Hepatitis C Virus multiple genotype infection

د ابتسام البوشي

مشفى دمشق

حالة سريرية

المريضة: ن.ر العمر: 45 سنة الحالة الإجتماعية: متزوجة السكن: درعا *

: سوابق مرضية *ESRD معالجة بالتحال الدموي

Anti-HCV : Neg AST: N ALT: N* مخبريا عند بدء التحال

عند اجراء مراقبة روتينية بعد 3 أشهر من بدء التحال الدموي *

أجري للمريضة * Anti-HCV وكان إيجابي أجري للمريضة * Pcr-HCVوكان ايجابي أجري للمريضة *Genotype-HCVوكان من النمطين (1-4)

Detection, genotyping and the RNA-HCV viral load

- The viral RNA was extracted from 200 ul sample of plasma using Roche MagNA pure (automated extraction).
- RNA sample were tested with negative control on Roter-Gene machine using Sacace Real Time HCV genotype.

Epidemiol. Infect. (2009), 137, 79–84. © 2008 Cambridge University Press doi:10.1017/S095026880800054X Printed in the United Kingdom

The unexpected discovery of a focus of hepatitis C virus genotype 5 in a Syrian province

Table 1. Geographical distribution of HCV genotypes in Syria

Genotype	North (n=207)	East (n=255)	Centre (n=67)	South $(n=95)$	Undetermined $(n=12)$	Total (n=636)
1	50 (24·1 %)	19 (7.5%)	44 (65%)	64 (67%)	4	181 (28.5%)
2	4 (2%)	0	1 (1.5%)	0	0	5 (0.8%)
3	2(1%)	3 (1%)	0	5 (5.3%)	1	11 (1.8%)
4	95 (46%)	227 (89%)	21 (31%)	25 (26.3)	7	375 (59%)
5	56 (27%)	6 (2.4)	1 (1.5%)	1(1%)	0	64 (10%)

Journal of Viral Hepatitis

Wiley-Blackwell, John Wiley & Sons

Prevalence of mixed genotype hepatitis C virus infections in the UK as determined by genotype-specific PCR and deep sequencing

- A. L. McNaughton, V. B. Sreenu, [...], and
- E. C. M. Leitch

Mixed HCV infection in UK

• The prevalence rate of mixed infection in the UK cohort of 506 individuals by PCR was 3.8%, with genotype 3 as the major genotype.

 The mixed infection rate obtained from PCR-NGS data was much higher.





Incident Hepatitis C Virus Genotype Distribution and Multiple Infection in Australian Prisons

Melanie R. Walker, Hui Li, Suzy Teutsch, Brigid Betz-Stablein, Fabio Luciani, Andrew R. Lloyd, Rowena A. Bull, on behalf of the HITS-investigators

In Australia

Most common genotype is 3.

• Multiple genotype infection is common in high-risk populations.

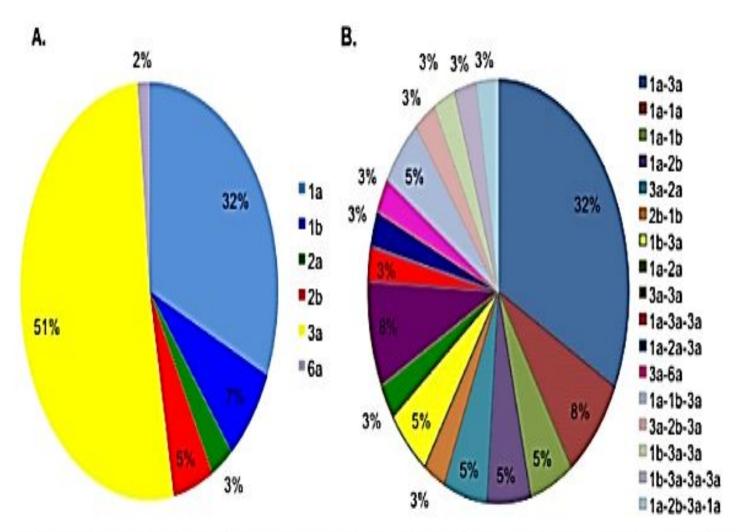


FIG 1 Distribution of HCV subtypes and multiple-infection combinations. (A) Distribution of subtypes in the primary incident HCV infection samples in the cohort, with different subtypes represented by different colors as shown in the key. (B) A varied set of multiple-infection combinations were observed. Each multiple-infection combination is represented by a different color as shown in the key, with the percentage of each set as a proportion of the total also represented.

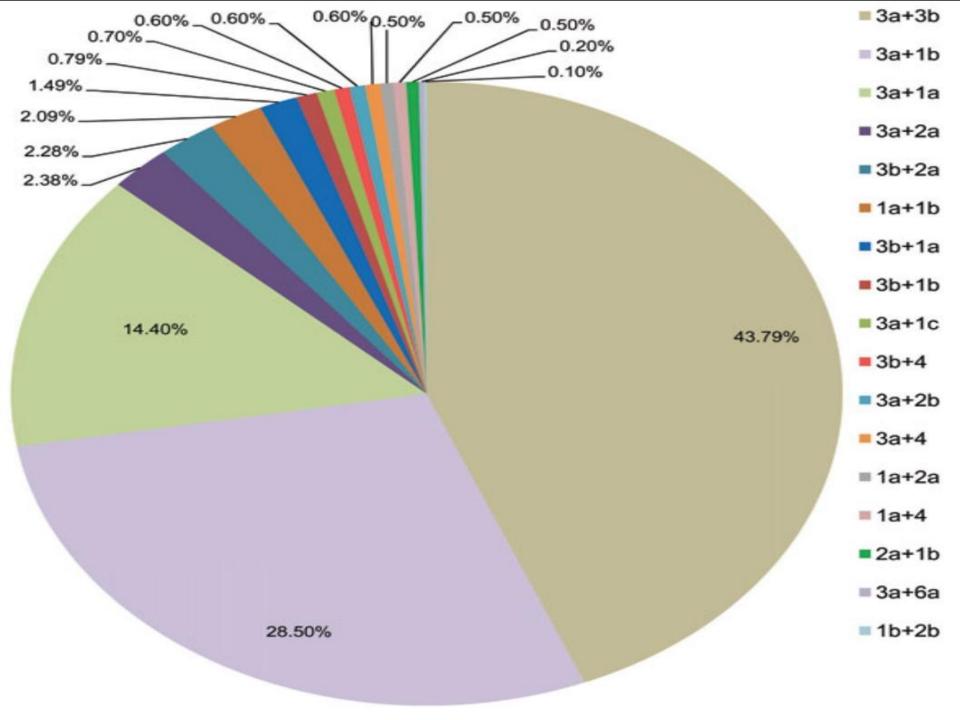
Volume 17, Number 8—August 2011

Letter

Mixed Genotype Infections with Hepatitis C Virus, Pakistan

Sadia Butt, Muhammad Idrees, , Irshad Ur Rehman, Haji Akbar, Muhammad Shahid, Samia Afzal, Saima Younas, and Iram Amin Author affiliations: Author affiliation: University of the Punjab, Lahore, Pakistan

In conclusion, the prevalence of HCV mixed genotype infections in Pakistan is higher than previously reported and higher among men



Why it's important to look for multi genotype HCV infection

- In patients serially infected by different HCV strains, one strain will prevail as the viremic virus.
- Under antiviral therapy, the displaced strain may influence the outcome of therapy.
- detection of inferior strains by serological assays before antiviral therapy may be important for choosing the adequate regimen.

Article | Open Access | Published: 25 June 2019

Infection with multiple hepatitis C virus genotypes detected using commercial tests should be confirmed using next generation sequencing

• HCV is a blood-borne pathogen, and transmission modes of the disease are parenteral and to a lesser extent sexual. and sharing toothbrushes/razors.

 Patients in hemodialysis units have an increased risk for contracting HCV.

• The ESRD patients with one genotype HCV infection have a high risk of developing infection with another HCV genotype(super infection) from dialysis centers.

- There are many reasons associated with HCV transmission within a facility including: the dialysis machine may have a role, transfusion during dialysis, the duration of hemodialysis.
- Failure to clean and disinfection of equipment and environmental surfaces, adherence to hand hygiene and glove use, vascular access care, and medication preparation and administration.

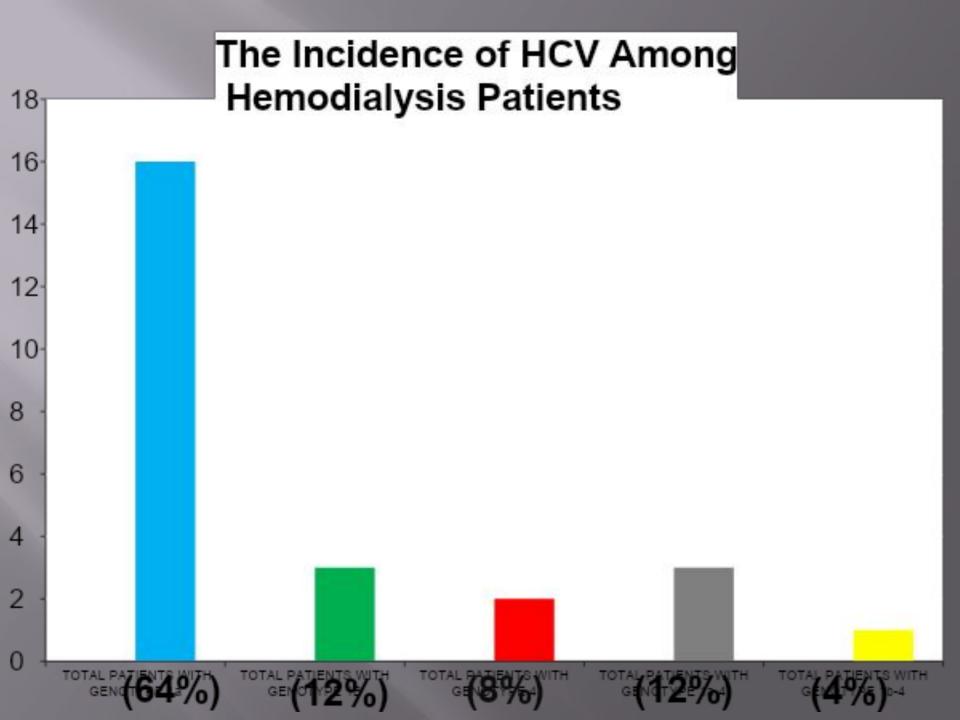
• All ESRD patients on dialysis should be screened for anti-HCV antibodies at the time of admission to dialysis center.



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Study in Damascus hospital

 We investigate the super infection with two genotypes Hepatitis C virus (HCV) infection in 25 patients with ESRD and found that 4 of 25 patients (16%) have two genotypes {genotypes 1a -4 found in 3 patients (12%) and genotypes 1b - 4 found in one patient (4%)} from different dialysis centers around Syria.



 Solutions to prevent transmission of HCV in dialysis facilities; adherence to recommended infection control and practicing are needed.

 training the dialysis staff routinely in a program for preventing infection, Follow CDC recommendations for HCV screening of hemodialysis patients.

evaluate the facility routinely for infection.

Conclusion

 We must do more researches in all around Syria to detect the multi genotype infection with HCV and to make precise percentage of the prevalence of all genotypes.

We must use the NGS routinely in Syria if possible

• Treat your patients with broad spectrum DAA.

• Follow CDC recommendations for HCV screening of hemodialysis patients.