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by Sikander Gill, Li Wenji, Fan Xu, Rajwant Gill, He Bin, Nick Bandy and Dong Liang

## Automation of a One-Step, Real-Time PCR-Based DNA Quantitative Kit for the Diagnosis of HBV

utomation of one-tube or one-step real-time PCR-based molecular diagnostics has distinct advantages for sample processing in preparative work. It saves time and effort, improves processing capacity, eliminates human error and ensures batch consistency of reagents and samples in the detection and diagnostic process of clinical samples. The VERSA 10 workstation (Aurora Biomed, Inc., Vancouver, B.C.) was validated for automated use for a Hepatitis B viral (HBV) DNA quantitative fluorescent diagnostic kit (Sansure Biotech, Changsha, Hunan, China).

Results show that the automated operation was performed with consistency, appreciable throughput and accuracy with no detectable cross-contamination. Ninety-six samples can be processed within 25 minutes using a total of 108 50-µL tips with one click on the software interface. The system includes a high-efficiency air filter hood, an automated shaker and liquid handling.

### Introduction

Hepatitis B is a significant global health problem: more than 780,000 people die each year due to complications associated with the virus. Its prognosis and diagnosis have been primarily based on the immunodetection of HBV surface antigen. The lack of detectable antibodies in human patients has necessitated the use of nucleic acid amplification-based assays for their sensitivity, specificity and tolerance of sequence variation.<sup>1,2</sup> In the last few years, the field of HBV molecular diagnostics and prognosis has included such tools as real-time PCR.<sup>3</sup> A one-tube or one-step real-time detection and quantification method for HBV DNA using nucleic acid sequence-based amplification further advances the benefits of these methods.<sup>4,5</sup> Sample processing and amplification are performed in one tube and the need to heat or centrifuge samples is eliminated.



Figure 1 – VERSA 10 workstation with HEPA/UV enclosure.

### **Materials and methods**

The validation of the kit was conducted on the VERSA 10 workstation (*Figure 1*) with four-channel liquid handling configuration as follows:

- 48 samples including 42 different concentration HBV serum samples, four standards, one negative and one positive control.
- 2. Hepatitis B viral DNA quantitative fluorescence diagnostic kit, 48 tests (*Figure 2*).
- 3. Software.
- 4. The following steps were automated in a PCR plate:
  - a. Addition of samples and reagents
  - Addition of 5 µL of lysis buffer: addition of 5 µL of sample and 40 µL of master mix and primer sets
  - c. Sensitivity test: 1.96 e+05 IU/mL of hepatitis sample, of high concentration, was diluted to 1.96 e+02 IU/mL low concentration to determine whether automation could meet kit performance requirements while dealing with low-concentration samples

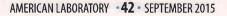




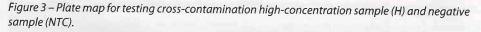
Figure 2 – Hepatitis B viral DNA quantitative fluorescence diagnostic kit.

- d. A cross-contamination test was designed as shown in the plate map (*Figure 3*). Mixing of the contents in the plate was done at 600 rpm for 10 minutes on the plate shaker housed on the deck
- The manual process was carried out in parallel to the automation for comparative evaluation as per the manufacturer's directions.
- The plate was manually transferred to the SLAN real-time PCR thermocycler for amplification (Hongshi, Shanghai, China).

### Results Sensitivity

The objective of this experiment was to examine whether the workstation can achieve kit performance requirements while dealing with low concentration samples. The amplification curves in *Figure 4a* and *b* indicate that the VERSA 10 procedure achieves better performance than the manual method when handling low concentration samples. As seen in Figure 4, automated sample preparation resulted in more consistent cycle threshold (ct) values and amplification curves, accounting for Sample processing and amplification are performed in one tube and the need to heat or centrifuge samples is eliminated.

	1	2	3	4	5	6
A	Н	NTC	Н	NTC	н	NTC
В	NTC	Н	NTC	н	NTC	Н
C	H	NTC	Н	NTC	Н	NTC
D	NTC	Н	NTC	н	NTC	Н
E	H	NTC	H	NTC	н	NTC
F	NTC	Н	NTC	Н	NTC	Н
G	Η	NTC	Н	NTC	Н	NTC
H	NTC	Н	NTC	Н	NTC	н



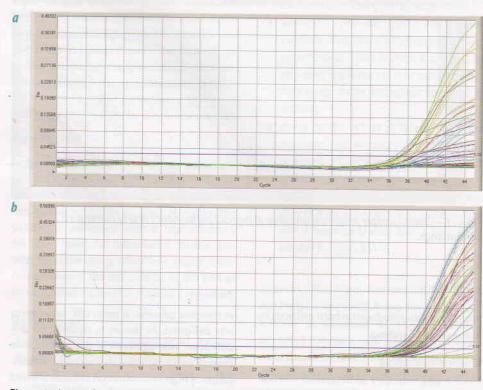


Figure 4 – Manual (a) and automated (b) comparison test for 48 samples.

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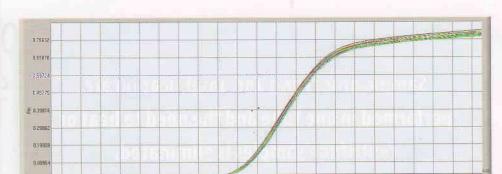
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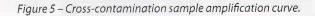
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**PCR-BASED DNA QUANTITATIVE KIT continued** 

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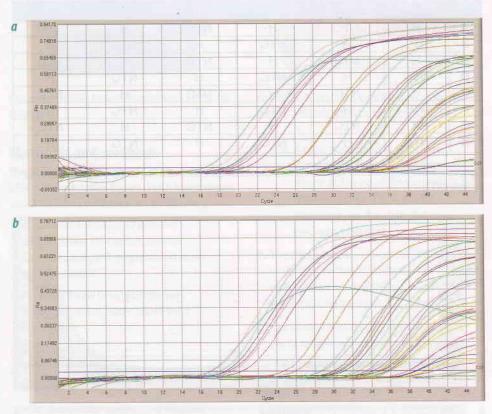


Figure 6 – Clinical sample amplification curve: a) manual performance and b) automated performance.

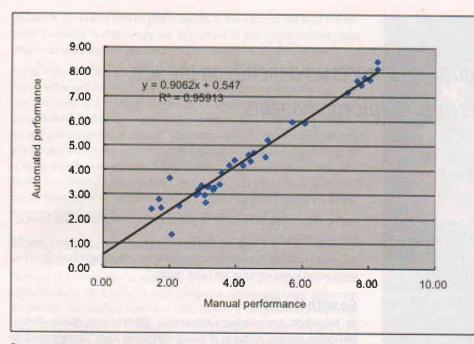
inconsistencies introduced in both methods by human liquid handling.

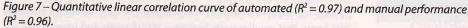
#### **Cross-contamination**

Since it was necessary to determine whether there is any cross-contamination during the

automated process, a cross-talk experiment was designed. As shown in *Figure 5*, no amplification was observed in the no template control (NTC) samples. Additionally, amplification of high-concentration samples showed very good repeatability.

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### **Clinical sample test results**

The purpose of this experiment was to test the performance requirements of the kit on VERSA 10 for dealing with clinical samples. Fortyeight samples were run in both automated and manual methods. The amplification curves and comparative quantitative analysis are presented in *Figures 6* and *7*, respectively. As shown in Figure 7, the quantitative linear correlation curve for the automated samples ( $R^2 = 0.97$ ) confirms similar performance to manual performance ( $R^2 = 0.96$ ) for the clinical samples.

### Conclusion

Automation offers higher consistency, throughput and accuracy than the manual process; can also provide time and energy savings; and offers confidence in the accuracy of results. The VERSA 10, equipped with a high-efficiency air filter hood, an automated shaker and with liquid handling, effectively avoids crosscontamination. The automation of this kit can be realized with one-click automatic operation.

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