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3 **Frequency of Dengue Virus Serotype 1 in Lahore by In-house**  
4 **assay**

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10  
11 **Abstract**

12 Dengue is an important systemic viral infection that is caused by the dengue  
13 virus. Ribonucleic acid (RNA) from dengue NS1 positive samples, collected  
14 randomly during dengue epidemic from October 2016 to October 2017 at  
15 Chughtai Lab, was extracted for nucleic acid. Both the detection and serotyping  
16 of dengue samples were performed using real-time PCR on Rotor Gene Q.  
17 From the 70 NS1 positive samples, 57 (81.4%) samples were confirmed to be  
18 positive for dengue virus RNA, while the remaining 13 (18.6%) were negative.  
19 Serotype 1 (DEN-1) were verified among all samples by in-house assay and  
20 using commercial kit FTD (Fast Track Diagnostics) dengue differentiation; it  
21 was concluded that our in-house assay is in 100% concordance with commercial  
22 kit. Serotype 2 (DEN-2) and serotype 3 (DEN-3) have been documented in  
23 Pakistan since 1994. But recent detection of serotype 1 in Pakistan is indicative  
24 of more severe dengue haemorrhagic fever in future due to reinfection.

25 **Keywords:** Dengue, Real-time PCR, Serotype.

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## 29 **Introduction**

30 Dengue is an important systemic viral infection that is caused by the dengue  
31 virus which is a single-stranded RNA virus belonging to the *Family*  
32 *Flaviviridae*. Dengue virus is classified into four antigenically distinct serotypes  
33 (DEN-1 to DEN-4).<sup>(1)</sup> A fifth serotype DEN-5 was identified in December 2013,  
34 in a farmer in Malaysia; however, it causes only mild febrile illness.<sup>(2)</sup> The main  
35 transmitting vectors of the virus are *Aedes aegypti* and *Aedes albopictus*  
36 mosquitoes.<sup>(2)</sup> The ideal conditions for this mosquito to breed are abundant  
37 rainfall and high humidity, during which the temperature of the surrounding  
38 environment reaches about 30°C. That is why dengue virus infection is common  
39 during September to December. Infection with one serotype of dengue provides  
40 lifelong protection against that serotype, but infection with another serotype  
41 may result in more serious disease.<sup>(3)</sup> Currently, more than 125 countries are  
42 known to be affected by the dengue viruses.

43 Dengue infection has been reported across the Americas, South-East Asia and  
44 Western Pacific regions, affecting millions of people.<sup>(4)</sup> After an incubation  
45 period of two to seven days, the patient experiences flu-like illness followed by  
46 fever, nausea, and vomiting, along with severe frontal and retro-orbital  
47 headache and muscle ache. The most severe and serious secondary infection,  
48 Dengue Haemorrhagic Fever (DHF) is characterised by fever, or recent history  
49 of acute fever with haemorrhagic manifestations and a platelet count of  
50  $100,000/\text{mm}^3$  or less along with objective evidence of “leaky capillaries”; the  
51 haematocrit is elevated (20% or more over baseline) with low albumin levels.<sup>5</sup>  
52 For diagnostic purposes, non-structural protein 1 (NS1) antigen detection and  
53 real-time reverse transcriptase PCR (qRT-PCR) are typically used to detect  
54 dengue viral genes in the viraemic phase on serum or plasma samples. From the  
55 fifth day onwards, detection of IgM and IgG antibodies by ELISA helps in  
56 establishing the diagnosis of infection with the dengue virus. There is no  
57 specific antiviral treatment for dengue and only supportive treatment with

58 analgesics and intravenous fluids is recommended.<sup>(6)</sup> Control of mosquito  
59 vectors by the use of insecticides, removal of artificial water containers and  
60 mosquito screening of household windows are very effective ways to reduce the  
61 outbreaks of dengue.<sup>(7)</sup> The first dengue vaccine, Dengavaxia by Sanofi Pasteur,  
62 was registered in several countries, in 2015-16, for use in people aged 9 to 45  
63 years living in dengue endemic areas. Several other vaccines are in the process  
64 of development. The present study was conducted to develop an in-house assay  
65 and find the prevalence of dengue serotype in Pakistan.

66 **Objective:** To develop an in-house assay for detection and serotyping of  
67 dengue virus by real-time PCR.

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#### 69 **Patients / Methods and Results**

70 A cross-sectional study was conducted on 70 suspected dengue cases that were  
71 sampled during October 2016 to October 2017 at the Molecular Biology and  
72 Virology Department, Chughtai Lab, Lahore, on patients belonging to different  
73 areas of Lahore.

74 Blood plasma was used to extract viral RNA using QIASymphony DSP  
75 Virus/Pathogen Midi Kit (Qiagen) on fully automated platform of  
76 QIASymphony SP (Qiagen) by following the manufacturer's instructions. The  
77 primers used for genotyping of the positive samples were designed according to  
78 the model used by Fatima et al in 2011.<sup>(8)</sup> The region of C-prM gene junction  
79 described by Lanciotti et al<sup>(9)</sup> was selected for genotyping as this is not very  
80 hyper variable and less prone to mutations. The probes for genotyping were  
81 designed by using the tool, Primer3, available online.

82 Detection and genotyping of plasma samples were performed on Rotor-Gene Q  
83 instrument (Qiagen) by adopting the one-step qRT-PCR strategy. In this regard,  
84 a reaction mixture of 20 $\mu$ l was prepared by adding 5 $\mu$ l of extracted viral RNA,  
85 1 $\mu$ l of forward and reverse primers (each 10 picomol/ $\mu$ l), 0.5 $\mu$ l of probe (10  
86 picomol/ $\mu$ l), 2.5 $\mu$ l of PCR water, and 10 $\mu$ l of master mix (Affymetrix USB

87 VeriQuest 2x). Furthermore, the thermal profile used was: an incubation at  
88 50°C for 15 minutes, afterward an initial denaturation at 95°C for 10 minutes  
89 followed by 35 cycles of denaturation at 94°C for 30 seconds, annealing at 52°C  
90 for 30 seconds and extension at 72°C for one minute. A final extension was  
91 given at 72°C for 10 minutes. Results of four serotypes, DEN 1-4, by in-house  
92 assay were further verified through the commercially available FTD dengue  
93 differentiation kit (Fast Track Diagnostics) and were analysed by using software  
94 SPSS 16.0.

95 From the analysis of 70 NS1 positive samples, dengue virus RNA was detected  
96 in almost 57 (81.3%) of the samples, while in the remaining 13 (18.6%) dengue  
97 virus RNA was not detected. (Figure 1). Furthermore, nearly 65 (92.9%) of  
98 these positive cases were sampled from Lahore, whereas 5 (7.1%) were from  
99 outside Lahore. Out of positive dengue patients 45 (64.28%) were males while  
100 25 (35.72%) were females. All these positive samples were subjected to  
101 genotyping by both the in-house assay and the commercially available FTD kit,  
102 and results showed 100% co-relation for serotype 1 (DEN-1). All the patients  
103 had DEN-1, and no other serotype was detected. Ten random positive samples  
104 were sent to the external lab dealing with the viral diagnosis and results were in  
105 100% concordance with the in-house assay.

106 A one way Anova test of independence was performed to check the relationship  
107 between positive cases with gender. The relationship between these variables  
108 was significant. The p-value was less than 0.05 which indicates that these  
109 variables are not independent of each other and that there is a statistically  
110 significant relationship between the categorical real-time PCR output and  
111 gender. The significance value was  $p = 0.048$  with a confidence interval of 95%,  
112 which shows the parameters selected for comparison in this study are relevant  
113 and enlighten the significance of this study.

114 Our country is at high risk of dengue infection due to overcrowding of cities,  
115 presence of stagnant water, a large number of refugees and people exposed to

116 mosquito bite. The first documented outbreaks in Karachi in 1994 and 1995  
117 were reported to be due to dengue serotypes 1 and 2, whereas in 2005 and 2006  
118 outbreaks serotypes 2 and 3 were documented.<sup>(10)</sup> Whereas, in Lahore serotypes  
119 2, 3 and 4 were reported to cause outbreak in 2008, and later in 2009 only  
120 serotypes 2 and 3 were responsible for major outbreaks in Lahore.<sup>(11)</sup> In the  
121 current study, serotype 1 was documented for causing outbreak in Lahore 2016,  
122 with a p-value of 0.048, showing relevance of this study. The serotype 1 has re-  
123 emerged after 21 years of its first evidence in Karachi, 1994 and then in 1998.  
124 DEN-3 and DEN-2 was reported in Karachi in 2006. Drift change in serotype  
125 poses a great risk of dengue haemorrhagic fever in population already exposed  
126 to other dengue serotype.

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### 128 **Conclusion**

129 Dengue serotype is changing quite dramatically over the time in Pakistan,  
130 because of which greater incidence of dengue haemorrhagic fever is predicted in  
131 population previously exposed to other dengue serotype.

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134 **Conflict of Interest:** None

135 **Sources of Funding:** None

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### 137 **References**

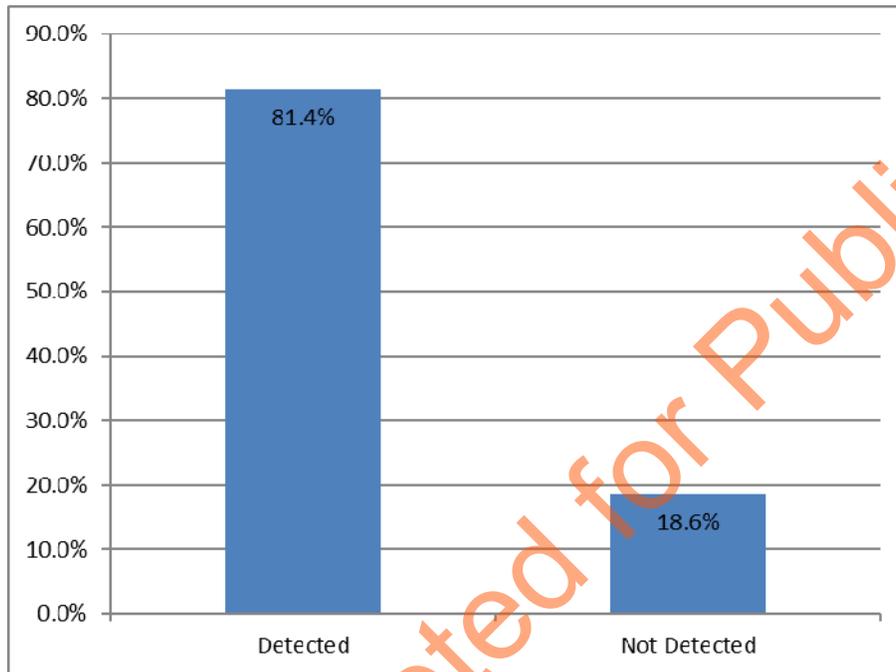
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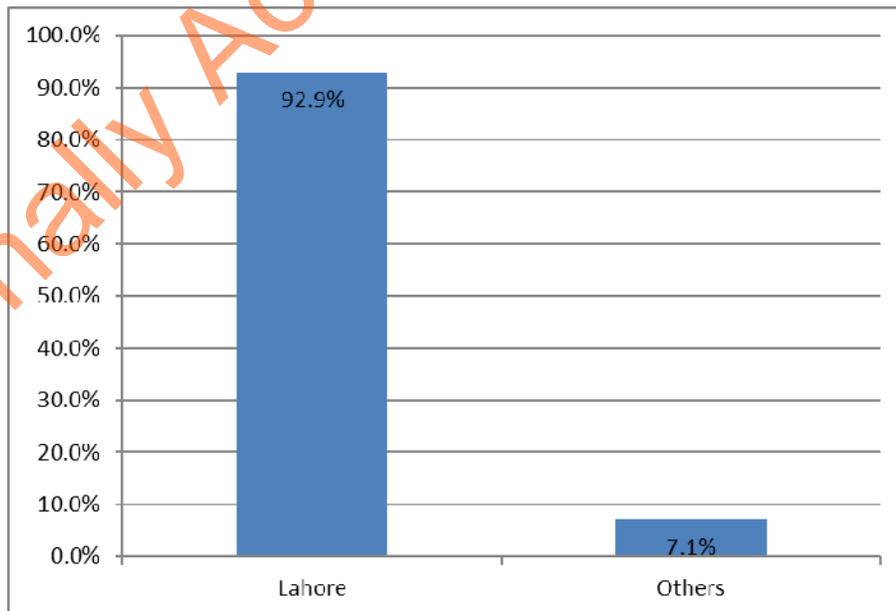
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**Figure 1: Real-time PCR results of suspected dengue samples. About 81.4 % of the samples were found to be RNA positive for dengue virus.**



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**Figure 2: Area wise distribution of RNA positive dengue cases. Almost 93% of cases sampled were from the Lahore region.**