Whole-Genome Sequences of 12 Clinical Strains of *Listeria monocytogenes*

Guojie Cao,a,b Jianmin Zhang,a Xuebin Xu,c Huiming Jin,c Xiaowei Yang,a Haijian Pan,a Marc Allard,d Eric Brown,d Jianghong Meng,a,b

Department of Food Science & Engineering, Shanghai Jiao Tong University, Shanghai, China; Department of Nutrition & Food Science and Joint Institute for Food Safety & Applied Nutrition, University of Maryland, College Park, Maryland, USA; Shanghai Municipal Center for Disease Control & Prevention, Shanghai, China; Division of Microbiology, Center for Food Safety & Applied Nutrition, Food and Drug Administration, College Park, Maryland, USA. G.C. and J.Z. contributed equally to this work.

*L. monocytogenes* is a foodborne pathogen of global concern due to the high mortality rate among immunocompromised patients. Whole-genome sequences of 12 strains of *L. monocytogenes* from humans were reported. The availability of these genomes should provide useful information on the evolutionary history and genetic diversity of *L. monocytogenes*.

Whole-genome sequences of 12 clinical strains of *L. monocytogenes* were generated using Nextera XT and MiSeq (Illumina, San Diego, CA) sequencing platforms. Genomic DNA was isolated from overnight cultures and sequenced. Genomes were assembled using Newbler 2.6 (罗氏, Branford, CT), including SHL001 (33 contigs, 2.95 Mb, 2,964 identified genes), SHL002 (22 contigs, 3.12 Mb, 301,779 bp contig size, and 2,806 identified genes), SHL003 (17 contigs, 3.01 Mb, 476,139 bp contig size, and 3,018 identified genes), SHL004 (18 contigs, 3.01 Mb, 579,300 bp contig size, and 3,113 identified genes), SHL005 (17 contigs, 2.88 Mb, 437,049 bp contig size, and 2,859 identified genes), SHL006 (30 contigs, 2.93 Mb, 476,139 bp contig size, and 2,959 identified genes), SHL007 (43 contigs, 2.98 Mb, 355,359 bp contig size, and 2,995 identified genes), SHL008 (26 contigs, 3.01 Mb, 293,078 bp contig size, and 2,990 identified genes), SHL009 (32 contigs, 2.87 Mb, 541,739 bp contig size, and 2,866 identified genes), SHL010 (84 contigs, 3.08 Mb, 259,950 bp contig size, and 3,112 identified genes), SHL011 (17 contigs, 2.87 Mb, 543,519 bp contig size, and 2,847 identified genes), SHL012 (23 contigs, 2.93 Mb, 355,398 bp contig size, and 2,906 identified genes), and SHL013 (20 contigs, 2.86 Mb, 358,858 bp contig size, and 2,806 identified genes). These sequences were annotated using the NCBI Prokaryotic Genomes Automatic Annotation Pipeline (10).

**Nucleotide sequence accession numbers.** The draft genome sequences of these 12 *L. monocytogenes* strains are available in GenBank under the accession numbers APIB00000000, APIC00000000, APIH00000000, APIJ00000000, APIG00000000, APIH00000000, APIF00000000, APIK00000000, APIL00000000, and APIM00000000.

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**REFERENCES**


