

Supplemental Table 1

Mitchell et al. Multi-laboratory evaluation of the Illumina iSeq platform for whole genome sequencing of *Salmonella*, *Escherichia coli*, and *Listeria*

Additional identifiers for isolates

Organism	V-CLASP ID/GenomeTrakr PT ID/NCBI Accession #	ST	Run #
<i>Salmonella enterica</i> ser. Albany	V-CLASP-D-59	292	
<i>Salmonella enterica</i> ser. Typhimurium	V-CLASP-D-01	19	
<i>Salmonella enterica</i> ser. Enteritidis	V-CLASP-D-40	11	1 [^] & 4*
<i>Salmonella enterica</i> ser. I 4,[5],12:i:-	V-CLASP-C-02	19	
<i>Salmonella enterica</i> ser. Albany	V-CLASP-D-59	292	
<i>Salmonella enterica</i> ser. Typhimurium	V-CLASP-D-01	19	
<i>Salmonella enterica</i> ser. Enteritidis	V-CLASP-D-40	11	2 [^]
<i>Salmonella enterica</i> ser. I 4,[5],12:i:-	V-CLASP-C-02	19	
<i>Salmonella enterica</i> ser. Derby	V-CLASP-D-18	40	
<i>Salmonella enterica</i> ser. Derby	V-CLASP-D-18	40	
<i>Salmonella enterica</i> ser. Newport	V-CLASP-D-12	5	3 [^]
<i>Salmonella enterica</i> ser. Newport	V-CLASP-D-12	5	
<i>Salmonella enterica</i> ser. Dublin	CP032396	10	
<i>Salmonella enterica</i> ser. Dublin	CP032393	10	
<i>Salmonella enterica</i> ser. Dublin	CP032390	10	5 [^]
<i>Salmonella enterica</i> ser. Dublin	CPO32387	10	
<i>Escherichia coli</i> ser. O157:H7	ECP19-198	11	
<i>Escherichia coli</i> ser. O157:H7	ECP19-598	11	1*
<i>Escherichia coli</i> ser. O157:H7	ECP19-798	11	
<i>Escherichia coli</i> ser. O157:H7	ECP19-2498	11	
<i>Listeria monocytogenes</i>	LMP18-H8393	3	
<i>Listeria monocytogenes</i>	LMP18-H2446	7	
<i>Escherichia coli</i> O9:H10	N17EC0320	540	
<i>Escherichia coli</i> O89:H9	N17EC0326	10	
<i>Escherichia coli</i> O83:H42	N17ECO616	1485	2* & 3*
<i>Escherichia coli</i> O117:H42	N17EC1164	2207	
<i>Listeria monocytogenes</i>	FDA00008248	554	
<i>Listeria monocytogenes</i>	FDA00006667	6	

[^]: Nextera XT; *: Nextera DNA Flex