

<u>Genome</u>	<u>Serotype</u>	<u>Pathovar</u>	<u>Phylogroup</u>	<u>SRA accession</u>	<u>Sequencing Platform</u>
C227-11	O104:H4	EAEC/STEC	B1	SRR341579	HiSeq
86-24	O157:H7	STEC	E	SRR1639082	HiSeq
C844-97	O157:H45	aEPEC	B2	SRR1772997	HiSeq
sakai	O157:H7	STEC	E	SRR587217	MiSeq
C260-92	O127:K-:H40	aEPEC	A	SRR1773018	HiSeq
KTE9	unknown	N/A	D	SRR640937	HiSeq

<u>read length</u>	<u>average positions for accurate placement &gt; 95%</u>	<u>read coverage for</u>
100	979	<0.01x
100	932	<0.01x
100	1062	<0.01x
250	361	<0.01x
100	73664	0.42x
100	2663	0.14x