

Supplementary Materials

Epidemiological and Genetic Characterization of Norovirus Outbreaks That Occurred in Catalonia, Spain, 2017–2019

Table S1. Prevalence of genotypes identified during the study period per year.

Genotype	N. of Outbreaks	2017 (%)	2018 (%)	2019 (%)
GI.1[P1]	5	0 (0%)	3 (8.82)	2 (5.13)
GI.3[P3]	6	0 (0%)	0 (0%)	6 (15.38)
GI.3[P13]	2	1 (3.7)	1 (2.94)	0 (0%)
GI.4[P4]	6	1 (3.7)	4 (11.76)	1 (2.56)
GI.5[P4]	1	0 (0%)	0 (0%)	1 (2.56)
GI.5[P5]	1	0 (0%)	1 (2.94)	0 (0%)
GI.6[P11]	3	1 (3.7)	2 (5.88)	0 (0%)
GII.1[P33]	1	1 (3.7)	0 (0%)	0 (0%)
Multiple GI genotypes ^a	1	1 (3.7)	0 (0%)	0 (0%)
GII.2[P16]	11	7 (25.93)	2 (5.88)	2 (5.13)
GII.3[P21]	1	0 (0%)	1 (2.94)	0 (0%)
GII.3[P30]	3	0 (0%)	0 (0%)	3 (7.69)
GII.4[P4] Sydney 2012	2	1 (3.7)	0 (0%)	1 (2.56)
GII.4[P16] Sydney 2012	11	4 (14.81)	3 (8.82)	4 (10.26)
GII.4[P31] Sydney 2012	14	4 (14.81)	2 (5.88)	8 (20.51)
GII.5[P40]	3	0 (0%)	0 (0%)	3 (7.69)
GII.6[P7]	6	0 (0%)	5 (14.71)	1 (2.56)
GII.8[P8]	1	0 (0%)	0 (0%)	1 (2.56)
GII.10[P16]	1	0 (0%)	0 (0%)	1 (2.56)
GII.14[P7]	1	0 (0%)	1 (2.94)	0 (0%)
GII.17[P17]	9	2 (7.41)	5 (14.71)	2 (5.13)
Multiple GII genotypes ^b	1	0 (0%)	1 (2.94)	0 (0%)
Multiple GI + GII genotypes ^c	3	2 (7.41)	1 (2.94)	0 (0%)
Untyped	7	2 (7.41)	2 (5.88)	3 (7.69)
Total	100	27	34	39

^a GI.3[P13]/GI.2[Puntyped]; ^b GII.2[P16]/GII.3[P30]; ^c GII.4[P16] Sydney 2012/GI.4[P4], GII.17[P17]/GI.3[P13] and GII.4[P4] Sydney 2012/GI.6[P11].