

# **Prevalence of Untypable HCV Variants in Different Districts of Punjab, Pakistan and a Case Report of its Treatment Strategy**



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# **Prevalence of untypable HCV variants in different districts of Punjab, Pakistan and a case report of its treatment strategy**

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**BY**

**AYESHA ZAFAR**

**ID**

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**Department of Life Sciences**



بِسْمِ اللَّهِ الرَّحْمَنِ الرَّحِيمِ

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**AYESHA ZAFAR**

# DEDICATION

I dedicate

My work to

MY BELOVED PARENTS

MR. AND MRS. ZAFAR TARIQ

And to my siblings for their endless support during my whole educational career.

## DECLARATION

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I AYESHA ZAFAR D/O MUHAMMAD ZAFAR TARIQ ID: 15001254009 Session 2015-2017 declare that the matter printed in the thesis dissertation titled “Prevalence of untypable HCV variants in different districts of Punjab, Pakistan and a case report of its treatment strategy” which I here with submit for the research qualification **M.S Degree In Biotechnology** to the Department of Life Sciences, School of Science, University of Management and Technology, Lahore, Pakistan is apart from the recognized assistance of my supervisors, my own work and has not been previously submitted as research work, thesis or publication in any form in any university, research institution etc. in Pakistan or abroad to obtain a research diploma or degree.

Dated: \_\_\_\_\_

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AYESHA ZAFAR

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## ABSTRACT

In Pakistan, around 11 million people are HCV infected with 6 circulating genotypes and subtypes in Pakistan and the viral burden is still on rise. HCV genotype determination is necessary to help the health care facilitators for recommending the appropriate antiviral therapy. This large scale study was carried out to investigate the epidemiological distribution of genotypes in different districts of Punjab, Pakistan and focuses on prominently increasing one (diagnostically untypable) and its possible antiviral therapy. We have also characterized an untypable HCV variant by sequence and phylogenetic analysis that showed the greatest homology of untypable HCV variant with 3a genotype from Pakistan (Unpublished data). A case study has been followed up with untypable HCV genotype for 24 weeks. Of the 8353 patients enrolled in study, the genotype 3 and subtype 3a (**n= 6650, 79.6%**) was most significantly prevalent throughout the study followed by undetermined genotype i.e untypable (**n= 1377, 16%**) by the diagnostic method of Ohno *et al.* The presence of untypable variants suggested that there are some novel or quasi species prevailing among HCV patients in local community. Case of patient with untypable genotype had been treated effectively with interferon alpha plus ribavirin therapy for 24 weeks and achieved SVR rate of 70-80 %. From the study, it can be concluded that untypable variants can be successfully treated with standard interferon therapy unless the mystery of untyped genotype has been solved with novel and effective diagnostic methods

## INTRODUCTION

### 1.1 Hepatitis

Viral hepatitis is illness of the liver caused by a virus. Hepatitis C is a long haul disorder, around 20-30 years with a moderate development. In hepatitis C, 80% of intense initial infections change to chronicity. The consequences of HCV disease can be resolved within a half year of disease exposure. Most of the HCV patients may develop cirrhosis and, eventually, hepatocellular carcinoma. Around 80% of HCV diseases prompt chronicity which are predominantly because of age, male sex, stoutness, liquor consumption and co-contamination with HBV (hepatitis B infection) or HIV (human immune-deficiency virus) (Vaquer *et al.*, 1994), development of viral quasi-species that escape immune resistance (Farci *et al.*, 2000) and deficient cell reactions (Bowen and Walker, 2005).