

Table 1 Patient characteristics according to hepatitis C virus genotype of infection

Variable	Genotype 1 (n = 79) (19.8%)		Genotype 4 (n = 256) (64.0%)		Genotypes 2/3 (n = 65) (16.3%)		P-value
Age [mean (SD) years]	44.9	(10.3)	47.5	(9.9)	46.2	(11.0)	0.51
Sex, male (no., %)	58	73.4	213	83.2	45	76.9	0.03
BMI [mean (SD) kg/m ²]	28.5	(4.8)	29.3	(4.4)	29.0	(6.1)	0.59
Risk factors present ^a (no. %)	59	74.7	164	64.1	35	53.8	0.06
High fibrosis score (F3,F4) (no. %)	26/66	39.3	74/233	31.8	20/56	36.4	0.68
ALT ^b (no. %)							0.69
Normal	19	24.1	53	20.7	16	24.6	
Near normal	29	36.7	94	36.7	19	29.2	
Abnormal	31	39.2	109	42.6	30	46.2	
Pretreatment viral load ^c (no. %)							
Low	43	54.4	173 ^d	67.6	37	56.9	0.001
High	36	45.6	81	31.6	28	43.1	
Geographic origin of infection (no. %)							
Middle East	41	51.9	31	12.1	15	23.1	0.001
Egypt	28	35.4	131	81.2	2	3.1	
South-East Asia	7	8.9	14	5.5	46	3.8	
Other	3	3.8	6	2.3	24	36.9	

^aAt least 1 known risk factor for transmission of infection identified.

^bNormal < 40 IU/mL; near normal 41–80 IU/mL; abnormal > 80 IU/mL.

^cHigh viral load > 700 × 10³ IU/mL; low viral load ≤ 700 × 10³ IU/mL.

^dData missing for 2 patients.

BMI = body mass index; SD = standard deviation; ALT = alanine aminotransferase.