

Erratum

Erratum to “Quantification of Human and Animal Viruses to Differentiate the Origin of the Fecal Contamination Present in Environmental Samples”

Sílvia Bofill-Mas, Marta Rusiñol, Xavier Fernandez-Cassi, Anna Carratalà, Ayalkibet Hundesa, and Rosina Girones

Laboratory of Viruses Contaminants of Water and Food, Department of Microbiology, Faculty of Biology, Avenida Diagonal 643, 08028 Barcelona, Catalonia, Spain

Correspondence should be addressed to Sílvia Bofill-Mas; sbofill@ub.edu

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The primer Q-PAdV-R was mismatched with probe Q-PAdV-P in Table 1. The table is corrected here.

TABLE 1: Oligonucleotide primers and probes used for the detection and quantification of viral indicators.

Primers and probes	Virus	Position ^a	Reference	Sequence (5'-3')
ADF		18869–18887		CWTACATGCACATCKCSGG
ADR	Human adenovirus (HAdV)	18919–18937	[26]	CRCGGGCRAAYTGCACCAG
ADP1		18889–18916		FAM-CCGGGCTCAGGTACTCCGAGGCGTCCT-BHQ1
JE3F		4317–4339		ATGTTGCCAGTGATGATGAAAA
JE3R	JC polyomavirus (JCPyV)	4251–4277	[27]	GGAAAGTCTTTAGGGTCTTCTACCTTT
JE3P		4313–4482		FAM-AGGATCCCAACACTCTACCCCCACCTAAAAAGA-BHQ1
QB-F1-1		2122–2144		CTAGATCCTACCCCTCAAGGGAAT
QB-R1-1	Bovine polyomavirus (BPyV)	2177–2198	[28]	TTACTTGGATCTGGACACCAAC
QB-P1-2		2149–2174		FAM-GACAAAGATGGTGTATCCTGTTGA-BHQ1
Q-PAdV-F		20701–20718		AACGGCCGCTACTGCAAG
Q-PAdV-R	Porcine adenovirus (PAdV)	20751–20768	[29]	AGCAGCAGGCTCTTGAGG
Q-PAdV-P		20722–20737		FAM-CACATCCAGGTGCCGC-BHQ1
qOv_F				CAGCTGYAGACATTGTGG
qOv_R	Ovine polyomavirus (OPyV)	VP1 ^b region	[30]	TCCAATCTGGGCATAAGATT
qOv_P				FAM-ATGATTACCAAGCCAGACAGTGGG-BHQ1
Q-PaV-F	Chicken/turkey parvovirus (ChPV/TuPV)	3326–3345		AGTCCACGAGATTGGCAACA
Q-PaV-R		3388–3407	[31]	GCAGGGTAAAGATTTCACG
Q-PaV-Pr		3356–3378		6FAM-AATTATTGAGATGGCGCCCACG-BHQ1

^aThe sequence positions are referred to strains J019171 (HAdV), NC_001699.1 (JCPyV), D13942 (BPyV), AJ237815 (PAdV), and GU214706 (ChPV/TuPV) from Genbank. ^bVP1: virion protein 1.