



## Eleven High-Quality Reference Genome Sequences and 360 Draft Assemblies of Shiga Toxin-Producing *Escherichia coli* Isolates from Human, Food, Animal, and Environmental Sources in Canada

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**ABSTRACT** We report high-quality closed reference genomes for 1 bovine strain and 10 human Shiga toxin (Stx)-producing *Escherichia coli* (STEC) strains from serogroups O26, O45, O91, O103, O104, O111, O113, O121, O145, and O157. We also report draft assemblies, with standardized metadata, for 360 STEC strains isolated from watersheds, animals, farms, food, and human infections.

Shiga toxin (Stx)-producing *Escherichia coli* (STEC) strains cause significant human enteric disease (1–3). Among >129 O serogroups, O157, O26, O45, O111, O103, O121, and O145 cause most infections (4–6). Non-O157 STEC strains are increasingly reported (5–7), with recent widespread STEC O121 and O103 outbreaks in Canada and the United States sourced to flour (8) and ground beef, respectively (<https://www.cdc.gov/ecoli/2019/o103-04-19/>).

To catalogue pangenomic diversity, we generated closed reference genomes for 11 routinely used lab control strains from the “top seven” STEC O serogroups, plus O91 and O104, and 360 draft assemblies for 129 distinct O serogroups from STEC culture collections (1980 to 2013) originating from watersheds, farms or foods ( $n = 238$ ), human infections ( $n = 74$ ), proficiency panels ( $n = 27$ ), and unknown sources ( $n = 32$ ). Prior to selection, isolates were traditionally serotyped at national or provincial reference labs, and *stx* gene presence/subtype was assessed by preestablished generic and differentiating *stx* PCR assays (9–11).

DNA extracted from 1-ml Luria-Bertani broth cultures grown overnight at 37°C using MasterPure complete DNA purification kits (Epicentre Technologies Corp., Chicago, IL, USA) was fragmented by an E210 ultrasonicator (Covaris, Inc., Woburn, MA, USA). TruSeq DNA library preparation v2 kit (Illumina, San Diego, CA, USA) libraries were shotgun sequenced using an Illumina GAIIx system ( $2 \times 150$ -bp paired-end cluster generation kit v4 and TruSeq SBS kit v5) or MiSeq platform ( $2 \times 300$  bp; v3 chemistry). Illumina reads were managed in the Integrated Rapid Infectious Disease Analysis (IRIDA) platform (12), assessed for quality ( $Q > 30$ ) using FastQC (13), and trimmed using Trimmomatic v0.34 (14). Overlapping reads merged with FLASH v1.2.11 (15) were *de novo* assembled using SPAdes v3.8.2 (16)/Shovill 0.9.0 (17). Postassembly quality control was achieved using QUAST v5.0.0 (number of contigs, < 500; reference coverage, 70%

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[closest polished genome of FWSEC0001-0011]) (18). Reference strains were augmented with  $2 \times 300$ -bp MiSeq (v3 chemistry) reads from mate pair ( $\sim 8$  kb) TruSeq libraries and with MinION Mk1b long reads from the rapid barcoding sequencing kit (SQK-RBK004; Oxford Nanopore Technologies Ltd., Oxford, UK) libraries. Albacore v2.3.0 base-called/quality-filtered long reads were *de novo* assembled using Canu v1.7 (19) and with quality-controlled Illumina mate pair reads as hybrid assemblies using Unicycler v0.4.4.0 (20). When assemblies appeared congruent in Mauve v20150226Build10 (21), Unicycler assemblies were used. Otherwise, mate pair reads were mapped to both assemblies using Bowtie 2 v2.3.4.1 (22) and BAM files assessed for coverage/connections using GAP5 v.1.2.14-r (23); long reads were mapped with BWA-MEM v0.7.17.1 (24) and assessed using Tablet v1.17.08.17 (25). Canu contigs were employed to scaffold/correct Unicycler contigs using the Staden package GAP4 (26, 27); all contigs were circularized and trimmed. Assemblies were Illumina read polished (5 rounds) using Bowtie 2/Pilon v1.20.1 (28). NCBI's default Web BLASTN (29) identified plasmid contigs and confirmed that *in silico* O-serogroup determinations were congruent with traditional lab determinations. Read depth was assessed using SAMtools idxstats (30). After functional annotation using NCBI's Prokaryotic Genome Annotation Pipeline (31), assemblies were reoriented to replication origin (*dnaA*) using Circlator v1.1.5 (32).

Illumina reference genome coverage ranged from  $65.6\times$  to  $130.7\times$  (average,  $96.1\times$ ); MinION coverage ranged from  $51.2\times$  to  $325.1\times$  (average,  $149.6\times$ ) (Table 1). Of 360 draft assemblies, 357 yielded scaffolds (average contigs, 153.0; average coverage depth,  $111.7\times$ ). Eleven reference chromosomes and all plasmids but one were circularized (0 to 3 plasmids per strain). The reference chromosomes (4,955,402 to 5,697,154 bp) contained 4,967 to 5,833 coding sequences (CDS), 22 rRNAs, 90 to 103 tRNAs, and 8 to 11 noncoding RNAs (ncRNAs), as well as bacteriophages. These genomic resources augment available data and are ideal for pathogenomics applications and machine learning.

**Data availability.** The standardized strain descriptions and accession numbers are presented in Table 1; the genomic data are publicly available in DDBJ/ENA/GenBank under BioProject no. [PRJNA287560](#) and in the Sequence Read Archive under accession no. [SRP155537](#). The versions described are the first versions.

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**TABLE 1** Characteristics and accession numbers for 111 high-quality STEC reference genomes and 360 STEC draft genome assemblies from 129 distinct serogroups sequenced for this study

Isolate identifier from this study	Traditional serogroup ( <i>in silico</i> prediction) <sup>a</sup>	STX subtype ( <i>in</i> <i>silico</i> prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no. <sup>c</sup>	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or plasmid(s), size (bp)	No. of contigs postassembly, (GenBank accession no.)	No. of CDS <sup>d</sup>	No. of Coverage (>X) (technology)	Genome size (all contigs) (bp)
FWSEC0001	O26:H11	STX <sub>1a</sub> , STX <sub>2a</sub>	Clinical, human	2002	Canada	GCF_005037725	Complete genome	4, chromosome, 5,697,154 (CP031922); plasmids, (CP031924), 4,148 (CP031925)	5,833	66,48 (Illumina), 320,55 (MinION)	5,803,315		
FWSEC0002	O145:NM (O145:H28)	STX <sub>1a</sub>	Clinical, human	2003	Canada	GCA_005037815	Complete genome	3, chromosome, 5,695,528 (CP031920), 2,031 (CP031921)	5,594	78.00 (Illumina), 124,40 (MinION)	5,790,057		
FWSEC0003	O45:H2	STX <sub>1a</sub>	Clinical, human	2005	Canada	GCA_005037845	Complete genome	3, chromosome, 5,532,455 (CP031916); plasmids, 95,228 (CP031917), 52,940 (CP031918)	5,612	100,74 (Illumina), 244,02 (MinION)	5,680,623		
FWSEC0004	O157:H7	STX <sub>1a</sub> , STX <sub>2a</sub>	Clinical, human	1987	Canada	GCA_005037735	Complete genome	3, chromosome, 5,406,250 (CP031913); plasmids, 92,754 (CP031914), 6,675 (CP031915)	5,420	130,71 (Illumina), 172,42 (MinION)	5,505,679		
FWSEC0005	O111:NM (O111:H8)	STX <sub>1a</sub> , STX <sub>2a</sub>	Clinical, human	2000	Canada	GCA_005037805	Complete genome	1, chromosome, 5,132,754 (CP031912); plasmids, none detected	4,967	108,41 (Illumina), 252,38 (MinION)	5,132,754		
FWSEC0006	O121:H19	STX <sub>2a</sub>	Clinical, human	2003	Canada	GCA_005037715	Complete genome	2, chromosome, 5,398,870 (CP031910); plasmid, 80,681 (CP031911)	5,344	103,62 (Illumina), 127,66 (MinION)	5,479,551		
FWSEC0007	O103:H2	STX <sub>1a</sub>	Clinical, human	2004	Canada	GCA_005037795	Complete genome	2, chromosome, 5,397,605 (CP031908); plasmid, 73,224 (CP031909)	5,447	103,12 (Illumina), 38,53 (MinION)	5,470,829		
FWSEC0008	O91:H21	STX <sub>2d</sub>	Clinical, human	1992	Germany	GCA_005037775	Complete genome	2, chromosome, 4,972,544 (CP031906); plasmid, 121,036 (CP031907)	4,971	98,41 (Illumina), 82,67 (MinION)	5,093,580		
FWSEC0009	O104:H4	STX <sub>2a</sub>	Clinical, human	2011	Germany	GCA_005014075	Complete genome	4, chromosome, 5,277,234 (CP031902); plasmids, 88,545 (CP031903), 75,669 (CP031904), 1,549 (CP031905)	5,329	65,55 (Illumina), 110,27 (MinION)	5,442,997		
FWSEC0010	O113:H21	STX <sub>2a</sub>	Clinical, human	1991	Canada	GCA_005014055	Complete genome	4, chromosome, 4,955,402 (CP031898); plasmids, 160,712 (CP031899), 7,769 (CP031900), 7,117 (CP031901)	5,006	94,45 (Illumina), 51,16 (MinION)	5,131,000		
FWSEC0011	O113:H21	STX <sub>2d</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2004	Canada	GCA_005171095	Chromosome 6, chromosome (CP031892); plasmids, 134,626 (CP031893), 115,288 in 4 scaffolds, <sup>e</sup> (66,896 (CP031894), 26,567 (CP031895), 12,383 (CP031896), 9,442 (CP031897))	5,173	108,45 (Illumina), 96,89 (MinION)	5,314,491			
FWSEC0021	O26:H11	STX <sub>1a</sub>	Clinical, human	1980	Canada	SRR7947278	RRCV000000000	Scaffold	208		92 (Illumina)	5,330,412	
FWSEC0022	O26:H11	STX <sub>1a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	1996	Canada	SRR7947279	RRCW000000000	Scaffold	240		89 (Illumina)	5,481,594	
FWSEC0023	O26:H11	STX <sub>1a</sub>	Clinical, human	1999	Switzerland	SRR7947276	RRCX000000000	Scaffold	223		130 (Illumina)	5,419,123	(Continued on next page)

**TABLE 1** (Continued)

	Isolate identifier from this study	Traditional serogroup ( <i>in silico</i> prediction) <sup>a</sup>	stx subtype ( <i>in</i> <i>silico</i> prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no. <sup>c</sup>	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or plasmid(s), size (bp)	No. of Coverage (>) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
FWSEC0024	O26:H11	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Ground beef, solid food, bovine, buildings (grocery/ retail/food store)	1999	Canada	SRR7947277	RRCY000000000	Scaffold	221	82 (Illumina)	5,315,394
FWSEC0025	O103:H2	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	1997	Canada	SRR7947274	RRCZ200000000	Scaffold	181	58 (Illumina)	5,214,405
FWSEC0026	O103:H2	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	1995	Canada	SRR7947275	RRDA000000000	Scaffold	198	56 (Illumina)	5,135,561
FWSEC0027	O103:H2	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	2001	Canada	SRR7947272	RRDB000000000	Scaffold	220	83 (Illumina)	5,413,960
FWSEC0028	O103:H2	STX <sub>1a</sub> , STX <sub>2a</sub>	STX <sub>1a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	2001	Switzerland	SRR7947273	RRDC000000000	Scaffold	159	82 (Illumina)	5,107,141
FWSEC0029	O45:H2	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	1994	Canada	SRR7947280	RRDD000000000	Scaffold	184	215 (Illumina)	5,378,562
FWSEC0030	O45:H2	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Clinical, human Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	1995	USA	SRR7947281	RRDE000000000	Scaffold	173	222 (Illumina)	5,134,981
FWSEC0031	O45:H2	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	1996	USA	SRR7947308	RRDF000000000	Scaffold	193	213 (Illumina)	5,496,950
FWSEC0032	O45:H2	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	1989	Canada	SRR7947309	RRDG000000000	Scaffold	165	199 (Illumina)	5,158,875
FWSEC0033	O111:H8	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	1993	Canada	SRR7947306	RRDH000000000	Scaffold	214	194 (Illumina)	5,281,723
FWSEC0034	O111:H8	STX <sub>1a</sub>	STX <sub>1a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2000	Canada	SRR7947307	RRDI000000000	Scaffold	209	177 (Illumina)	5,270,854
FWSEC0035	O111:NM	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	2000	Switzerland	SRR7947304	RRDJ000000000	Scaffold	179	212 (Illumina)	5,159,316
FWSEC0036	O111:NM	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	2002	Canada	SRR7947305	RRDK000000000	Scaffold	187	222 (Illumina)	5,281,033
FWSEC0039	O121:H19	STX <sub>2a</sub>	STX <sub>2a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	2002	Switzerland	SRR7947302	RRDL000000000	Scaffold	127	196 (Illumina)	5,130,319
FWSEC0040	O121:H19	STX <sub>2a</sub>	STX <sub>2a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	2004	Canada	SRR7947303	RRDM000000000	Scaffold	150	221 (Illumina)	5,182,763
FWSEC0041	O145:NM (O145:H28)	STX <sub>1a</sub>	STX <sub>1a</sub>	Clinical, human Clinical, human Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	1997	Switzerland	SRR7947300	RRDN000000000	Scaffold	192	196 (Illumina)	5,358,040
FWSEC0042	O145:H25	STX <sub>2a</sub>	STX <sub>2a</sub>	Clinical, human Clinical, human Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	1998	Switzerland	SRR7947301	RRDO000000000	Scaffold	174	189 (Illumina)	5,245,327
FWSEC0043	O145:NM (O145:H28)	STX <sub>2a</sub>	STX <sub>2a</sub>	Clinical, human Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	1999	Argentina	SRR7947248	RRDP000000000	Scaffold	169	190 (Illumina)	5,178,179
FWSEC0044	O145:NM (O145:H28)	STX <sub>1a</sub> , STX <sub>2a</sub>	STX <sub>1a</sub> , STX <sub>2a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	2001	Switzerland	SRR7947247	RRDQ000000000	Scaffold	162	177 (Illumina)	5,208,478
FWSEC0045	O91:NM (O91:H14)	STX <sub>1a</sub> , STX <sub>2b</sub>	STX <sub>1a</sub> , STX <sub>2b</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	1994	USA	SRR7947246	RRDR000000000	Scaffold	252	186 (Illumina)	5,459,036
FWSEC0046	O91:NM (O91:H14)	STX <sub>1a</sub> , STX <sub>2b</sub>	STX <sub>1a</sub> , STX <sub>2b</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	1994	USA	SRR7947245	RRDS000000000	Scaffold	251	191 (Illumina)	5,528,103
FWSEC0047	O91:H21	STX <sub>2d</sub>	STX <sub>1a</sub> , STX <sub>2a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	2001	USA	SRR7947244	RRDT000000000	Scaffold	109	210 (Illumina)	4,899,053
FWSEC0048	O91:H21	STX <sub>2d</sub>	STX <sub>1a</sub> , STX <sub>2a</sub>	Clinical, human Domesticated livestock, bovine, animal manure, agricultural (farm)	2001	Canada	SRR7947243	RRDU000000000	Scaffold	133	165 (Illumina)	5,100,887

(Continued on next page)

**TABLE 1** (Continued)

	Isolate identifier from this study	Traditional serogroup ( <i>in silico</i> prediction) <sup>a</sup>	stX subtype ( <i>in</i> <i>silico</i> prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no.	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or plasmid(s), size (bp)	No. of Coverage (>1) CDSe (technology)	Genome size (all contigs) (bp)
FWSEC0049	O113:H4	STX <sub>1αv</sub> ; STX <sub>2d</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	1996	Canada	SRR7947242	RRDV00000000	Scaffold	170	226 (Illumina)	4,951,257	4,987,637
FWSEC0050	O113:H4	STX <sub>1αv</sub> ; STX <sub>2d</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	1996	Canada	SRR7947241	R RDV00000000	Scaffold	196	139 (Illumina)	4,951,257	
FWSEC0051	O113:H21	STX <sub>2d</sub>	Clinical, human	1989	Canada	SRR7947250	R RDV00000000	Scaffold	122	196 (Illumina)	5,251,997	
FWSEC0052	O128:NM (O:128:H2)	STX <sub>1c</sub>	Clinical, human	1996	Germany	SRR7947249	R RDV00000000	Scaffold	196	196 (Illumina)	5,398,882	
FWSEC0053	O128:H2	STX <sub>1c</sub> STX <sub>1</sub> (partial); STX <sub>1αv</sub>	Clinical, human	1988	Switzerland	SRR7947252	R RDV00000000	Scaffold	213	240 (Illumina)	5,388,481	
FWSEC0054	O128:H10	STX <sub>1a</sub> (STX <sub>2</sub> ), STX <sub>1a</sub> (KAT)	Clinical, human	1999	Switzerland	SRR7947253	R REA00000000	Scaffold	141	187 (Illumina)	4,942,536	
FWSEC0055	O113:H21	STX <sub>2a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2000	Canada	SRR7947254	R REB00000000	Scaffold	120	199 (Illumina)	4,949,372	
FWSEC0057	O185:H28	STX <sub>1αv</sub> ; STX <sub>2a</sub>	River water	2012	Canada	SRR7947255	R REC00000000	Scaffold	419	46 (Illumina)	4,862,678	
FWSEC0058	O153:NM (O:153:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7947256	R REH00000000	Scaffold	500	5,232,124		
FWSEC0062	O182/O1:09:NM (O:109:H16)	STX <sub>2a</sub> ; STX <sub>2g</sub>	River water	2012	Canada	SRR7947260	R REH00000000	Scaffold	158	154 (Illumina)	5,214,988	
FWSEC0063	O182/O1:09:NM (O:109:H16)	STX <sub>2a</sub> ; STX <sub>2g</sub>	River water	2012	Canada	SRR7947261	R REH00000000	Scaffold	195	127 (Illumina)	5,205,797	
FWSEC0064	O8:H25	STX <sub>1αv</sub> ; STX <sub>2a</sub>	Stream water	2012	Canada	SRR7947273	R REJ00000000	Scaffold	160	77 (Illumina)	4,836,164	
FWSEC0066	O8:H19	STX <sub>2c</sub>	River water	2012	Canada	SRR7947275	R REJ00000000	Scaffold	135	72 (Illumina)	5,198,072	
FWSEC0067	O8:H19	STX <sub>2c</sub>	Municipal drain water	2012	Canada	SRR7947274	R REN00000000	Scaffold	144	99 (Illumina)	5,172,338	
FWSEC0069	O84:NM (O:84:H2)	STX <sub>1a</sub>	Solid stool, clinical, human	1999	Canada	SRR7947276	R REO00000000	Scaffold	194	185 (Illumina)	5,368,100	
FWSEC0070	O121:NM (O:121:H19)	STX <sub>2a</sub>	Solid stool, clinical, human	2004	Canada	SRR7947279	R REP00000000	Scaffold	133	129 (Illumina)	5,000,333	
FWSEC0071	O156:NM (O:156:H25)	STX <sub>1a</sub>	Solid stool, clinical, human	2004	Canada	SRR7947278	R REQ00000000	Scaffold	160	64 (Illumina)	5,054,573	
FWSEC0072	O26:H11	STX <sub>1a</sub>	Proficiency panel isolate, human	2005	Denmark	SRR7947381	R RRER00000000	Scaffold	205	97 (Illumina)	5,362,334	
FWSEC0073	O145:H28	STX <sub>1a</sub>	Proficiency panel isolate, human	2005	Denmark	SRR7947380	R RER00000000	Scaffold	142	103 (Illumina)	5,343,649	
FWSEC0074	O75:H8	STX <sub>1c</sub> ; STX <sub>2b</sub>	Clinical, human Solid stool, clinical, human	2005	Canada	SRR7947223	R RET00000000	Scaffold	153	79 (Illumina)	5,748,156	
FWSEC0075	O26:H11	STX <sub>1a</sub>	Solid stool, clinical, human	2005	Canada	SRR7947224	R REU00000000	Scaffold	239	81 (Illumina)	5,396,470	
FWSEC0076	O1:H20 (O:1:H21)	STX <sub>2a</sub>	Clinical, human Proficiency panel isolate, human	2006	Canada	SRR7947221	R REV00000000	Scaffold	81	92 (Illumina)	5,009,613	
FWSEC0077	O38:H26	STX <sub>1c</sub> ; STX <sub>2b</sub>	Proficiency panel isolate, human Solid stool, clinical, human	2006	Denmark	SRR7947222	R REV00000000	Scaffold	108	69 (Illumina)	5,138,266	
FWSEC0078	O51:H49	STX <sub>2e</sub>	Proficiency panel isolate, human Solid stool, clinical, human	2006	Denmark	SRR7947227	R REX00000000	Scaffold	201	102 (Illumina)	5,378,358	
FWSEC0079	O55:H7	STX <sub>2d</sub>	Solid stool, clinical, human	2006	Canada	SRR7947228	R REY00000000	Scaffold	125	90 (Illumina)	5,285,787	
FWSEC0080	O119:H25 (O:119:H25)	STX <sub>1a</sub> STX <sub>2c</sub> (partial); STX <sub>2d</sub> (SRS72); STX <sub>2c</sub>	Food	2006	Canada	SRR7947225	R REZ00000000	Scaffold	156	90 (Illumina)	5,146,192	
FWSEC0081	O NT:NM (O:177:H25)	STX <sub>2d</sub> (KAT)	Solid stool, clinical, human	2006	Canada	SRR7947226	R RFA00000000	Scaffold	185	136 (Illumina)	5,008,627	

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**TABLE 1** (Continued)

Isolate identifier from this study	Traditional serogroup (in silico prediction) <sup>a</sup>	stX subtype (in silico prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no.	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or plasmid(s), size (bp)		No. of Coverage (>) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
									(GenBank accession no.)	No. of contigs postassembly, (all contigs)		
FWSEC0082	O21:H8	stX <sub>2a</sub>	Proficiency panel isolate	2007	Denmark	SRR7947229	RRFB000000000	Scaffold	188	53 (Illumina)	4,960,003	
FWSEC0083	O28ab:NM (O28:H9)	stX <sub>2a</sub>	Proficiency panel isolate	2007	Denmark	SRR7947230	RRFC000000000	Scaffold	142	99 (Illumina)	4,996,535	
FWSEC0084	O26:H11	stX <sub>1a</sub>	Solid stool, clinical, human	2007	Canada	SRR7947334	RRFD000000000	Scaffold	229	106 (Illumina)	5,294,352	
FWSEC0086	O110:H28	stX <sub>2a</sub>	Unknown specimen type	2009	USA	SRR7947332	RRFF000000000	Scaffold	107	74 (Illumina)	4,944,331	
FWSEC0087	O104:H7 (O104:H19)	stX <sub>2b</sub>	Unknown specimen type	2009	USA	SRR7947331	RRFG000000000	Scaffold	237	128 (Illumina)	5,466,655	
FWSEC0088	O76:H19	stX <sub>1c</sub> stX <sub>2b</sub>	Proficiency panel isolate	2009	Denmark	SRR7947338	RRFH000000000	Scaffold	138	113 (Illumina)	5,327,960	
FWSEC0089	O113:H4	stX <sub>1c</sub> stX <sub>2b</sub>	Proficiency panel isolate	2009	Denmark	SRR7947337	RRFI000000000	Scaffold	134	71 (Illumina)	5,110,727	
FWSEC0091	O128ab:H2 (O128:H2)	stX <sub>1c</sub> stX <sub>2b</sub>	Proficiency panel isolate	2009	Denmark	SRR7947335	RRFJ000000000	Scaffold	195	89 (Illumina)	5,567,620	
FWSEC0092	O91:H14	stX <sub>2b</sub>	Proficiency panel isolate	2009	Denmark	SRR7947330	RRFK000000000	Scaffold	101	87 (Illumina)	5,105,837	
FWSEC0093	O146:H28	stX <sub>2b</sub>	Proficiency panel isolate	2009	Denmark	SRR7947329	RRFL000000000	Scaffold	159	127 (Illumina)	5,363,081	
FWSEC0094	O26:H11	stX <sub>1a</sub> , stX <sub>2a</sub>	Solid stool, clinical, human	2009	Canada	SRR7947355	RRFM000000000	Scaffold	209	113 (Illumina)	5,482,763	
FWSEC0095	O116:H NT (O116:H21)	stX <sub>1a</sub> , stX <sub>2a</sub>	Unknown specimen type	2009	USA	SRR7947356	RRFN000000000	Scaffold	137	107 (Illumina)	5,147,136	
FWSEC0096	O8:H19	stX <sub>1a</sub> , stX <sub>2d</sub>	Unknown specimen type	2009	USA	SRR7947357	RRFO000000000	Scaffold	81	93 (Illumina)	4,890,473	
FWSEC0097	O130:H11	stX <sub>2a</sub>	Unknown specimen type	2009	USA	SRR7947358	RRFP000000000	Scaffold	90	81 (Illumina)	5,054,940	
FWSEC0098	O6:H34	stX <sub>2a</sub>	Unknown specimen type	2009	USA	SRR7947351	RRFQ000000000	Scaffold	93	77 (Illumina)	5,170,195	
FWSEC0100	O26:H11	stX <sub>1a</sub>	Solid stool, clinical, human	2009	Canada	SRR7947353	RRFR000000000	Scaffold	201	156 (Illumina)	5,402,804	
FWSEC0101	O8:H16	stX <sub>1a</sub> , stX <sub>2a</sub>	Solid stool, clinical, human	2009	Canada	SRR7947354	RRFS000000000	Scaffold	70	133 (Illumina)	4,891,263	
FWSEC0102	O98:H29 (O98:H21)	stX <sub>1a</sub>	Clinical, human	2009	Canada	SRR7947359	RRFT000000000	Scaffold	160	147 (Illumina)	5,223,295	
FWSEC0103	O165:NM (O165:H25)	stX <sub>1a</sub> , stX <sub>2a</sub>	Solid stool, clinical, human	2010	Canada	SRR7947360	RRF000000000	Scaffold	166	141 (Illumina)	4,927,569	
FWSEC0104	O26:H11	stX <sub>1a</sub>	Unknown specimen type	2010	USA	SRR7947289	RRFV000000000	Scaffold	203	138 (Illumina)	5,283,037	
FWSEC0105	O156:H25	stX <sub>1a</sub>	Unknown specimen type	2010	USA	SRR7947288	RRFW000000000	Scaffold	146	82 (Illumina)	5,029,646	
FWSEC0106	O165:H25	stX <sub>1a</sub> , stX <sub>2a</sub>	Unknown specimen type	2010	USA	SRR7947291	RRFX000000000	Scaffold	154	93 (Illumina)	4,919,952	
FWSEC0107	O79:H7	stX <sub>2c</sub>	Unknown specimen type	2010	USA	SRR7947290	RRFY000000000	Scaffold	105	45 (Illumina)	4,882,294	
FWSEC0108	O39:H49	stX <sub>1a</sub> , stX <sub>2a</sub>	Unknown specimen type	2010	USA	SRR7947285	RRFZ000000000	Scaffold	101	88 (Illumina)	5,008,177	
FWSEC0109	O26:NM (O26:H11)	stX <sub>1a</sub>	Clinical, human	2010	Canada	SRR7947284	RRGA000000000	Scaffold	241	94 (Illumina)	5,339,476	
FWSEC0110	O103:H2	stX <sub>1a</sub>	Profoundly panel isolate	2010	Canada	SRR7947340	RRGB000000000	Scaffold	192	131 (Illumina)	5,341,967	
FWSEC0111	O174:H8	stX <sub>1c</sub> , stX <sub>2b</sub>	Profoundly panel isolate	2010	Denmark	SRR7947339	RRGC000000000	Scaffold	115	109 (Illumina)	5,029,489	

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**TABLE 1** (Continued)

Isolate identifier from this study	Traditional serogroup (in silico prediction) <sup>a</sup>	stx subtype (in silico prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no.	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or plasmid(s), size (bp)		No. of Coverage (>10%) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
									(GenBank accession no.)	117 (Illumina)		
FWSEC0112	O41:H26	STX <sub>1d</sub>	Proficiency panel isolate	2010	Denmark	SRR7947361	RGD000000000	Scaffold	120	72 (Illumina)	5,255,259	
FWSEC0113	O6:H2 (O103:H2)	STX <sub>1a</sub>	Unknown specimen type	2010	Canada	SRR7947328	RGD000000000	Scaffold	184	76 (Illumina)	5,146,693	
FWSEC0114	O139:H1	STX <sub>2e</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947318	RGD000000000	Scaffold	135	79 (Illumina)	5,161,156	
FWSEC0115	O171:H2	STX <sub>2c</sub> (partial), STX <sub>2</sub> (SRST2), STX <sub>2d</sub> (KATT)	Proficiency panel isolate	Missing	Denmark	SRR7947319	RGD000000000	Scaffold	151	86 (Illumina)	4,929,810	
FWSEC0116	O91:H21	STX <sub>2d</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947316	RGD000000000	Scaffold	97	65 (Illumina)	4,910,022	
FWSEC0117	O145:H34	STX <sub>2f</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947317	RGD000000000	Scaffold	105	83 (Illumina)	5,348,740	
FWSEC0118	O2:H25	STX <sub>2g</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947314	RGD000000000	Scaffold	135	97 (Illumina)	5,423,256	
FWSEC0119	O146:H21	STX <sub>1c</sub> , STX <sub>2a</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947315	RGD000000000	Scaffold	186	73 (Illumina)	5,305,349	
FWSEC0120	O154:H31	STX <sub>1d</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947282	RGD000000000	Scaffold	81	87 (Illumina)	5,397,345	
FWSEC0121	O22:H8	STX <sub>1c</sub> , STX <sub>2b</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947283	RGD000000000	Scaffold	193	99 (Illumina)	5,036,092	
FWSEC0122	O48:H21	STX <sub>1a</sub> , STX <sub>2a</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947310	RGD000000000	Scaffold	105	161 (Illumina)	5,239,138	
FWSEC0123	O174:H21	STX <sub>2c</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947311	RGD000000000	Scaffold	101	92 (Illumina)	5,011,013	
FWSEC0124	O118:H12	STX <sub>2b</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947236	RGD000000000	Scaffold	146	82 (Illumina)	5,126,562	
FWSEC0125	O73:H18 (O17:H18)	STX <sub>2d</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947235	RGD000000000	Scaffold	72	108 (Illumina)	5,592,160	
FWSEC0126	O8:k85ab:H rough (O8:H19)	STX <sub>1d</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947238	RGD000000000	Scaffold	77	80 (Illumina)	5,355,188	
FWSEC0127	O128ab:NM (O128:H2)	STX <sub>2f</sub>	Proficiency panel isolate	Missing	Denmark	SRR7947237	RGD000000000	Scaffold	164	72 (Illumina)	5,293,073	
FWSEC0128	O183:H21 (O183:H18)	STX <sub>1a</sub> , STX <sub>2a</sub>	Solid stool, clinical, human	2010	Canada	SRR7947232	RGD000000000	Scaffold	85	70 (Illumina)	5,212,364	
FWSEC0129	O71:H11	STX <sub>1a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947231	RGD000000000	Scaffold	215	112 (Illumina)	5,431,187	
FWSEC0131	O118:H16	STX <sub>1a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947233	RGD000000000	Scaffold	216	86 (Illumina)	4,960,664	
FWSEC0132	O177:NM (O177:H25)	STX <sub>2c</sub>	Solid stool, clinical, human	2011	Canada	SRR7947240	RGD000000000	Scaffold	204	102 (Illumina)	5,308,084	
FWSEC0133	O26:H11	STX <sub>1a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947239	RGD000000000	Scaffold	229	102 (Illumina)	4,775,607	
FWSEC0135	O49:NM (O49:H16)	STX <sub>2a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947389	RGD000000000	Scaffold	186	83 (Illumina)	5,325,986	
FWSEC0136	O103:H21 (O103:H2)	STX <sub>1a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947386	RGD000000000	Scaffold	172	78 (Illumina)	5,127,832	
FWSEC0137	O103:H2 (O103:H25)	STX <sub>1a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947387	RGD000000000	Scaffold	173	69 (Illumina)	5,264,166	
FWSEC0138	O103:H2	STX <sub>1a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947384	RGD000000000	Scaffold	157	65 (Illumina)	5,449,882	
FWSEC0139	O26:H21 (O26:H11)	STX <sub>1a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947385	RGD000000000	Scaffold	225			

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**TABLE 1** (Continued)

Isolate identifier from this study	Traditional serogroup (in silico prediction) <sup>a</sup>	stx subtype (in silico prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no.	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or plasmid(s), size (bp)	No. of Coverage (>) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
FWSEC0140	O121:H1 (O121:H19)	STX <sub>2a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947382	RRHF00000000	Scaffold	143	71 (Illumina)	5,127,687
FWSEC0141	O69:H11	STX <sub>1a</sub>	Solid stool, clinical, human	2010	Canada	SRR7947383	RRHG00000000	Scaffold	214	96 (Illumina)	5,495,125
FWSEC0142	O26:H21 (O26:H11)	STX <sub>1a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947390	RRHH00000000	Scaffold	229	81 (Illumina)	5,277,599
FWSEC0143	O103:H2	STX <sub>1a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947391	RRHI00000000	Scaffold	225	44 (Illumina)	5,201,971
FWSEC0144	O26:H11	STX <sub>1a</sub>	Unknown specimen type	2011	Canada	SRR7947367	RRHJ00000000	Scaffold	234	72 (Illumina)	5,303,448
FWSEC0146	O103:H25	STX <sub>1a</sub>	Clinical, human	2011	Canada	SRR7947365	RRHL00000000	Scaffold	181	149 (Illumina)	5,390,238
FWSEC0147	O103:H2	STX <sub>1a</sub>	Clinical, human	2011	Canada	SRR7947364	RRHM00000000	Scaffold	161	113 (Illumina)	5,304,488
FWSEC0150	O1:H rough (O1:H20)	STX <sub>2a</sub>	Clinical, human	2011	Canada	SRR7947369	RRHP00000000	Scaffold	83	134 (Illumina)	5,049,946
FWSEC0151	O111:NM (O111:H8)	STX <sub>1a</sub>	Solid stool, clinical, human	2001	Canada	SRR7947368	RRHQ00000000	Scaffold	189	144 (Illumina)	5,347,332
FWSEC0152	O111:NM (O111:H8)	STX <sub>1a</sub>	Solid stool, clinical, human	2010	Canada	SRR7947363	RRHR00000000	Scaffold	179	118 (Illumina)	5,193,004
FWSEC0153	O121:H19	STX <sub>2a</sub>	Solid stool, clinical, human	2010	Canada	SRR7947362	RRHS00000000	Scaffold	166	156 (Illumina)	5,078,658
FWSEC0154	O121:H19	STX <sub>2a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947347	RRHT00000000	Scaffold	169	145 (Illumina)	5,218,466
FWSEC0155	O121:H19	STX <sub>2a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947348	RRHU00000000	Scaffold	182	112 (Illumina)	5,077,668
FWSEC0156	O121:H19	STX <sub>2a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947349	RRHY00000000	Scaffold	159	178 (Illumina)	5,156,591
FWSEC0157	O111:NM (O111:H8)	STX <sub>1a</sub>	Solid stool, clinical, human	2010	Canada	SRR7947350	RRHW00000000	Scaffold	192	183 (Illumina)	5,287,703
FWSEC0158	O111:NM (O111:H8)	STX <sub>1a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947343	RRHX00000000	Scaffold	196	210 (Illumina)	5,346,730
FWSEC0159	O111:NM (O111:H8)	STX <sub>1a</sub> , STX <sub>2a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947344	RRHY00000000	Scaffold	244	183 (Illumina)	5,359,848
FWSEC0160	O121:H19	STX <sub>2a</sub>	Solid stool, clinical, human	2011	Canada	SRR7947345	RRHZ00000000	Scaffold	167	173 (Illumina)	5,118,519
FWSEC0161	O121:H19	STX <sub>2a</sub>	Clinical, human	2011	Canada	SRR7947346	RRIAO00000000	Scaffold	161	219 (Illumina)	5,114,368
FWSEC0233	O8:H19	STX <sub>2c</sub>	River water	2012	Canada	SRR7612229	RRIIC00000000	Scaffold	70	111 (Illumina)	4,959,778
FWSEC0234	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612228	RRIC00000000	Scaffold	139	118 (Illumina)	5,310,469
FWSEC0235	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612231	RRID00000000	Scaffold	135	116 (Illumina)	5,308,750
FWSEC0236	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612230	RRIE00000000	Scaffold	133	105 (Illumina)	5,313,318
FWSEC0237	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612225	RRIF00000000	Scaffold	134	87 (Illumina)	5,310,915
FWSEC0238	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612224	RRIG00000000	Scaffold	141	89 (Illumina)	5,316,450
FWSEC0239	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612227	RRIH00000000	Scaffold	133	102 (Illumina)	5,317,769
FWSEC0241	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612233	RRIJ00000000	Scaffold	133	96 (Illumina)	5,314,331
FWSEC0242	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612232	RRIK00000000	Scaffold	150	68 (Illumina)	5,311,201
FWSEC0244	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612362	RRIM00000000	Scaffold	243	121 (Illumina)	5,568,249
FWSEC0245	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612181	RRIN00000000	Scaffold	145	51 (Illumina)	5,309,767
FWSEC0246	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612182	RRIO00000000	Scaffold	135	114 (Illumina)	5,317,036
FWSEC0247	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612175	RRIP00000000	Scaffold	131	51 (Illumina)	5,309,640
FWSEC0248	O4 (O135)NM (O4:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612176	RRIQ00000000	Scaffold	131	125 (Illumina)	5,212,374
FWSEC0249	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612177	RRIR00000000	Scaffold	132	80 (Illumina)	5,309,552
FWSEC0251	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612223	RRIT00000000	Scaffold	132	72 (Illumina)	5,317,712
FWSEC0252	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612304	RRIU00000000	Scaffold	144	143 (Illumina)	5,310,446
FWSEC0253	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612506	RRIV00000000	Scaffold	139	79 (Illumina)	5,307,818
FWSEC0254	O45NM (O45:H2)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612505	RRIW00000000	Scaffold	135	49 (Illumina)	5,282,918

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TABLE 1 (Continued)

Isolate identifier from this study	Traditional serogroup ( <i>in silico</i> prediction) <sup>a</sup>	stx subtype (in <i>silico</i> prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no. SRR7612503	WGS accession no. <sup>c</sup> RRX00000000	Assembly level	No. of contigs postassembly, chromosome and/or plasmid(s), size (bp)	No. of Coverage (>) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
FWSEC0255	O4:NM (O45:H2)	STX <sub>2f</sub> STX <sub>2a</sub>	River water Ground beef, solid food, bovine, buildings (grocery/ retail/food store)	2012	Canada Canada	SRR7612504 SRR7612503	RRJZ00000000 RRY00000000	Scaffold Scaffold	135 106	88 (Illumina) 143 (Illumina)	5,314,657 4,939,455
FWSEC0256	O9:NM (O45:H2)	O9:NM (O45:H2)	River water Ground beef, solid food, bovine, buildings (grocery/ retail/food store)	2010	Canada Canada	SRR7612504 SRR7612503	RRJZ00000000 RRY00000000	Scaffold Scaffold	135 106	88 (Illumina) 143 (Illumina)	5,314,657 4,939,455
FWSEC0257	O9:1:H14	STX <sub>1a</sub>	River water	2011	Canada	SRR7612510	RRJZ00000000	Scaffold	96	120 (Illumina)	5,291,420
FWSEC0258	O11:1:H8	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	2011	Canada	SRR7612509	RRJA00000000	Scaffold	147	81 (Illumina)	5,104,080
FWSEC0259	O1:H20	STX <sub>2a</sub>	River water	2011	Canada	SRR7612508	RRJB00000000	Scaffold	67	122 (Illumina)	5,067,012
FWSEC0260	O11:1:H8	STX <sub>1a</sub> , STX <sub>2a</sub>	Ground beef, solid food, bovine, buildings (grocery/ retail/food store)	2011	Canada	SRR7612507	RRJC00000000	Scaffold	150	76 (Illumina)	5,110,497
FWSEC0261	O11:NM (O11:H8)	STX <sub>1a</sub>	River water	2011	Canada	SRR7612512	RRJD00000000	Scaffold	137	107 (Illumina)	5,137,935
FWSEC0262	O9:1:H14	STX <sub>1a</sub>	River water	2011	Canada	SRR7612511	RRJE00000000	Scaffold	101	107 (Illumina)	5,325,101
FWSEC0263	O1:H20	STX <sub>1a</sub>	River water	2011	Canada	SRR7612467	RRJF00000000	Scaffold	72	114 (Illumina)	5,026,605
FWSEC0264	O26:H11	STX <sub>1a</sub>	River water	2011	Canada	SRR7612468	RRJG00000000	Scaffold	186	136 (Illumina)	5,305,929
FWSEC0265	O103:H2	STX <sub>1a</sub>	River water	2011	Canada	SRR7612465	RRJH00000000	Scaffold	115	85 (Illumina)	5,327,519
FWSEC0266	O1:H20	STX <sub>1a</sub>	River water	2012	Canada	SRR7612466	RRJL00000000	Scaffold	70	82 (Illumina)	5,064,420
FWSEC0267	O103:H22	STX <sub>1a</sub>	River water	2012	Canada	SRR7612471	RRJU00000000	Scaffold	141	122 (Illumina)	5,327,547
FWSEC0268	O1:H20	STX <sub>1a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2012	Canada	SRR7612472	RRJK00000000	Scaffold	49	98 (Illumina)	4,974,621
FWSEC0269	O1:H20	STX <sub>1a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2012	Canada	SRR7612469	RRJL00000000	Scaffold	64	106 (Illumina)	5,021,634
FWSEC0270	O103:H12	STX <sub>1a</sub>	Water	2012	Canada	SRR7612470	RRJM00000000	Scaffold	135	67 (Illumina)	5,301,031
FWSEC0271	O103:H12	STX <sub>1a</sub>	Water	2012	Canada	SRR7612463	RRJN00000000	Scaffold	143	97 (Illumina)	5,362,142
FWSEC0272	O174:H21	STX <sub>2d</sub>	Water	2012	Canada	SRR7612464	RRJQ00000000	Scaffold	143	5,247,426	121 (Illumina)
FWSEC0273	O26:H11	STX <sub>1a</sub>	Water	2012	Canada	SRR7612473	RRJP00000000	Scaffold	180	72 (Illumina)	5,246,449
FWSEC0274	O145:F:M (O145:H28)	STX <sub>2a</sub>	Water	2012	Canada	SRR7612559	RRJC00000000	Scaffold	139	86 (Illumina)	5,169,990
FWSEC0275	O26:H11	STX <sub>1a</sub>	Water	2012	Canada	SRR7612336	RRJU00000000	Scaffold	182	83 (Illumina)	5,238,710
FWSEC0276	O26:NM (O26:H11)	STX <sub>1a</sub>	Water	2012	Canada	SRR7612335	RRJS00000000	Scaffold	179	85 (Illumina)	5,384,889
FWSEC0277	O26:H11	STX <sub>1a</sub>	Water	2012	Canada	SRR7612338	RRJT00000000	Scaffold	201	81 (Illumina)	5,371,254
FWSEC0278	O104:H7	STX <sub>2a</sub>	Water	2012	Canada	SRR7612337	RRJU00000000	Scaffold	58	57 (Illumina)	4,996,585
FWSEC0279	O103:H2	STX <sub>1a</sub>	Water	2012	Canada	SRR7612340	RRJV00000000	Scaffold	128	96 (Illumina)	5,432,377
FWSEC0280	O103:H2	STX <sub>1a</sub>	Water	2013	Canada	SRR7612339	RRJW00000000	Scaffold	120	113 (Illumina)	5,200,889
FWSEC0281	O118:H16	STX <sub>1a</sub>	Water	2013	Canada	SRR7612556	RRJX00000000	Scaffold	178	81 (Illumina)	5,483,599
FWSEC0282	O9:1:H21	STX <sub>1a</sub> , STX <sub>2d</sub> , STX <sub>2c</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2013	Canada	SRR7612555	RRJY00000000	Scaffold	83	92 (Illumina)	5,033,695
FWSEC0283	O1:H20	STX <sub>1a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2013	Canada	SRR7612244	RRJZ00000000	Scaffold	62	108 (Illumina)	5,089,114
FWSEC0284	O174:H21	STX <sub>2a</sub>	Ground beef, solid food, bovine, buildings (grocery/ retail/food store)	2013	Canada	SRR7612245	RRKA00000000	Scaffold	82	95 (Illumina)	5,077,035
FWSEC0285	O103:H11	STX <sub>1a</sub>	Water	2013	Canada	SRR7612089	RRKB00000000	Scaffold	193	81 (Illumina)	5,304,443

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**TABLE 1** (Continued)

Isolate identifier from this study	Traditional serogroup ( <i>in silico</i> prediction) <sup>a</sup>	stx subtype ( <i>in</i> <i>silico</i> prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no.	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or plasmid(s), size (bp)	No. of Coverage (>) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
FWSEC0286	O103:H25	stX <sub>1a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2013	Canada	SRR7612095	RRKC000000000	Scaffold	154	98 (Illumina)	5,289,604
FWSEC0287	O103:H25	stX <sub>1a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2013	Canada	SRR7612248	RRKD000000000	Scaffold	156	76 (Illumina)	5,276,158
FWSEC0289	O153:H25	stX <sub>1a</sub> , stX <sub>2a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2013	Canada	SRR7612250	RRKE000000000	Scaffold	90	67 (Illumina)	4,909,247
FWSEC0290	O113:H21	stX <sub>2a</sub> (partial), stX <sub>2a</sub> (SRST2), stX <sub>2a</sub> (KAT)	Domesticated livestock, bovine, animal manure, agricultural (farm)	2013	Canada	SRR7612251	RRKF000000000	Scaffold	112	97 (Illumina)	5,070,644
FWSEC0291	O113:H21	stX <sub>1a</sub> , stX <sub>2a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2013	Canada	SRR7612105	RRKG000000000	Scaffold	86	99 (Illumina)	5,165,323
FWSEC0292	O177:NM (O177:H25)	stX <sub>2c</sub>	River water	2012	Canada	SRR7612106	RRKH000000000	Scaffold	179	78 (Illumina)	4,883,321
FWSEC0293	O157:NM (O157:H7)	stX <sub>1a</sub> , stX <sub>2c</sub>	River water	2012	Canada	SRR7612096	RRKL000000000	Scaffold	119	72 (Illumina)	5,277,187
FWSEC0294	O26:H11	stX <sub>1a</sub>	River water	2012	Canada	SRR7612281	RRKL000000000	Scaffold	190	93 (Illumina)	5,310,274
FWSEC0296	O111:NM (O111:H8)	stX <sub>1a</sub> , stX <sub>2a</sub>	River water	2013	Canada	SRR7612341	RRKM000000000	Scaffold	139	124 (Illumina)	5,314,455
FWSEC0297	O111:H8	stX <sub>1a</sub> , stX <sub>2a</sub>	River water	2013	Canada	SRR7612092	RRKM000000000	Scaffold	161	54 (Illumina)	5,358,475
FWSEC0298	O111:NM (O111:H8)	stX <sub>1a</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612091	RRKN000000000	Scaffold	140	99 (Illumina)	5,289,100
FWSEC0299	O111:H8	stX <sub>1a</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612090	RRK000000000	Scaffold	140	128 (Illumina)	5,281,401
FWSEC0300	O26:H11	stX <sub>1a</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612557	RRKP000000000	Scaffold	180	112 (Illumina)	5,390,046
FWSEC0301	O26:H11	stX <sub>1a</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612283	RRKQ000000000	Scaffold	192	82 (Illumina)	5,385,968
FWSEC0302	O121:H19	stX <sub>2a</sub>	Domesticated livestock, bovine ( <i>Bos taurus</i> )	1992	Canada	SRR7612282	RRKR000000000	Scaffold	124	108 (Illumina)	5,029,461
FWSEC0303	O121:H19	stX <sub>2a</sub>	Domesticated livestock, bovine ( <i>Bos taurus</i> )	1992	Canada	SRR7612533	RRKS000000000	Scaffold	120	75 (Illumina)	5,030,995
FWSEC0304	O121:H19	stX <sub>2a</sub>	Clinical, human	1999	Switzerland	SRR7612534	RRKT000000000	Scaffold	108	108 (Illumina)	5,112,326
FWSEC0305	O121:H19	stX <sub>2a</sub>	Clinical, human	1999	Switzerland	SRR7612531	RRKL000000000	Scaffold	118	71 (Illumina)	5,167,571
FWSEC0319	O163:H19	stX <sub>2a</sub>	Drain water, waste water, culvert	2013	Canada	SRR7612530	RRKW000000000	Scaffold	82	102 (Illumina)	5,024,293
FWSEC0320	O163:H19	stX <sub>2a</sub>	Drain water, waste water, culvert	2013	Canada	SRR7612527	RRKX000000000	Scaffold	85	80 (Illumina)	5,121,870
FWSEC0321	O163:NM (O163:H19)	stX <sub>2a</sub>	Drain water, waste water, culvert	2013	Canada	SRR7612528	RRKY000000000	Scaffold	207	42 (Illumina)	4,918,527
FWSEC0322	O163:NM (O163:H19)	stX <sub>2a</sub>	Drain water, waste water, culvert	2013	Canada	SRR7612535	RRKZ000000000	Scaffold	84	135 (Illumina)	5,019,235
FWSEC0323	O163:H missing (O163:H19)	stX <sub>2a</sub>	Drain water, waste water, culvert	2013	Canada	SRR7612336	RRLA000000000	Scaffold	83	94 (Illumina)	5,021,924
FWSEC0324	O163:H19	stX <sub>2a</sub>	Drain water, waste water, culvert	2013	Canada	SRR7612172	RRLB000000000	Scaffold	85	106 (Illumina)	5,119,734
FWSEC0325	O163:H19	stX <sub>2a</sub>	Drain water, waste water, culvert	2013	Canada	SRR7612171	RRLC000000000	Scaffold	91	109 (Illumina)	5,211,926

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**TABLE 1 (Continued)**

Isolate identifier from this study	Traditional serogroup ( <i>in silico</i> prediction) <sup>a</sup>	stx subtype ( <i>in</i> <i>silico</i> prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no.	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or chromosome(s), size (bp)	No. of Coverage (>1) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
FWSEC0327	O163:H19	STX <sub>2a</sub>	Drain water, waste water, culvert	2013	Canada	SRR7612173	RRL000000000	Scaffold	85	108 (Illumina)	5,107,873
FWSEC0329	O163:H19	STX <sub>2a</sub>	Drain water, waste water, culvert	2013	Canada	SRR7612167	RRLG000000000	Scaffold	84	110 (Illumina)	5,019,985
FWSEC0330	O163:H19	STX <sub>2a</sub>	Drain water, waste water, culvert	2013	Canada	SRR7612170	RRLH000000000	Scaffold	86	96 (Illumina)	5,219,978
FWSEC0332	O54 (O57):H21	STX <sub>1d</sub>	River water	2013	Canada	SRR7612180	RRLJ000000000	Scaffold	75	93 (Illumina)	4,981,787
FWSEC0333	O54 (O57):H21	STX <sub>1d</sub>	River water	2013	Canada	SRR7612537	RRLK000000000	Scaffold	75	134 (Illumina)	4,920,129
FWSEC0334	O54 (O57):H21	STX <sub>1d</sub>	River water	2013	Canada	SRR7612276	RRLK000000000	Scaffold	71	121 (Illumina)	4,982,034
FWSEC0335	O8:H19	STX <sub>1a</sub> , STX <sub>2d</sub>	Stream water	2013	Canada	SRR7612277	RRLL000000000	Scaffold	95	90 (Illumina)	5,341,011
FWSEC0336	O118:H2	STX <sub>1a</sub>	River water	2013	Canada	SRR7612278	RRLM000000000	Scaffold	149	105 (Illumina)	5,256,612
FWSEC0337	O8:H19	STX <sub>2a</sub> , STX <sub>2d</sub>	River water	2012	Canada	SRR7612279	RRLN000000000	Scaffold	90	111 (Illumina)	5,212,166
FWSEC0338	O168:H18	STX <sub>2a</sub>	River water	2012	Canada	RRL000000000	Scaffold	111	152 (Illumina)	5,276,982	
FWSEC0339	O8:H19	STX <sub>2a</sub> , STX <sub>2d</sub>	River water	Missing	Canada	SRR7612273	RRLP000000000	Scaffold	88	119 (Illumina)	4,912,080
FWSEC0340	O116:H25	STX <sub>2d</sub>	Stream water	2012	Canada	SRR7612274	RRLQ000000000	Scaffold	85	102 (Illumina)	4,957,319
FWSEC0341	O5:NM	STX <sub>1a</sub>	Stream water	2012	Canada	SRR7612275	RRLR000000000	Scaffold	143	131 (Illumina)	4,991,314
FWSEC0342	O69:H11	STX <sub>1a</sub>	River water	2013	Canada	SRR7612270	RRLS000000000	Scaffold	209	76 (Illumina)	5,350,561
FWSEC0344	O5:NM (O5:H9)	STX <sub>1a</sub>	Stream water	2013	Canada	SRR7612401	RRLU000000000	Scaffold	144	99 (Illumina)	4,994,048
FWSEC0346	O22:H8	STX <sub>2d</sub>	River water	2013	Canada	SRR7612399	RRLX000000000	Contig	99	102 (Illumina)	4,982,956
FWSEC0347	O174:H21	STX <sub>2c</sub>	River water	2013	Canada	SRR7612398	RRLY000000000	Scaffold	102	113 (Illumina)	5,055,323
FWSEC0348	O163:NM (O163:H19)	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	Missing	Canada	SRR7612405	RRLY000000000	Scaffold	80	126 (Illumina)	5,022,312
FWSEC0350	O163:NM (O163:H19)	STX <sub>1a</sub> , STX <sub>2a</sub>	Stream water	Missing	Canada	SRR7612403	RRMA000000000	Scaffold	78	165 (Illumina)	5,022,555
FWSEC0351	O163:NM (O163:H19)	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	Missing	Canada	SRR7612402	RRMB000000000	Scaffold	80	117 (Illumina)	5,021,604
FWSEC0352	O rough:NM (O163:H19)	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	Missing	Canada	SRR7612397	RRMC000000000	Scaffold	78	105 (Illumina)	5,020,306
FWSEC0353	O163:NM (O163:H19)	STX <sub>1a</sub> , STX <sub>2a</sub>	Canal water	Missing	Canada	SRR7612396	RRMD000000000	Scaffold	82	124 (Illumina)	5,021,023
FWSEC0354	O rough:NM (O163:H19)	STX <sub>1a</sub> , STX <sub>2a</sub>	Canal water	Missing	Canada	SRR7612083	RRME000000000	Scaffold	80	121 (Illumina)	5,019,161
FWSEC0355	O163:H19	STX <sub>1a</sub> , STX <sub>2a</sub>	Canal water	Missing	Canada	SRR7612084	RRMF000000000	Scaffold	76	132 (Illumina)	5,022,505
FWSEC0356	O69:H11	STX <sub>1a</sub>	River water	2013	Canada	SRR7612081	RRMG000000000	Scaffold	213	99 (Illumina)	5,417,563
FWSEC0357	O rough:NM (O163:H19)	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	Missing	Canada	SRR7612082	RRMH000000000	Scaffold	81	153 (Illumina)	5,022,476
FWSEC0358	O rough:NM (O163:H19)	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	Missing	Canada	SRR7612087	RRMI000000000	Scaffold	77	126 (Illumina)	5,021,430
FWSEC0359	O rough:NM (O163:H19)	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	Missing	Canada	SRR7612088	RRMJ000000000	Scaffold	87	156 (Illumina)	5,019,056
FWSEC0360	O rough:H21 (O113:H21)	STX <sub>2d</sub>	River water	2013	Canada	SRR7612085	RRMK000000000	Scaffold	102	105 (Illumina)	5,129,184
FWSEC0361	O113:H21	STX <sub>2d</sub>	River water	2013	Canada	SRR7612086	RRML000000000	Scaffold	136	31 (Illumina)	5,244,379
FWSEC0362	O5:NM (O5:H9)	STX <sub>1a</sub>	River water	2013	Canada	SRR7612079	RRMM000000000	Scaffold	146	149 (Illumina)	4,983,571
FWSEC0363	O8:H19	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	2013	Canada	SRR7612080	RRMN000000000	Scaffold	95	108 (Illumina)	4,909,165
FWSEC0364	O174:H21	STX <sub>2a</sub>	River water	2013	Canada	SRR7612237	RRMO000000000	Scaffold	124	88 (Illumina)	5,039,528
FWSEC0365	O128:H2	STX <sub>1c</sub> , STX <sub>2b</sub>	Stream water	2013	Canada	SRR7612238	RRMP000000000	Scaffold	132	87 (Illumina)	5,412,442
FWSEC0367	O NT:H19 (O8:H19)	STX <sub>1a</sub> , STX <sub>2a</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612240	RRMT000000000	Scaffold	83	92 (Illumina)	4,857,754
FWSEC0369	O98:NM (O98:H21)	STX <sub>1a</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612240	RRMT000000000	Scaffold	155	85 (Illumina)	5,326,975
FWSEC0370	O98:NM (O98:H21)	STX <sub>1a</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612243	RRMU000000000	Scaffold	162	96 (Illumina)	5,322,293
FWSEC0371	O157:H7	STX <sub>1a</sub> , STX <sub>2a</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612242	RRMV000000000	Scaffold	108	127 (Illumina)	5,209,164
FWSEC0372	O163:H19	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	Missing	Canada	SRR7612235	RRMW000000000	Scaffold	78	67 (Illumina)	5,022,836

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TABLE 1 (Continued)

Isolate identifier from this study	Traditional serogroup ( <i>in silico</i> prediction) <sup>a</sup>	stx subtype (in <i>silico</i> prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no. <sup>c</sup>	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or mid(s), size (bp)	No. of Coverage (>) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
FWSEC0373	O165:H25	STX <sub>1α</sub> , STX <sub>2α</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612234	RMMX00000000	Scaffold	163	49 (Illumina)	4,945,961
FWSEC0374	O165:NM (O165:H25)	STX <sub>1α</sub> , STX <sub>2α</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612235	RMMY00000000	Scaffold	159	36 (Illumina)	4,948,312
FWSEC0375	O157:H7	STX <sub>1α</sub> , STX <sub>2α</sub>	Stream water	2013	Canada	SRR7612236	RMMZ00000000	Scaffold	128	238 (Illumina)	5,305,861
FWSEC0376	O88:H25	STX <sub>1α</sub> , STX <sub>2α</sub>	River water	2013	Canada	SRR7612337	RRNA00000000	Scaffold	84	156 (Illumina)	4,833,021
FWSEC0377	O103:H2	STX <sub>1α</sub>	River water	2013	Canada	SRR7612338	RRNB00000000	Scaffold	140	145 (Illumina)	5,304,691
FWSEC0378	O103:H25	STX <sub>1α</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612339	RRNC00000000	Scaffold	165	96 (Illumina)	5,304,528
FWSEC0379	O26:H11	STX <sub>1α</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612330	RND00000000	Scaffold	178	108 (Illumina)	5,366,964
FWSEC0380	O165:NM (O165:H25)	STX <sub>2α</sub>	River water	2013	Canada	SRR7612331	RNE00000000	Scaffold	162	146 (Illumina)	4,966,985
FWSEC0381	O174:H8	STX <sub>1α</sub> , STX <sub>2b</sub>	Stream water	2013	Canada	SRR7612332	RNF00000000	Scaffold	141	129 (Illumina)	5,277,367
FWSEC0382	O128:H2	STX <sub>1α</sub> , STX <sub>2b</sub>	Stream water	Missing	Canada	SRR7612333	RNF00000000	Scaffold	143	86 (Illumina)	5,383,076
FWSEC0383	O163:H19	STX <sub>1α</sub> , STX <sub>2α</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612334	RNH00000000	Scaffold	83	103 (Illumina)	5,099,503
FWSEC0384	O128:H2	STX <sub>1α</sub> , STX <sub>2b</sub>	River water	2013	Canada	SRR7612462	RNI00000000	Scaffold	164	81 (Illumina)	5,533,994
FWSEC0385	O111:NM (O111:H8)	STX <sub>1α</sub> , STX <sub>2α</sub>	River water	2013	Canada	SRR7612461	RNI00000000	Scaffold	183	88 (Illumina)	5,370,982
FWSEC0386	O8:H9	STX <sub>2d</sub>	Water, agricultural (irrigation ditch)	2013	Canada	SRR7612460	RNK00000000	Scaffold	53	91 (Illumina)	5,789,001
FWSEC0387	O103:H25	STX <sub>1α</sub> , STX <sub>2α</sub>	Canal water	2013	Canada	SRR7612459	RNL00000000	Scaffold	164	178 (Illumina)	5,178,285
FWSEC0388	O103:H25	STX <sub>1α</sub>	River water	Missing	Canada	SRR7612458	RNN00000000	Scaffold	155	112 (Illumina)	5,168,250
FWSEC0389	O103:H2	STX <sub>1α</sub>	River water	2013	Canada	SRR7612457	RNN00000000	Scaffold	171	104 (Illumina)	5,301,480
FWSEC0390	O103:H2	STX <sub>1α</sub>	River water	2013	Canada	SRR7612456	RNN00000000	Scaffold	157	73 (Illumina)	5,236,779
FWSEC0391	O103:H2	STX <sub>1α</sub> , STX <sub>2d</sub>	River water	2013	Canada	SRR7612455	RNP00000000	Scaffold	182	93 (Illumina)	5,424,966
FWSEC0392	O163:NM (O163:H19)	STX <sub>1α</sub> , STX <sub>2α</sub>	Domesticated livestock, animal manure, bovine (dairy liquid), dairy farm	1992	Canada	SRR7612454	RNQ00000000	Scaffold	77	109 (Illumina)	5,022,374
FWSEC0394	O157:H7	STX <sub>1α</sub> , STX <sub>2α</sub>	Waste water	2010	Canada	SRR7612453	RNR00000000	Scaffold	113	82 (Illumina)	5,252,251
FWSEC0395	O111:H8	STX <sub>1α</sub>	Waste water	2010	Canada	SRR7612155	RNS00000000	Scaffold	104	97 (Illumina)	5,003,932
FWSEC0397	O174:H1	STX <sub>2d</sub>	Waste water	2011	Canada	SRR7612154	RNV00000000	Scaffold	76	74 (Illumina)	4,91,063
FWSEC0398	O52:H5	STX <sub>1c</sub>	Waste water	2011	Canada	SRR7612153	RNV00000000	Scaffold	100	102 (Illumina)	4,784,323
FWSEC0399	O91:H14	STX <sub>1α</sub>	Intake water	2011	Canada	SRR7612151	RNW00000000	Scaffold	93	102 (Illumina)	5,324,693
FWSEC0400	O113:H4	STX <sub>2d</sub>	Intake water	2011	Canada	SRR7612152	RNX00000000	Contig	128	107 (Illumina)	4,863,421
FWSEC0401	O113:H21	STX <sub>2d</sub>	Waste water	2011	Canada	SRR7612149	RNY00000000	Scaffold	96	118 (Illumina)	5,086,552
FWSEC0402	O171:H2	STX <sub>2c</sub>	Intake water	2011	Canada	SRR7612150	RNZ00000000	Scaffold	111	93 (Illumina)	5,233,546
FWSEC0404	O121:H10	STX <sub>2e</sub>	River water	2011	Canada	SRR7612148	RROA00000000	Scaffold	123	71 (Illumina)	5,095,240
FWSEC0405	O157:H7	STX <sub>1α</sub> , STX <sub>2α</sub>	River water	2011	Canada	SRR7612312	RROB00000000	Scaffold	127	49 (Illumina)	5,340,005
FWSEC0406	O157:H7	STX <sub>2c</sub>	Intake water	2011	Canada	SRR7612311	RROC00000000	Scaffold	130	110 (Illumina)	5,239,223
FWSEC0407	O128:NM (O128:H2)	STX <sub>1c</sub> , STX <sub>2b</sub>	Waste water	2011	Canada	SRR7612314	RROD00000000	Scaffold	132	105 (Illumina)	5,345,000
FWSEC0408	O91:NM (O91:H14)	STX <sub>1α</sub>	River water	2012	Canada	SRR7612313	RROE00000000	Scaffold	127	106 (Illumina)	5,346,527
FWSEC0409	O103:H25	STX <sub>1α</sub>	River water	2012	Canada	SRR7612308	RROF00000000	Scaffold	159	106 (Illumina)	5,262,960
FWSEC0410	O157:H7	STX <sub>1α</sub> , STX <sub>2α</sub>	River water	2012	Canada	SRR7612307	RROG00000000	Scaffold	145	117 (Illumina)	5,343,022
FWSEC0411	O121:H19	STX <sub>2a</sub>	Stream water	2012	Canada	SRR7612310	RROH00000000	Scaffold	118	115 (Illumina)	5,000,817
FWSEC0413	O157:H7	STX <sub>2a</sub>	Waste water	2012	Canada	SRR7612306	RROI00000000	Scaffold	117	129 (Illumina)	5,237,889
FWSEC0414	O153:NM (O153:H2)	STX <sub>2f</sub>	Intake water	2012	Canada	SRR7612305	RROJ00000000	Scaffold	167	130 (Illumina)	5,262,169
FWSEC0415	O157:H7	STX <sub>2a</sub>	Waste water	2012	Canada	SRR7612351	RROK00000000	Scaffold	125	117 (Illumina)	5,347,671
FWSEC0416	O113:H11	STX <sub>2a</sub>	Waste water	2012	Canada	SRR7612350	RROL00000000	Scaffold	98	115 (Illumina)	5,003,403
FWSEC0417	O157:H7	STX <sub>1α</sub>	River water	2012	Canada	SRR7612353	RROM00000000	Scaffold	122	118 (Illumina)	5,277,352
FWSEC0418	O139:H19	STX <sub>1α</sub> , STX <sub>2c</sub>	River water	2012	Canada	SRR7612352	RRON00000000	Scaffold	93	110 (Illumina)	5,051,814
FWSEC0419	O26:H11	STX <sub>1α</sub>	Intake water	2012	Canada	SRR7612347	RROO00000000	Scaffold	189	115 (Illumina)	5,475,738

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**TABLE 1** (Continued)

Isolate identifier from this study	Traditional serogroup ( <i>in silico</i> prediction) <sup>a</sup>	stx subtype ( <i>in</i> <i>silico</i> prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no.	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or chromosome(s), size (bp)		No. of Coverage (>) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
									No. of contigs postassembly, chromosome and/or chromosome(s), size (bp)	No. of contigs postassembly, chromosome and/or chromosome(s), size (bp)		
FWSEC0420	O157:H7	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	2012	Canada	SRR7612546	RROP000000000	Scaffold	111	118 (Illumina)	5,419,843	
FWSEC0421	O22:H NT (O22:H8)	STX <sub>2f</sub>	River water	2012	Canada	SRR7612556	RROQ000000000	Scaffold	109	112 (Illumina)	5,104,544	
FWSEC0422	O103:H2	STX <sub>1a</sub>	River water	2013	Canada	SRR7612548	RROR000000000	Scaffold	128	114 (Illumina)	5,192,306	
FWSEC0423	O91:NM (O91:H14)	STX <sub>1a</sub>	River water	2013	Canada	SRR7612543	RRC000000000	Scaffold	139	112 (Illumina)	5,391,718	
FWSEC0424	O182 (O109:H5) (O109:H5)	STX <sub>1a</sub>	River water	2013	Canada	SRR7612542	RCT000000000	Scaffold	86	158 (Illumina)	5,195,233	
FWSEC0425	O5:NM (O5:H9)	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	2013	Canada	SRR7612195	RROU000000000	Scaffold	159	123 (Illumina)	5,100,963	
FWSEC0426	O128:H2	STX <sub>1c</sub> , STX <sub>2b</sub>	River water	2013	Canada	SRR7612421	RROV000000000	Scaffold	175	121 (Illumina)	5,573,765	
FWSEC0427	O103:H2	STX <sub>1a</sub>	River water	2013	Canada	SRR7612418	RROV000000000	Scaffold	145	112 (Illumina)	5,298,328	
FWSEC0428	O26:H11	STX <sub>1a</sub>	River water	2013	Canada	SRR7612419	RROV000000000	Scaffold	182	137 (Illumina)	5,443,552	
FWSEC0429	O1:H20	STX <sub>1a</sub>	River water	2013	Canada	SRR7612416	RROY000000000	Scaffold	63	144 (Illumina)	5,071,067	
FWSEC0430	O157:H7	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	2013	Canada	SRR7612417	RZOZ000000000	Scaffold	128	88 (Illumina)	5,348,101	
FWSEC0431	O103:H11	STX <sub>1a</sub>	River water	2013	Canada	SRR7612414	RPA000000000	Scaffold	183	99 (Illumina)	5,303,799	
FWSEC0432	O5:NM (O5:H9)	STX <sub>1a</sub>	River water	2013	Canada	SRR7612415	RRPB000000000	Scaffold	159	91 (Illumina)	5,118,324	
FWSEC0433	O45:NM (O45:H2)	STX <sub>2f</sub>	River water	2013	Canada	SRR7612183	RPPC000000000	Scaffold	147	94 (Illumina)	5,227,140	
FWSEC0434	O45:NM (O45:H2)	STX <sub>2f</sub>	River water	2013	Canada	SRR7612184	RPPD000000000	Scaffold	152	83 (Illumina)	5,227,905	
FWSEC0436	O153:NM (O153:H2)	STX <sub>2f</sub>	River water	2013	Canada	SRR7612296	RPPE000000000	Scaffold	158	105 (Illumina)	5,323,485	
FWSEC0437	O45:NM (O45:H2)	STX <sub>2f</sub>	River water	2013	Canada	SRR7612295	RPPE000000000	Scaffold	156	110 (Illumina)	5,184,594	
FWSEC0438	O113:H4	STX <sub>2d</sub>	River water	2013	Canada	SRR7612294	RPGE000000000	Contig	122	99 (Illumina)	4,852,452	
FWSEC0439	O5:NM (O5:H9)	STX <sub>2f</sub>	River water	2013	Canada	SRR7612183	RPPD000000000	Scaffold	164	5,106,616	5,227,905	
FWSEC0440	O157:H7	STX <sub>1a</sub> , STX <sub>2a</sub>	River water	2013	Canada	SRR7612301	RPI000000000	Scaffold	125	109 (Illumina)	5,285,150	
FWSEC0441	O5:NM (O5:H19)	STX <sub>1c</sub>	River water	2013	Canada	SRR7612299	RPIJ000000000	Scaffold	120	123 (Illumina)	5,279,452	
FWSEC0442	O136:NM (O136:H12)	STX <sub>1a</sub>	River water	2013	Canada	SRR7612298	RPKD000000000	Scaffold	211	108 (Illumina)	5,301,182	
FWSEC0443	O5:NM (O5:H9)	STX <sub>1a</sub>	River water	2013	Canada	SRR7612303	RPL000000000	Scaffold	165	99 (Illumina)	5,108,519	
FWSEC0446	O6:H7 (O6:H11)	STX <sub>2a</sub>	Unknown specimen type	2001	Canada	SRR7612142	RPM000000000	Scaffold	81	76 (Illumina)	4,843,537	
FWSEC0447	O26:H11	STX <sub>1a</sub>	Unknown specimen type	Not collected	Not collected	SRR7612143	RPNP000000000	Scaffold	197	74 (Illumina)	5,379,480	
FWSEC0449	O145:NM (O145:H49)	STX <sub>1a</sub>	Unknown specimen type	Not collected	Not collected	SRR7612137	RRPP000000000	Scaffold	137	95 (Illumina)	4,833,826	
FWSEC0451	O103:H2	STX <sub>1a</sub>	Unknown specimen type	Not collected	Not collected	SRR7612139	RRPQ000000000	Scaffold	129	120 (Illumina)	5,163,904	
FWSEC0464	O121:NM (O121:H20)	STX <sub>1a</sub>	Unknown specimen type	Not collected	Canada	SRR7612121	RRPZ000000000	Scaffold	62	71 (Illumina)	5,029,443	
FWSEC0465	O113:H4 (O113:H20)	STX <sub>1a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	Not collected	Canada	SRR7612124	RRQA000000000	Scaffold	61	77 (Illumina)	5,068,214	
FWSEC0502	O91:H10	STX <sub>2</sub>	Unknown specimen type	Not collected	Not collected	SRR7612438	RRRH000000000	Scaffold	113	62 (Illumina)	5,055,432	
FWSEC0508	O26:H11	STX <sub>1a</sub>	Unknown specimen type	1984	Canada	SRR7947323	RRRI000000000	Scaffold	202	127 (Illumina)	5,273,275	
FWSEC0510	O103:H2	STX <sub>1a</sub>	Unknown specimen type	1997	Not collected	SRR7947325	RRRK000000000	Scaffold	166	303 (Illumina)	5,217,653	
FWSEC0511	O103:H2	STX <sub>1a</sub>	Unknown specimen type	Not collected	Not collected	SRR7947324	RRRL000000000	Scaffold	213	59 (Illumina)	5,271,027	
FWSEC0512	O103:H2	STX <sub>1a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	1999	Canada	SRR7947327	RRRM000000000	Scaffold	170	303 (Illumina)	5,353,158	
FWSEC0514	O103:H2	STX <sub>1a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	2004	Canada	SRR7612433	RRR000000000	Scaffold	141	76 (Illumina)	5,323,892	

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**TABLE 1** (Continued)

Isolate identifier from this study	Traditional serogroup (in silico prediction) <sup>a</sup>	stx subtype (in silico prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no.	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or plasmid(s), size (bp) (GenBank accession no.)	No. of Coverage (>) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
FWSEC0515	O139:H19	STX <sub>1a</sub> , STX <sub>2a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612432	RRP000000000	Scaffold	74	71 (Illumina)	5,020,809
FWSEC0516	O26:NM (O26:H11)	STX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612435	RRQ000000000	Scaffold	168	87 (Illumina)	5,327,060
FWSEC0517	O138:H38	STX <sub>1a</sub> , STX <sub>2a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612434	RRR000000000	Scaffold	117	109 (Illumina)	5,052,530
FWSEC0519	O139:H19	STX <sub>1a</sub> , STX <sub>2a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7947313	RRS000000000	Scaffold	597	31 (Illumina)	4,805,470
FWSEC0521	O26:NM (O26:H11)	STX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612321	RRR000000000	Scaffold	178	84 (Illumina)	5,321,164
FWSEC0523	O6:H34	STX <sub>1a</sub> , STX <sub>2a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612319	RRRU000000000	Scaffold	92	81 (Illumina)	4,880,556
FWSEC0526	O88:H25	STX <sub>1a</sub> , STX <sub>2a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7947312	RRV000000000	Scaffold	599	30 (Illumina)	4,689,586
FWSEC0527	O109:H5	STX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612317	RRRW000000000	Scaffold	74	44 (Illumina)	5,213,197
FWSEC0529	O88:H25	STX <sub>2a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612315	RRRX000000000	Scaffold	95	81 (Illumina)	4,927,775
FWSEC0530	O rough:H38 (O134:H38)	STX <sub>2c</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612316	RRRY000000000	Scaffold	69	94 (Illumina)	4,954,025
FWSEC0532	O168:H48	STX <sub>2a</sub>	Unknown specimen type	2012	Canada	SRR7947294	RRZ000000000	Scaffold	383	49 (Illumina)	5,191,428
FWSEC0534	O26:H11	STX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	Not collected	Canada	SRR7947295	RSA000000000	Scaffold	244	79 (Illumina)	5,309,969
FWSEC0535	O111:H missing (O111:H11)	STX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	Not collected	Canada	SRR7947292	RSB000000000	Scaffold	317	172 (Illumina)	5,583,793
FWSEC0536	O111:H11	STX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	Not collected	Canada	SRR7947293	RRSC000000000	Scaffold	306	245 (Illumina)	5,754,316

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**TABLE 1** (Continued)

	Isolate identifier from this study	Traditional serogroup ( <i>in silico</i> prediction) <sup>a</sup>	stX subtype ( <i>in</i> <i>silico</i> prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no.	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or plasmid(s), size (bp)	No. of Coverage (>1) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
FWSEC0337	O26:H missing (O26:H11)	stX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	Not collected	Canada	SRR7947298	RRS000000000	Scaffold	203	231 (Illumina)	5,274,187	
FWSEC0338	O26:H missing (O26:H11)	stX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	Not collected	Canada	SRR7947299	RRSE000000000	Scaffold	223	187 (Illumina)	5,309,671	
FWSEC0339	O111:H missing (O111:H11)	stX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	Not collected	Canada	SRR7947296	RRSF000000000	Scaffold	319	150 (Illumina)	5,736,056	
FWSEC0540	O111:H missing (O111:H11)	stX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	Not collected	Canada	SRR7947297	RRSG000000000	Scaffold	318	141 (Illumina)	5,736,086	
FWSEC0541	O111:H missing (O111:H11)	stX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	Not collected	Canada	SRR7947286	RRSI000000000	Scaffold	297	154 (Illumina)	5,759,118	
FWSEC0542	O111:H missing (O111:H11)	stX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	Not collected	Canada	SRR7947287	RRI000000000	Scaffold	327	97 (Illumina)	5,716,208	
FWSEC0543	O5:H missing (O5:H9)	stX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	Not collected	Canada	SRR7947271	RRSJ000000000	Scaffold	381	56 (Illumina)	5,198,826	
FWSEC0544	O111:H missing (O111:H11)	stX <sub>1a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	Not collected	Canada	SRR7947270	RRSK000000000	Scaffold	285	162 (Illumina)	5,698,696	
FWSEC0550	O22:H8	stX <sub>1a</sub> , stX <sub>2a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612323	RRL000000000	Scaffold	160	95 (Illumina)	4,924,792	
FWSEC0553	O117:H16	stX <sub>1a</sub>	Domesticated livestock, bovine, animal milk, agricultural (farm)	Not collected	Canada	SRR7947269	RRSM000000000	Scaffold	102	484 (Illumina)	5,129,595	
FWSEC0554	O26:H11	stX <sub>1a</sub>	Domesticated livestock, bovine, animal milk, agricultural (farm)	Not collected	Canada	SRR7947268	RRSN000000000	Scaffold	272	59 (Illumina)	5,256,407	
FWSEC0555	O22:H8	stX <sub>1a</sub> , stX <sub>2c</sub>	Domesticated livestock, bovine, animal milk, agricultural (farm)	Not collected	Canada	SRR7612324	RRSO000000000	Scaffold	116	50 (Illumina)	5,150,247	
FWSEC0563	O5:NM (O5:H9)	stX <sub>1a</sub>	River water	2012	Canada	SRR7612214	RRSU000000000	Scaffold	169	118 (Illumina)	5,346,500	
FWSEC0576	O165:NM (O165:H25)	stX <sub>1a</sub> , stX <sub>2a</sub>	Unknown specimen type	Not collected	Canada	SRR7612216	RRSW000000000	Scaffold	158	145 (Illumina)	4,934,747	
FWSEC0595	O26:NM (O26:H11)	stX <sub>2a</sub>	Stream water	2012	Canada	SRR7612366	RRSY000000000	Scaffold	90	118 (Illumina)	4,933,762	
FWSEC0597	O130:H11 (O130:H38)	stX <sub>1a</sub> , stX <sub>2a</sub>	Unknown specimen type	2012	Canada	SRR7612365	RRSZ000000000	Scaffold	123	132 (Illumina)	5,047,716	

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**TABLE 1** (Continued)

	Isolate identifier from this study	Traditional serogroup (in silico prediction) <sup>a</sup>	stx subtype (in silico prediction) <sup>b</sup>	Source	Yr of isolation	Country of isolation	SRA accession no.	WGS accession no. <sup>c</sup>	Assembly level	No. of contigs postassembly, chromosome and/or plasmid(s), size (bp) (GenBank accession no.)	No. of Coverage (>) CDS <sup>d</sup> (technology)	Genome size (all contigs) (bp)
FWSEC0611	O139:H19	STX <sub>1a</sub> , STX <sub>2a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612372	RRTF000000000	Scaffold	82	141 (Illumina)	5,012,638	
FWSEC0614	O182:NM (O182:H25)	STX <sub>2a</sub>	Stream water, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612501	RRTK000000000	Scaffold	149	103 (Illumina)	5,089,306	
FWSEC0620	O6:H34	STX <sub>2c</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612497	RRTK000000000	Scaffold	77	109 (Illumina)	4,882,643	
FWSEC0621	O136:H12	STX <sub>1a</sub>	Unknown specimen type	2012	Canada	SRR7612498	RRTL000000000	Scaffold	196	58 (Illumina)	5,260,253	
FWSEC0622	O136:H16	STX <sub>1a</sub>	Unknown specimen type	2012	Canada	SRR7612495	RRTM000000000	Scaffold	130	109 (Illumina)	5,284,097	
FWSEC0623	O136:H16 (O136:H12)	STX <sub>1a</sub>	Unknown specimen type	2012	Canada	SRR7612496	RRTN000000000	Scaffold	195	126 (Illumina)	5,217,336	
FWSEC0624	O26:H11	STX <sub>1a</sub>	Unknown specimen type	2012	Canada	SRR7612493	RRTQ000000000	Scaffold	179	216 (Illumina)	5,286,972	
FWSEC0625	O113:H4	STX <sub>1a</sub> , STX <sub>2d</sub>	Unknown specimen type	2012	Canada	SRR7612494	RRTP000000000	Scaffold	161	81 (Illumina)	4,907,794	
FWSEC0626	O139:H19	STX <sub>1a</sub> , STX <sub>2a</sub>	Rectal swab, domesticated livestock, bovine, agricultural (farm)	2012	Canada	SRR7612134	RRTQ000000000	Scaffold	83	86 (Illumina)	5,015,665	
FWSEC0627	O113:NM (O113:H121)	STX <sub>2a</sub>	Unknown specimen type	2012	Canada	SRR7612133	RRTR000000000	Scaffold	103	94 (Illumina)	4,992,326	
FWSEC0629	O130:H38	STX <sub>1a</sub> , STX <sub>2a</sub>	Domesticated livestock, farm animal, porcine, animal manure, agricultural (farm)	Not collected	Not collected	SRR7947266	RRTS000000000	Scaffold	264	60 (Illumina)	5,029,373	
FWSEC0631	O69:H11	STX <sub>1a</sub>	Unknown specimen type	2012	Canada	SRR7947264	RRTT000000000	Scaffold	494	50 (Illumina)	5,209,350	
FWSEC0632	O111:H missing (O111:H8)	STX <sub>1a</sub> , STX <sub>2a</sub>	Domesticated livestock, bovine, animal manure, agricultural (farm)	Not collected	Not collected	SRR7947263	RRTU000000000	Scaffold	269	80 (Illumina)	5,455,050	

<sup>a</sup>NT, nontypeable; NM, nonmotile, by traditional lab determination. Isolates were traditionally serotyped at national or provincial reference labs prior to selection for sequencing. Serogroups were then determined algorithmically using ECTyper ([https://github.com/phac-nml/ecoli\\_serotyping](https://github.com/phac-nml/ecoli_serotyping)) under default settings (min\_percentIdentity 90; percentLength 50). When traditional and *in silico* serogroup calls varied, both are reported. It is not unusual for genotypically positive strains to be nonmotile under lab determination conditions.

<sup>b</sup>stx allele subtype PCR determination (33) (with algorithmic determination, where noted). Genome assemblies underwent a default BLASTN search (90% threshold for %ID; 60% minimum length) against the Virulence Finder 2.0 Web-based tool (<https://cge.cbs.dtu.dk/services/VirulenceFinder/>) (34). Reads were mapped using SRST2 (35) and the K-mer analysis toolkit (KAT) sect function (36) to the Flemming Scheutz-curated *E. coli* virulence database. Although all isolates had been previously determined as genotypically positive for at least one stx subtype by PCR, both the A and B subunits of each Stx complex (stx<sub>1</sub> or stx<sub>2</sub>) had to be detected during *in silico* screening to be classified as genotypically present.

<sup>c</sup>WGS, whole-genome sequencing.

<sup>d</sup>Coding sequences (CDS) for reference genomes FWSEC001 through FWSEC011 only.

<sup>e</sup>A linear sequence in FWSEC001 that could not be circularized owing to unresolved or collapsed repeats.

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