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

Received 13 May 2014; Accepted 21 July 2014; Published 28 August 2014

Copyright © 2014 Sílvia Bofill-Mas et al. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

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	Human adenovirus (HAdV)	18869–18887	[26]	CWTACATGCACATCKCSGG
ADR		18919–18937		CRCGGGCRAAYTGCACCAG
ADP1		18889–18916		FAM-CCGGGCTCAGGTACTCCGAGGCGTCCT-BHQ1
JE3F	JC polyomavirus (JCPyV)	4317–4339	[27]	ATGTTTGCCAGTGATGATGAAAA
JE3R		4251–4277		GGAAAGTCTTTAGGGTCTTCTACCTTT
JE3P		4313–4482		FAM-AGGATCCCAACACTCTACCCACCTAAAAAGA-BHQ1
QB-F1-1	Bovine polyomavirus (BPvV)	2122–2144	[28]	CTAGATCCTACCCTCAAGGGAAT
QB-R1-1		2177–2198		TTACTTGGATCTGGACACCAAC
QB-P1-2		2149–2174		FAM-GACAAAGATGGTGTGTATCCTGTTGA-BHQ1
Q-PAdV-F	Porcine adenovirus (PAdV)	20701–20718	[29]	AACGGCCGCTACTGCAAG
Q-PAdV-R		20751–20768		AGCAGCAGGCTCTTGAGG
Q-PAdV-P		20722–20737		FAM-CACATCCAGGTGCCGC-BHQ1
qOv_F	Ovine polyomavirus (OPyV)	VP1 ^b region	[30]	CAGCTGYAGACATTGTGG
qOv_R				TCCAATCTGGGCATAAGATT
qOv_P				FAM-ATGATTACCAAGCCAGACAGTGGG-BHQ1
Q-PaV-F	Chicken/turkey parvovirus (ChPV/TuPV)	3326–3345	[31]	AGTCCACGAGATTGGCAACA
Q-PaV-R		3388–3407		GCAGGTAAAGATTTTCACG
Q-PaV-Pr		3356–3378		6FAM-AATTATTCGAGATGGCGCCACG-BHQ1

^aThe sequence positions are referred to strains J01917.1 (HAdV), NC_001699.1 (JCPyV), D13942 (BPvV), AJ237815 (PAdV), and GU214706 (ChPV/TuPV) from Genbank. ^bVP1: virion protein 1.