

**Table 2** Epidemiological and clinic features of infectious diseases among immigrant patients according to region of origin, Italy, 2015

Epidemiological and clinic features	Total No.	Origin								P-value <sup>a</sup>	
		Africa (Af)	East Africa	West Africa	Central Africa	Asia (As)	Pakistan	Bangladesh	India		Other countries
HBsAg+	23	17	6	4	7	6	2	2	1	1	0.03 <sup>b</sup>
HBeAg+	7	3	2	1		4	1	1	1	1	
HBV-DNA+, HBsAg+, OBI	17	13	4	5	4	4		2	1	1	0.03 <sup>b</sup>
Genotype	12										
A	3	3	2	1							
B	1						1				
C	1									1	
D	2				2						
E	5		1	3	1						0.02 <sup>c</sup>
CHB	12	10	2	3	5	2	1			1	0.03 <sup>b</sup>
OBI	77	51	19	16	16	26	11	3	5	7	0.03 <sup>b</sup>
HCV+	8	3	1	2		5	1	1	3		0.02 <sup>d</sup>
CHC	3					3	1		2		
Genotype	3										
1						2	1		1		
3						1	1				
HIV+	2	1				1					
Virological coinfection	5	2 (1 B, 1 C)				2 (1 B, 1 C)					
LTBI TST+	21	15	3	7	5	6	1	3		2	0.02 <sup>b</sup>
TB & hepatitis coinfection	5	3	1	1	1	2		2			

OBI = anti-HBc; CHB = chronic active HBV; LTBI = latent tuberculosis infection; TST = tuberculin skin test.

<sup>a</sup>When the value of P is not specified, the parameters compared are not statistically significant.

<sup>b</sup>Af vs As.

<sup>c</sup>E vs all.

<sup>d</sup>As vs Af.