

Electronic Supplementary Material

Human enteric viruses – potential indicators for enhanced monitoring of recreational water quality

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Table S3. Sequencing results of randomly selected amplicons detected during the seasonal study.

Month	Site	BLAST	Match	E-Value	Identity	Sequence
October	F+ 2, 13	JF719736.1	Enterobacteria phage Qbeta isolate Qbeta_2_FR	1.00E-49	98%	CCGCGTGGGG TAAATCCCAC TAACGGTGTT GCCGCGCTTT CACAAGCGGG TGCAGTTCCT GCGCTGGAGA AGCGTGTTAC CGTTTCGGTA TCTCAGCCTT CTCGCAATCG TAAGAA
October	EnV 4	DQ995647.1	Human coxsackievirus A22 strain ban99-10427	6.00E-54	94%	A CATGGTGTGA AGAGTCTATT GAGTACTTG AGAGTCCTCC GGCCCCTGAA TGCGGCTAAT CCTAACCATG GAGCAAGTGC TTACTAACCA GTAAGTAGCT TGTCGTAACG CGTAAGTCCG TGGCGGAACC GACTATTTTG GA
October	EnV 4	GU236263.1	Human echovirus 24 isolate 23927 5' UTR	6.00E-59	96%	CCAAAGTAGT CGGTTCCGCT GCAGAGTTGC CCGTTACGAC AGACTGCCCA CTGGCTCGTA GGTATCTGCT CCGCAGTTAG GATTAGCCGC ATTCAGGGGC CGGAGGACTA CTAAC TAGCT CAATATACTC TTCACACCAT GT
October	EnV 4	AY302549.1	Human echovirus 25 strain JV-4 complete genome	3.00E-47	92%	C CAAGGTAGTC GGATCCGCTG CAGAGTTGCC CGTTACCACA GACTGCCAC TGGCTCGTAT GTATCTGCTC CGCAGTTAGG ATTAACCGCA TTCAGGGGCC GGAGGACTAC TAACTATCTC AGTAGACTCT TCACACCATG T

October	EnV 14	GQ126860.1	Human coxsackievirus B5 5' UTR	3.00E-67	100%	CCAAAGTA GTCGGTTCCG CTGCAGAGTT GCCCGTTACG ACACACTGCC CCCTGGCTTG AGGGTGTGTG CTCCGCAGTT AGGATTAGCC GCATTCAGGG GCCGGAGGAT TACCAACTAG CTCAATAGAC TCTTCACACC ATGTA
April	EnV 7	GQ126860.1	Human coxsackievirus B5 5' UTR	2.00E-25	100%	CCAAGTAGTC GGTTCCGCTG CAGAGTTGCC CGTTACGACA CACTGCCCCC TGGCTTGAGG GTGTGTGCAA
October	NoVI 11	AB112128.1	Human norovirus Saitama strain: NV/Saitama T83GI/02/JP	6.00E-31	98%	CC TTAGACGCCA TCATCATTTA CATAATCGGG TAGGAGATTG CGATCTCCTG CCCACAAGCT CAAGTCATGG AACCGCATCC
April	NoVII 6	JX989075.1	Norovirus Hu/GII.6/GZ2010-L96/Guangzhou/CHN/2011, complete genome	1.00E-39	98%	TCGACGCCAT CTTCATTAC ACTCTCGGGA GCAAGATTGC GATCGCCCTC CCATGTGCTG AGGTCAGAGA ATCTCATCCA CCTAAACATC GACTCTTGA