STUDY OF THE PREVELANCE OF URTICARIA IN CHRONIC RENAL DIALYSIS EGYPTIAN PATIENTS WITH HCV INFECTION

Thesis

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INTRODUCTION

Hepatitis C virus (HCV) is a major health problem; it is the main cause of transfusion-associated hepatitis and is also seen in intravenous drug abuse, organ transplant and hemodialysis patients, and health care workers (Neil et al., 2003).

In addition to hepatitis, cirrhosis and hepatocellular carcinoma (HCC), several extrahepatic manifestations (EHM) have been reported in the natural history of HCV infection. According to different studies, 40-74% of patients infected with HCV might develop at least one EHM during the course of the disease (Galossi et al., 2007).

A significant proportion of these extrahepatic manifestations (EHM) disorders involve the skin (*Davis et al.*, 2003). Various cutaneous eruptions have been described in the setting of HCV infection (*Neil et al.*, 2003).

Urticaria is a common disorder that affects up to 20% of the population at some point during their lifetime. Aggravating factors include drugs, foods, additives, connective tissue disorders and infections. It is well established that Hepatitis B virus causes urticaria. Whether Hepatitis C infection causes urticaria or not is still debated with reports both in favor of and against this (Siddique et al., 2004).

AIM OF THE WORK

To analyze the frequency of urticaria in chronic renal dialysis Egyptian patients with HCV infection.

LIST OF ABBREVIATIONS

AASLD American Association for the Study of Liver

Diseases

ALT Alanine aminotransferase AST Aspartate aminotransferase

CDC Center of Disease Control

CDNA Complementary DNA

CG Cryoglobulinemia

CI Confidence Interval

CKD Chronic kidney disease

CLD Chronic liver disease

DNA Deoxyribonucleic acid

EDHS Egyptian Demographic Health Survey

EIA Enzymatic immune assay

EHM Extra hepatic manifestations

ERBP European Renal Best Practice Guidelines

ESRD End stage renal disease

ETR End of treatment response

EVR Early viral response

FDA Food and Drug Administration

GFR Glomerular filtration rate

GN Glomerulonephritis

HAI History activity index

HAV Hepatitis A virus

HBV Hepatitis B virus

HCC Hepatocellular carcinoma

HCM Hypertrophic cardiomyopathy

HCV Hepatitis C virus

HCVcAg Hepatitis C core antigen

HD Hemodialysis

HIV Human immune deficiency virus

HTLV Human T-lymphotropic virus

IDE Integrated development environmentIDSA Infectious Disease Society of America

IFN Interferon

Ig Immunoglobulin IR Insulin resistance

IRES Internal ribosome entry site

IVDU Intra venous drug users

KDIGO Kidney disease improving global outcome KDOQI Kidney disease outcomes quality initiative

LHF Lassa haemorrhagic fever MENA Middle East North Africa MHD Maintenance hemodialysis

MICS Malnutrition inflammation cachexia syndrome

MIEC Medanta independent ethics committee

MPGN Membrano proliferative glomerulonephritis MPGN Mesangio proliferative glomerulo nephritis

MU Million units

NANB Non- A Non – B

NAT Nucleic acid testing

NHANES National Health and Nutrition Examination Survey

NHL Non-Hodgkin lymphoma

NSAID Non steroid anti- inflammatory drugs

OR Odd ratio

PAT Parentral antischistosomal therapy

PBMC Peripheral blood monocytes

PBMC Periphral blood mononuclear cell

PCR Polymerase chain reaction

PCT Porphyria cutanea tarda

PD Peritoneal Dialysis

PN Peripheral neuropathy

RA Rheumatoid arthritis

RBV Ribavirin

RdRp RNA dependant RNA polymerase

RF Rheumatoid Factor

RIBA Reverse immuno- blot assay

RNA Ribonucleic acid

RR Relative risk

RRT Renal replacement therapy

RT-PCR Reverse transcriptase polymerase chain reaction

RVR Rapid virological response

SLE Systemic lupus eryromatosus

SOC Standard of care

SPT Skin prick test

SS Sjogren's syndrome

SVR Sustained viral response

TIPSS Transjugular intrahepatic portosystemic shunt

TMA Transcription mediated amplificationUSPHS United States Public Health Service

UTR Untranslated region

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HEPATITIS C VIRUS (HCV)

Definition

HCV, first identified in 1989, is strictly a blood-borne RNA viral infection in the family Flaviviridae. Humans are the only reservoir for this viral infection. HCV infection most often leads to an asymptomatic chronic state, which can later progress to active liver disease, liver failure, or primary hepatocellular carcinoma. Treatment of HCV is costly, beyond the reach of most patients in less-developed countries, requires 48 or more weeks to complete, and has serious adverse effects and low efficiency. HCV in a family member can be socially and economically detrimental (Millera and Abu Rhaddad, 2010).

Egypt reports the highest prevalence of HCV worldwide, ranging from 11% to more than 14% among regions and demographic groups. In the U.S., the number of new cases of infection with HCV has declined over the last 10 years from a peak of some 200,000 annually to about 19,000 in 2006. Up to 85% of newly infected people fail to clear the virus and become chronically infected. In the U.S., more than three million people are chronically infected with HCV. Infection is most common among people who are 40 to 60 years of age, reflecting the high rates of infection in the 1970s and 1980s. There are 8,000 to 10,000 deaths each year in the U.S. related to HCV. HCV is the leading cause of liver transplantation in the U.S and is a risk factor for liver cancer (*Dubuisson*, 2007)

HCV is a small single-stranded RNA virus with a lipid envelope (E) containing glycoproteins (E1 and E2) and a core with a genome consisting of 9500 nucleotides. HCV components are both structural (core, E1, and E2) and nonstructural (NS; P7, NS2, NS3, NS4A, NS4B, NS5A, and NS5B). The nonstructural genes encode various enzymes including a polymerase responsible for replication of HCV (KDIGO Guidelines, 2008).

The structural proteins include the core (C), which forms the viral nucleocapsid, and the envelope glycoproteins E1 and E2. They are released by host-cell signal peptidases. The structural proteins are separated from the nonstructural proteins by the short membrane peptide p7, thought to be a viroporin. The nonstructural (NS) proteins NS2 to NS5B are involved in polyprotein processing and viral replication. The proteolytic processing of NS polyprotein part is complex and requires two distinct proteinases: the NS2- NS3 zinc-dependent metalloproteinase, and the NS3 serine proteinase located in the N-terminal region of NS3. The NS2-NS3 proteinase appears to be dedicated solely to cleavage at the NS2/NS3 site that occurs rapidly and by a conformation-dependent, autocatalytic mechanism (*Penin et al.*, 2004).

The remaining NS proteins are released by the NS3 proteinase associated with its cofactor, NS4A. The C-terminal region of NS3 protein includes RNA helicase and NTPase activities. NS4B is an integral membrane protein of unknown function. NS5A is a polyphosphorylated protein of unknown function, and NS5B is the RNA-dependent RNA polymerase

(RdRp). The existence of one or more previously unknown HCV proteins potentially synthesized by ribosomal frame shift has been suggested recently *(Choi et al., 2003)*.

Although this basic structure is common to all hepatitis C viruses, there are at least six distinctly different strains of the virus which have different genetic profiles (genotypes). In Egypt genotype 4 is the most common form of HCV. In the U.S., genotype 1 is the most common form of HCV. Even within a single genotype there may be some variations (genotype 1a and 1b, for example). Genotyping is important to guide treatment because some viral genotypes respond better to therapy than others. The genetic diversity of HCV is one reason that it has been difficult to develop an effective vaccine since the vaccine must generate viral proteins from each genotype (*Dev et al., 2002*).

HCV Prevalence in Egypt

The highest HCV prevalence in the world occurs in Egypt, where the prevalence of infection increases steadily with age, and high rates of infection are observed among persons in all age groups (*Perz et al., 2006*). This pattern indicates an increased risk in the distant past followed by an ongoing high risk for acquiring HCV infection, although there are regional differences in average overall prevalence (*Medhat et al., 2002*).

In 1992, when HCV antibody testing became widely available, the prevalence of HCV in Egypt was reported to be 10.8% among first-time blood donors (*Millera and Abu Raddad*, 2010). Since this discovery, many prevalence estimates of HCV