Validation of anti HCV-hepatitis c identification in blood donors using PCR: case study at Tuban East Java district, Indonesia.

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Abstract

The validation of virus-free blood donors is often neglected possibly related to the high cost process analyzing. While, validity of the medical status of hepatitis C virus (HCV) is crucial for blood donors quality. ELISA is common method for HCV detection, however inaccurate result possibly occurred during the window period. Hence, the validation of this method using molecular test is still required. The samples were collected from 100 blood donors consist of 85 males (85%) and 15 females (15%) from Tuban region, East Java Indonesia. The ELISA kit test resulted 100% of negative HCV. Surprisingly, after validated using PCR there were found 6 blood samples (6%) positively infected HCV. This study encourages who one concerned to propose that PCR-based validation for screening blood donors is necessarily required.

Keywords: Blood donors, ELISA, PCR, Anti HCV virus, Tuban.

Introduction

Virus infection caused by hepatitis C virus (HCV) is still an issue on global health. Hepatitis C virus is an RNA virus from hepacivirus family, single stranded and have similarity with flavivirus [1,2]. Since the hepatitis C virus can be transmitted from blood transfusion (90% from cases), the evaluation of HCV in the blood among donors is crucially important [2,3]. ELISA is common methods for HCV detection, however inaccurate result possibly occurred during the window period, hence the validation by re-testing this method using molecular test using Polymerase Chain Reaction (PCR) is still required [4,5].

This research would be useful for the local government as scientific recommendation to include the PCR validation test into their Standard Operating Procedure (SOP) of screening blood donors. This case report was conducted in Tuban as a representative region in East Java, Indonesia.

Material and Methods

Sample collection

Blood samples were collected from 100 donors in Tuban region according to standard procedure from the Indonesian Red Cross Society.

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Hepatitis C identification

Serum from 100 blood donors was then subjected to Anti-HCV identification using Enzyme-Linked Immunosorbent Assay (ELISA), following the instructions (HCV Combo, Biorad, Germany).

Furthermore, the serum samples with negative anti-HCV were prepared for retesting using PCR analysis. Prior to PCR analysis, total RNA-virus in the serum was extracted using RNeasy mini kit -QIAGEN protocols.

The extracted RNA virus was then converted into complementary DNA (cDNA) refers to ReverTra Ace[®] qPCR Master mix (Toyobo) as previously described by Jadid et al. [6]

for further PCR amplification. The reaction was performed within two rounds of PCR. Detail primers in Table 1 [7].

Table 1. Details of the primers for HCV amplification.

166	TGGGGATCCCGTATGATACCCGCTGCTTTGA	8230-8260	+	Outer
167R	GGCGGAATTCCTGGTCATAGCCTCCGTGAA	8601-8630	-	
HC23	TTTGACTCAACCGTCACTGA	8256-8275	+	Inner
HC24	CTCAGGCTCGCCGCATCCTC	8577-8596	-	
HC26	CTCAGGTTCCGCTCGTCCTC	8577-8596	-	
HC15	ACTGTCACTGAACAGGACAT	8265-8284	+	Inner
HC16	GCTCTATCCTCATCGACGCC	8568-8587	-	
HC28	CACGAGCATGGTGCAGTCCCGGAGC	8507-8531	-	
HC32	AGGTAGCACGTCAGCGTGTTTCC	8454-8476	-	
HC34	TAGCACGTCATGGTGTTTCCCAT	8451-8473	-	

PCR used "GeneAmp PCR System 2400 (Perkin Elmer)" Every PCR conducted using 40 cycles. Before the cycle hot start conducted with temperature of 94°C for 5 min. Condition of the PCR cycles describe below: denaturation at 94°C for 60 s, annealing temperature for 60 s and elongation at 72°C for 75 s.

Amplification result visualization

PCR results were visualized by gel electrophoresis which consist of TBE buffer 0.5X, 2% of agarose gel and supplemented with ethidium bromide. The 1000 bp DNA ladder HAE III Digest was used as DNA marker.

Data analysis

Data population of HCV infected status of blood samples by ELISA methods were qualitatively grouped for further investigation using PCR methods. Positive HCV determination based on PCR were analysed descriptively refer to visual DNA-virus bands.

Result and Discussions

The 100 blood samples were obtained from 85 male and 15 female donors. Blood sample HCV detection (positive and negative status) by ELISA was exhibited in following Table 2.

The ELISA analysis indicated that all donors (100%) were negative HCV. These negative HCV were then administered with PCR analysis.

The PCR results showed that 5 samples (5.8%) among 85 negative samples of male 1 sample (6.7%) among 15 negative samples of male were detected positively infected by HCV. The PCR visualizations of HCV in blood sample were represented Figure 1, which positive one was shown in sample BD-10 (300 bp).

The positive HCV that resulted by PCR analysis mainly depends on the choice of selected primer. In this study, we used primer which was designed based on the sequences of non-structural protein 5B (NS5B) of hepatitis C virus (HCV) [8]. NS5B have a polymerase activity which responsible for viral genome replication [9]. Moreover, due to its capacity, the NS5B has been developed for antivirus therapy. Some studies reported NS5B region can be used to determine genotype and subtype of hepatitis C virus as well as an effective learning tool for molecular HCV epidemiology [10,11]. This case study demonstrated that 6% detection by ELISA were inaccurate. Since the HCV transmission is mediated by blood, this 6% value considerably crucial and highly risk. Accordingly, the validation using PCR for screening blood donors in Tuban region is necessarily required.

Table 2.	Result of and	ti-HCV and	lvsis using	ELISA.
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Gender	Anti HCV		PCR		
	Ratio Negative/Positive	Percentage of negative HCV	Ratio Negative/Positive	Percentage of negative HCV	
Male	85/100	85%	5/85	5.8%	
Female	15/100	15%	1/15	6.7%	
Total	100	100 %	6/100	6%	

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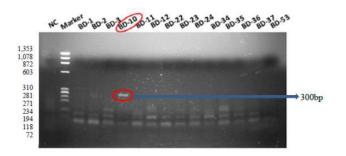


Figure 1. The PCR visualization of HCV in blood sample. The positive HCV shown in sample BD-10 (circled). NC: Negative Control.

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Conclusion

According to HCV analysis using ELISA toward 100 blood donors sample, there was resulted 100% negative HCV. After retested by PCR analysis, there were found 6 blood samples (6%) positively infected HCV. This study encourages who one concerned to propose that PCR-based validation for screening blood donors is necessarily required.

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