of Hepatitis B Surface Antigen in Various Patient Populations

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1 **ABSTRACT**

2

3 **Background:** Rapid diagnostic tests (RDT) have been developed for the detection of
4 hepatitis B surface antigen (HBsAg). They represent a promising alternative to enzyme
5 immunoassays and a powerful tool for large-scale screening and diagnosis of HBV
6 infection, especially in regions without easy access to serological and molecular testing.

7 **Objectives:** The aims of the present study were to evaluate the characteristics and clinical
8 performance of a new CE-marked HBsAg RDT, DRW-HBsAg v2.0 assay (Diagnostics for the
9 Real World™, Ltd, USA), in various patient populations, including those chronically infected
10 with HBV, patients with severe acute hepatitis of unknown origin and pregnant women
11 with unknown HBV serological status at delivery. **Results:** The lower limit of detection of
12 the assay, evaluated in 21 clinical samples, ranged from 0.30 ± 0.07 to 0.97 ± 0.26
13 international units/mL (using Abbott Architect as a reference), depending on the HBV
14 genotype. The assay tested positive in 100% of patients with chronic hepatitis B, 96.3% of
15 HBsAg-positive acute hepatitis patients, and 95.2% of HBsAg-positive pregnant women. Its
16 specificity was 98.8% in HBsAg-negative patients, 98.7% in HBsAg-negative patients with
17 acute hepatitis of unknown origin and 97.8% in HBsAg-negative pregnant women. Amino
18 acid substitutions in the HBsAg major hydrophilic region did not affect HBsAg detection by
19 DRW-HBsAg v2.0. **Conclusions:** The new DRW-HBsAg v2.0 assay is a simple, rapid, easy-
20 to-run and highly sensitive assay that can be used in both high- and low-risk populations
21 for the diagnosis of HBsAg carriage. It appears to be a promising new tool for large-scale
22 screening and diagnosis of HBV infection.

23

1 **BACKGROUND**

2

3 Approximately 240 million people are chronically infected with hepatitis B virus
4 (HBV) worldwide.¹ Each year, over 600,000 people die from HBV-related chronic liver
5 disease through end-stage cirrhosis, liver failure and hepatocellular carcinoma (HCC).²
6 Hepatitis B is a global health problem, with geographic disparities in terms of prevalence.
7 In high endemicity areas, 70% to 90% of the population becomes HBV-infected before the
8 age of 40, and 8% to 20% of individuals are chronic HBV carriers. In contrast, in low
9 endemicity areas, the chronic carriage rate is less than 2%.

10 Because chronic HBV infection is often asymptomatic, the vast majority of infected
11 individuals are unaware of their liver disease, including in high medical standard areas. For
12 instance, in France, more than 45% of infected individuals ignore their infection.³ Early
13 diagnosis and treatment can reduce the risk of development of long-term complications
14 and prevent further transmission. Hepatitis B surface antigen (HBsAg) is the key marker
15 for screening and laboratory diagnosis of HBV infection. Rapid diagnostic tests (RDT) have
16 been developed and widely used for viral infection screening. A high number of RDTs with
17 satisfactory performance for the detection of antibodies to human immunodeficiency virus
18 (HIV) have received product license in the United States and Europe. In contrast, very few
19 rapid tests for HBsAg meet the analytical sensitivity of 0.130 international units (IU)/mL
20 required by the regulatory authorities. Indeed, these RDTs in general lack the analytical
21 sensitivity compared to enzyme immunoassays (EIA), the reference techniques for HBsAg
22 detection in plasma or serum specimens.⁴

23

24 **OBJECTIVES**

1 The aims of the present study were to evaluate the test characteristics and clinical
2 performance of a newly developed HBsAg RDT, DRW-HBsAg v2.0 assay (Diagnostics for the
3 Real World™, Ltd, Sunnyvale, USA), in various populations of patients.

4 5 **STUDY DESIGN**

6 7 **Materials**

8
9 Plasma and serum samples were obtained from HBsAg-positive and -negative
10 individuals with various conditions.

11 Group A was comprised of 335 HBsAg-negative individuals with different
12 serological profiles for HBV infection, including no markers of past infection (n=138),
13 vaccination profile (n=94), isolated anti-HBc antibodies (n=29), or markers of resolved
14 past infection (n=74). Among them, 34 were HIV-seropositive and 48 were HCV-
15 seropositive.

16 Group B was comprised of 203 patients with chronic HBV infection, all of whom had
17 quantifiable HBsAg by EIA (mean HBsAg titer: $3.85 \pm 0.40 \text{ Log}_{10}$ international units (IU)/mL;
18 range: 2.40-5.10), anti-HBc antibodies, and detectable HBV DNA (mean HBV DNA level:
19 $3.95 \pm 1.45 \text{ Log}_{10}$ IU/mL; range: 1.80-8.75). This group comprised 49 patients infected with
20 HBV genotype A, 18 with genotype B, 23 with genotype C, 65 with genotype D, and 48 with
21 genotype E.

22 Group C included 20 blood donors with chronic HBV infection infected with a
23 dominant variant bearing one or several amino acid substitutions in the major hydrophilic
24 region (MHR) containing the “a” determinant of the S gene. Among them, 3 were infected
25 with HBV genotype A, 4 with genotype B, one with genotype C, 8 with genotype D and 4

1 with genotype E. All had detectable/quantifiable HBsAg by EIA ($2.75 \pm 1.00 \text{ Log}_{10} \text{ IU/mL}$;
2 range: -0.90 to 4.30) with one donor who had an HBsAg titer above the upper limit of
3 quantification of the assay ($>5.4 \text{ Log}_{10} \text{ IU/mL}$). HBV DNA was detectable in 60.0% of them.

4 Group D was comprised of 408 patients with severe acute hepatitis; 27 of them had
5 detectable/quantifiable HBsAg ($2.90 \pm 0.95 \text{ Log}_{10} \text{ IU/mL}$, range: -0.50 - 4.40), and HBV DNA
6 was detectable in 81.5% of them. The remaining patients were HBsAg-negative by EIA.

7 Group E included 802 pregnant women with unknown HBsAg status at delivery; 21
8 of them had detectable/quantifiable HBsAg by EIA (mean $3.00 \pm 0.90 \text{ Log}_{10} \text{ IU/mL}$, range:
9 0.15 - $4.10 \text{ Log}_{10} \text{ IU/mL}$), and HBV DNA was detectable in 80.0% of them.

10

11 **Laboratory measurements**

12

13 Serum HBsAg levels were quantified by means of the Architect HBsAg assay (Abbott
14 Diagnostic, Chicago, Illinois), after a 1:100 dilution, as recommended by the manufacturer.
15 The dynamic range of quantification of this assay is 0.05 to 250.0 IU/mL. Samples with
16 HBsAg levels $<0.05 \text{ IU/mL}$ at 1:100 dilution were retested undiluted, while those with
17 HBsAg $>250.0 \text{ IU/mL}$ at 1:100 dilution were retested at a final dilution of 1:999, according
18 to the manufacturer's instructions. The use of the Architect HBsAg assay as the reference
19 test was justified by previous reports showing it is accurate, precise, reproducible and well
20 calibrated to the World Health Organization (WHO) HBsAg standard.^{5, 6} We confirmed its
21 good calibration by testing 5 members of the WHO HBsAg subtype adw2 genotype A
22 reference panel (code: 03/262, NIBSC, UK) 5 times in two different experiments (data not
23 shown).

1 Serum HBV DNA levels were measured by means of a real-time PCR assay
2 (*m2000_{RT}/m2000_{SP}*, Abbott Diagnostic).⁷ Results were expressed in IU/mL. The lower limit
3 of detection of the assay is 15 IU/mL.

4 The HBV genotype was determined by directly sequencing a portion of the
5 overlapping genes encoding HBsAg and the B and C subdomains of the HBV reverse
6 transcriptase. Sequence analysis was performed as previously described.⁸⁻¹⁰

7 The amino acid substitutions in the MHR were identified by directly sequencing a
8 portion of the S gene according to a previously described method.¹¹ Amino acid sequences
9 were compared to a consensus sequence of region 100-178 of HBsAg using reference
10 sequences of different subgenotypes from GenBank.

11 **HBsAg RDT**

12
13 The DRW-HBsAg v2.0 assay is a new rapid immunochromatographic test for the
14 qualitative detection of HBsAg that is performed manually. The test consists of a test strip
15 (nitrocellulose membrane) and a single-use tube containing two lyophilized reagent beads.
16 One reagent bead contains a hapten-labeled mouse monoclonal antibody to HBsAg
17 (primary antibody), and the other contains a mouse monoclonal antibody to hapten
18 conjugated to colloidal gold (secondary antibody). Eighty μ L of human serum or plasma is
19 added to the tube, which is then mixed gently before insertion of the test strip. The sample
20 mixture flows up the test strip by capillary action, and the visual result is read after 30 min.
21 The development of a purple line in the test line region indicates the presence of HBsAg in
22 the clinical specimen. The absence of a colored line indicates the absence of HBsAg or its
23 presence at a concentration below the limit of detection of the test. Each test strip also
24 incorporates a built-procedural control for reagent stability and test functionality, and
25 indicates that the full reagents have properly flowed up the test strip.

1 The test result is read visually according to the manufacturer's instructions.
2 However, in this study, an automatic readout by means of ESE-Quant Lateral Flow Reader
3 (ESE GmbH, Germany) was evaluated and compared with visually read results. The strips
4 were visually analyzed by two independent investigators and automatically with the ESE-
5 Quant Lateral Flow Reader. A cutoff optical density value of 20 was established with a
6 standard panel: an optical density value <20 indicated the absence of HBsAg, whereas
7 values >25 indicated the presence of HBsAg. A grey zone was defined between 20 and 25.

8

9 **Statistical analysis**

10

11 Descriptive statistics are shown as the mean \pm standard deviation (SD). The
12 sensitivities, specificities and the positive and negative predictive values of the RDT DRW-
13 HBsAg v2.0 for the detection of HBsAg were calculated.

14

15 **RESULTS**

16

17 **Analytical sensitivity in clinical samples of different HBV genotypes**

18

19 To determine the analytical sensitivity of the DRW-HBsAg v2.0 assay, serial
20 dilutions of clinical samples from 21 patients from group B, infected with HBV genotypes A
21 to E, were tested once. The lower limit of detection was estimated as the lowest amount of
22 HBsAg that could be detected by the DRW-HBsAg v2.0 assay. As shown in Table 1, the
23 lower limit of detection of the assay ranged from 0.30 ± 0.07 IU/mL to 0.97 ± 0.26 (using
24 Architect HBsAg assay as a comparator), depending on the HBV genotype.

25

1 **Specificity**

2
3 The specificity of the DRW-HBsAg v2.0 was assessed by testing the 335 clinical
4 specimens from group A, the 381 HBsAg-negative samples from group D (acute hepatitis)
5 and the 781 HBsAg-negative samples from group E (pregnant women at delivery).

6 Of the 335 HBsAg-negative specimens from group A, 4 tested HBsAg-positive with
7 DRW-HBsAg v2.0 (specificity: 98.8%; 95% confidence interval [95%CI], 96.9%-99.5%).
8 Among these 4 false-positive results, two patients were HBV seronegative (no markers of
9 past or ongoing infection), and two individuals had been vaccinated and harbored anti-HBs
10 antibodies only. The serological status for HCV and HIV did not interfere with the
11 specificity of the DRW-HBsAg v2.0 assay.

12 Among the 381 HBsAg-negative samples from patients with acute hepatitis of
13 unknown origin, there were 5 false-positive results with DRW-HBsAg v2.0 (specificity:
14 98.7%; 95%CI: 96.9%-99.4%). Among the 781 HBsAg-negative samples from pregnant
15 women with unknown HBV status at delivery, there were 17 false-positive results with
16 DRW-HBsAg v2.0 (specificity: 97.8%; 95%CI: 96.6%-98.7%). None of these individuals was
17 HBsAg-positive in EIA (lower limit of detection: 0.05 IU/mL) or HBV DNA-positive with
18 real-time PCR (lower limit of detection: 15 IU/mL).

19 20 **Clinical sensitivity, accuracy and influence of the HBV genotype**

21
22 Among 203 patients chronically infected with HBV genotypes A to E with
23 detectable/quantifiable HBsAg by EIA and HBV DNA by real-time PCR, all tested positive
24 for HBsAg by the DRW-HBsAg v2.0 assay (clinical sensitivity: 100%; 95%CI: 98.1%-100%).
25 HBsAg was not detected by DRW-HBsAg v2.0 in one out of 27 patients with severe acute

1 hepatitis ascribed to HBV due to the presence of HBsAg by EIA (sensitivity: 96.3%; 95%CI:
2 81.7%-99.3%). This patient was infected with HBV genotype D, HBsAg level was 0.32
3 IU/mL, anti-HBc IgM was present and HBV DNA level was 4.5 Log₁₀ IU/mL. Similarly, DRW-
4 HBsAg v2.0 failed to detect HBsAg in one out of 21 pregnant women found to be HBsAg-
5 positive by EIA (sensitivity: 95.2%; 95%CI: 77.3%-99.1%). This patient had an HBsAg level
6 of 0.70 IU/mL. The presence of HBsAg was confirmed by two different EIA assays and
7 neutralized, and HBV DNA level was undetectable (<15 IU/mL).

8

9 **Influence of amino acid substitutions in HBsAg MHR**

10

11 Table 2 shows the amino acid positions with substitutions in the MHR in the 20
12 blood donors from group C identified as HBsAg mutants. HBsAg was detected in all of them
13 by means of DRW-HBsAg v2.0, regardless of the amino acid substitution(s). A low signal
14 was observed in one case, with an HBsAg titer of 0.13 IU/mL.

15

16 **Comparison between visual and automatic readouts**

17

18 The results of the automated readout were compared to those obtained by visual
19 analysis. Among 1,768 patients included in the study, 157 (8.9%) had discrepant results
20 between visual and automatic readouts (Table 3). The test was interpreted as positive with
21 both reading methods in all of the HBsAg-positive patients from group B (203 patients with
22 chronic HBV infection), group C (20 blood donors infected with a dominant variant bearing
23 substitutions in the MHR), group D (27 patients with severe acute hepatitis B) and group E
24 (21 pregnant women with HBV infection). A false-positive result was obtained with the
25 automated strip reader in 25 patients from group A, 63 patients from group D and 69

1 pregnant women from group E. Only 8 out of 161 discrepant results fell within the gray
2 zone of the automated reader. All of them were HBsAg-negative with the Architect assay
3 and had undetectable HBV DNA with the *m2000* real-time PCR assay.

4

5

6 **DISCUSSION**

7

8 In the last 20 years, the availability and use of point-of-care (POC) tests have
9 markedly increased and expanded to all fields of medicine. In the setting of infectious
10 diseases, most existing POC tests consist of immunoassays, including agglutination or
11 immunochromatographic strips (strip tests). These methods can be used to detect HBsAg.
12 Other non-immunological POC tests based on nucleic acid detection and eventually
13 quantification are currently in development. Rapid diagnostic tests remain promising
14 alternatives to EIA-based methods. Indeed, they offer the advantages of simplicity, limited
15 need for instrumentation, minimal training required, and rapid performance at room
16 temperature.¹² Therefore, these assays represent a powerful tool for large-scale screening
17 and subsequent diagnosis. They are of particular value to improve access to care in regions
18 without easy access to serological and molecular biology testing. They can also be used at
19 the bedside or in remote care centers in any area of the world.

20 The results of the present study show that the newly developed DRW-HBsAg v2.0
21 rapid test has a better analytical sensitivity than other commercially available point-of-care
22 HBV tests. Indeed, its lower limit of detection was <1.0 IU/mL of HBsAg, whereas among 19
23 HBsAg rapid assays recently tested, the lower limit of detection was ≥ 1.5 IU/mL in all cases
24 and >4.0 IU/mL in most of them.⁴ The analytical sensitivity required by regulatory agencies
25 is 0.130 IU/mL of HBsAg, subtype adw2, genotype A (NIBSC 00/588). To obtain CE mark,

1 the manufacturer has provided evidence for reaching the sensitivity requirement. In
2 addition, each lot of the DRW-HBsAg v2.0 assay was submitted to the Paul Ehrlich Institute
3 for sensitivity and specificity testing prior to its release and met the CE-mark requirement.
4 In the present work, the analytical sensitivity of DRW-HBsAg v2.0 in 23 clinical samples
5 ranging in genotype from A to F was slightly above this threshold, using the Architect
6 HBsAg assay as a reference. Our study also demonstrated the improved clinical sensitivity
7 of DRW-HBsAg v2.0 for different HBV genotypes, as compared to the first-generation DRW-
8 HBsAg assay.¹³ Indeed, among 271 HBsAg-positive patients tested in this study, DRW-
9 HBsAg v2.0 failed to detect HBsAg in only two cases, both containing a low level of HBsAg
10 (<1.0 UI/mL).

11 The genetic variability of the S gene can impair the diagnostic performance of HBsAg
12 assays, but the frequency of such variants in HBV-infected patients is not well established
13 and varies according to the HBV endemicity.¹⁴ We confirmed here the manufacturer's claim
14 that DRW-HBsAg v2.0 was able to detect HBsAg in patients infected with HBV variants
15 bearing various amino acid substitutions or combinations of substitutions in this region
16 (Table 2).

17 Assay sensitivity is a key criterion for large-scale HBsAg testing. However, test
18 specificity is also crucial because it influences the requirement for result confirmation in
19 areas where resources for such confirmation are likely to be limited. In this study, we
20 evaluated the specificity of DRW-HBsAg v2.0 in a large number of HBsAg-negative samples
21 (n=1497 overall). Our results showing that specificity of the assay was $\geq 98.0\%$ indicated an
22 effective balance between specificity and sensitivity. Specificity was optimal with visual
23 reading in our experiments. Overall, as shown in Table 3, visual reading was found to be
24 more reliable than automatic readout by means of ESE-Quant Lateral Flow Reader, which
25 needs to be further optimized.

1 In conclusion, our study shows that the new, second version of the DRW-HBsAg
2 rapid test is a simple, rapid, easy-to-run and highly sensitive assay that can be used in both
3 high- and low-risk populations for the diagnosis of HBsAg carriage with a 99.3% accuracy.
4 Thus, RDTs appear as promising new tools for large-scale screening and diagnosis of HBV
5 infection in the clinical setting.

6

7 **FUNDING**

8

9 The DRW-HBsAg v2.0 assay kits were kindly provided by Diagnostics for the Real
10 World™, Ltd, Sunnyvale, California USA.

11

12 **COMPETING INTERESTS**

13

14 T.L. and L.N. are scientists employed by Diagnostics for the Real World Ltd (DRW);
15 H.L, in addition to her academic position, is also the President and CEO of DRW. The other
16 authors have no conflict of interest to disclose.

17

18 **ETHICAL APPROVAL**

19 Patients gave their consent to the use of their residual samples.

20

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20

21

Table 1. Analytical sensitivity of DRW-HBsAg v2.0, according to the HBV genotype (using Architect HBsAg assay as a comparator).

HBV genotype	N	Mean±SD lowest HBsAg level detected (IU/mL)
Genotype A	5	0.38±0.12
Genotype B	4	0.52±0.17
Genotype C	4	0.30 ±0.07
Genotype D	4	0.40±0.10
Genotype E	4	0.97±0.26

Table 2. HBsAg titers (Architect HBsAg assay) in the 20 blood donors from group C with amino acid substitutions in the major hydrophilic region (MHR) containing the “a” determinant of the S gene.

Blood donors	Genotype	HBsAg titer (Log ₁₀ IU/mL)	Amino acid substitution(s) in the MHR
1	E	4.1	D144E
2	D	2.5	T131I
3	D	0.6	T131N, M133I, T134H
4	C	3.9	I126N
5	D	2.8	T118V, A128V, Q129L, S171F, W172*stop
6	A	3.9	M133T
7	D	0.7	Q101R, M103I, T118K, S143L, F158L, W172L
8	D	3.7	Y134N
9	B	3.3	P120S, F134I
10	B	2.9	T126A
11	A	3.4	Q129H, D144A
12	D	2.3	T118A
13	B	4.3	G129H
14	D	1.8	G130R
15	D	2.4	G130R
16	E	4.0	G145R
17	E	2.8	S143L
18	B	-0.9	M133T, T116N
19	E	3.3	G145A
20	A	>5.4	T116N

Table 3. Visual reading *versus* automatic readouts by means of ESE-Quant Lateral Flow Reader for HBsAg detection with DRW-HBsAg v2.0 in the different study groups.

		Visual readout	
		Positive	Negative
Group A: HBsAg-negative individuals (n=335)			
Automatic readout	Positive	<u>4</u>	<u>22</u>
	Gray zone	0	3
	Negative	0	306
Group B: chronic HBV infection (n=203)			
Automatic readout	Positive	203	0
	Gray zone	0	0
	Negative	0	0
Group C: blood donors (n=20)			
Automatic readout	Positive	20	0
	Gray zone	0	0
	Negative	0	0
Group D: severe acute hepatitis (n=408)			
Automatic readout	Positive	<u>31</u>	63
	Gray zone	0	0
	Negative	0	<u>314</u>
Group E: pregnant women (n=802)			
Automatic readout	Positive	<u>37</u>	64
	Gray zone	0	5
	Negative	0	<u>696</u>