




## Correction to: Genetic Diversity Among Genogroup II Noroviruses and Progressive Emergence of GII.17 in Wastewaters in Italy (2011–2016) Revealed by Next-Generation and Sanger Sequencing

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The original version of this article unfortunately contained a mistake. The presentation of Table 1 was incorrect. The corrected table is given below. The original article has been corrected.

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The original article can be found online at <https://doi.org/10.1007/s12560-017-9328-y>.

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**Table 1** Primers used in the present study

Region	Primer name	Primer (5'–3')	Length (bp)	Cycle	References
<i>Real-time RT-(q)PCR for GII NoV</i>					
ORF1–ORF2 junction	QNIF2	ATGTTTCAGRTGGATGAGRTTCTCWGA	–	One-step real-time	Kageyama et al. (2003), Loisy et al. (2005)
	COG2R	TCGACGCCATCTTCATTCACA			
	QNIFs	FAM-AGCACGTGGGAGGGCGATCG-TAMRA			
<i>Broad-range GII NoV nested RT-PCR</i>					
ORF2	COG2F	CARGARBCNATGTTYAGRTGGATGAG	398	1st cycle	Kageyama et al. (2003), Kojima et al. (2002)
	G2SKR	CCRCCNGCATRHCCRTTRTACAT			
	G2SKF	CNTGGGAGGGCGATCGCAA	344	2nd round	
	G2SKR	CCRCCNGCATRHCCRTTRTACAT			
<i>Specific GII.17 NoV nested RT-PCR</i>					
ORF2	2063	GCTCCATCTAATGATGGTG	368	1st cycle	La Rosa et al. (2017)
	2062	TCCACTGGGAAATTTGGC			
	2104	TGGTCTCGTACCAGAGGG	310	2nd round	
	2064	GATCTTCCGGCAGTGAAC			
<i>GII.17 ORF1-ORF2 junction nested RT-PCR</i>					
ORF1–ORF2 junction	GIIPF800 M	GATGCWGAYTAYTCYMGNTGGGA	2354	1st cycle	Koo et al. (2017)
	GIICR1450	ACCCARGMNTCAAAYCTRAART			
	GIIPF750 M	CNGCHHTAGARRTNATGGT	1048	2nd round	
	GII-R1 M	CCRCCNGCATRNCCRTTRTACAT			

Real-time probe labeled at the 5'-end with 6-carboxyfluorescein (FAM) and at the 3'-end with 6-carboxytetramethylrhodamine (TAMRA)